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(54) **USE OF ANTI-CGRP ANTIBODIES AND ANTIBODY FRAGMENTS TO PREVENT OR INHIBIT PHOTOPHOBIA OR LIGHT A VERSION IN SUBJECTS IN NEED THEREOF, ESPECIALLY MIGRAINE SUFFERERS**

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None
See application file for complete search history.

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(57) **ABSTRACT**

The present invention is directed to methods of inhibiting or preventing photophobia in subjects in need thereof using anti-CGRP antibodies or antibody fragments that inhibit photophobia, especially CGRP-associated photophobia. These antibodies and fragments are useful in treating different disorders associated with photophobia such as migraine, cluster headaches and the like.

The present invention also provides assays using transgenic Nestin/Ramp1 rodents, utilizing a CGRP model light aversive behavior model for identifying therapeutically effective anti-CGRP antibodies and fragments thereof having binding specificity for CGRP which inhibit or prevent photophobia in subjects in need thereof. The present invention is specifically directed to methods for identifying therapeutically effective antibodies and fragments thereof having binding specificity for CGRP that may be used to treat CGRP associated disorders such as migraine. Specifically, this invention relates to assays and therapies using the antibodies described herein to inhibit or prevent photophobia, and binding fragments thereof, comprising the sequences of the V_H, V_L and CDR polypeptides described herein, and the polynucleotides encoding them.

10 Claims, 70 Drawing Sheets

Specification includes a Sequence Listing.

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Figure 1A

Ab1

Ab1 Heavy chain (chimera) Full length protein sequence.

QSLEESGGRLVTPGTPLTLTCTVSGLDLSSYYMQWVRQAPGKGLEWIGVIGINDNTYYASWAKGRFTISRASSTTVDLKMTS
LTTEDTATYFCARGDIWGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV
LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTP
EVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTIS
KAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
QGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 4)

Ab1 Variable region heavy chain (chimera) protein sequence.

QSLEESGGRLVTPGTPLTLTCTVSGLDLSSYYMQWVRQAPGKGLEWIGVIGINDNTYYASWAKGRFTISRASSTTVDLKMTS
LTTEDTATYFCARGDIWGPGTLVTVSS (SEQ ID NO: 3)

Ab1 Variable region heavy chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QSLEESGGRLVTPGTPLTLTCTVSGLDLS**SYYMQ**WVRQAPGKGLEWIGVIGINDNTYYASWAKGRFTISRASSTTVDLKMTS
LTTEDTATYFCAR*GDI*WGPGTLVTVSS (SEQ ID NOS: 8, 9, 10, respectively)

Ab1 Variable region heavy chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCTGACACTCACCTGCACAGTCTCTGGACTCG
ACCTCAGT**AGCTACTACATGCAA**TGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATTGGTATTA
ATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGCCTCGTCGACCACGGTGGATCTGA
AAATGACCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGA*GGGGACATC*TGGGGCCCAGGCACCCTCGT
CACCGTCTCGAGC (SEQ ID NO: 143)

Figure 1B

Ab1 Heavy chain (chimera) Full length DNA sequence.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCTGACACTCACCTGCACAGTCTCTGGACTCG
ACCTCAGTAGCTACTACATGCAATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATTGGTATTA
ATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGCCTCGTCGACCACGGTGGATCTGA
AAATGACCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGGGGCCCAGGCACCCTCG
TCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCAC
AGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCTGACCAG
CGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGC
AGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTTGAGCCC
AAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCTCTTCC
CCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAG
ACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAGCAG
TACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGC
AAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACCACAG
GTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATC
CCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTGGACT
CCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTC
CGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 144)

Ab1 Light chain (chimera) Full length protein sequence.

QVLTQTASPVSAAVGSTVTINCQASQSVYDNNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDLECAD
AATYYCLGSYDCSSGDCFVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 2)

Ab1 Variable region light chain (chimera) protein sequence.

QVLTQTASPVSAAVGSTVTINCQASQSVYDNNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDLECAD
AATYYCLGSYDCSSGDCFVFGGGTEVVVKR (SEQ ID NO: 1)

Figure 1C

Ab1 Variable region light chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQTASPVSAAVGSTVTINC**QASQSVYDNNYLA**WYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDLECA
DAATYYC*LGSYDCSSGDCFI*FGGGTEVVVKR (SEQ ID NOS: 5, 6, 7, respectively)

Ab1 Variable region light chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATGATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGAC
CTGGAGTGTGCCGATGCTGCCACTTACTACTGT*CTAGGCAGTTATGATTGTAGTAGTGGTGATTGTTTTGTT*TTCGGCGGAG
GGACCGAGGTGGTGGTCAAACGT (SEQ ID NO: 141)

Ab1 Light chain (chimera) Full length DNA sequence.

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGCCAGGCCAGTCAG
AGTGTTTATGATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTACAT
CCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGACCT
GGAGTGTGCCGATGCTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATTGTTTTGTTTTCGGCGGAG
GGACCGAGGTGGTGGTCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATC
TGGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCC
CTCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTG
ACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTC
ACAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 142)

Figure 2A

Ab2

Ab2 Heavy chain (humanized) Full length protein sequence – mammalian produced.

EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGVIGINDNTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV
HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL
MISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPA
PIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 14)

Ab2 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGVIGINDNTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSS (SEQ ID NO: 13)

Ab2 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGLDLS**SYYMQ**WVRQAPGKGLEWVGVIGINDNTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCAR*GDI*WGQGTLVTVSS (SEQ ID NOS: 18, 19, 20, respectively)

Ab2 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAC
TCGACCTCAGT**AGCTACTACATGCAA**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
TCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGA*GGGGACATC*TGGGGCCAAGGGAC
CCTCGTCACCGTCTCGAGC (SEQ ID NO: 153)

Figure 2B

Ab2 Heavy chain (humanized) Full length DNA sequence – mammalian produced.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAC
TCGACCTCAGTAGCTACTACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
TCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 154)

Ab2 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSSGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQ
ESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 12)

Ab2 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSSGDCFVFGGGTKVEIKR (SEQ ID NO: 11)

Figure 2C

Ab2 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQSVYDNNYLA**WYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCSSGDCFV*FGGGTKVEIKR (SEQ ID NOS: 15, 16, 17, respectively)

Ab2 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATGATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTAGGCAGTTATGATTGTAGTAGTGGTGATTGTTTTGTT*TTCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 151)

Ab2 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
GTGTTTATGATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTACATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 152)

Figure 3A
Ab3

Ab3 Heavy chain (humanized) Full length protein sequence – yeast produced.

EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGVIGINDNTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV
HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDARVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL
MISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPA
PIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 24)

Ab3 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGVIGINDNTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSS (SEQ ID NO: 23)

Ab3 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGLDLS**SYYMQ**WVRQAPGKGLEWVGVIGINDNTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCAR*GDI*WGQGTLVTVSS (SEQ ID NOS: 28, 29, 30, respectively)

Ab3 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAC
TCGACCTCAGT**AGCTACTACATGCAA**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
TCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGA*GGGGACATC*TGGGGCCAAGGGAC
CCTCGTCACCGTCTCGAGC (SEQ ID NO: 163)

Figure 3B

Ab3 Heavy chain (humanized) Full length DNA sequence – yeast produced.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAC
TCGACCTCAGTAGCTACTACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
TCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACGCGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 164)

Ab3 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSSGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQ
ESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 22)

Ab3 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSSGDCFVFGGGTKVEIKR (SEQ ID NO: 21)

Figure 3C

Ab3 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQSVYDNNYLA**WYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCSSGDCFV*FGGGTKVEIKR (SEQ ID NOS: 25, 26, 27, respectively)

Ab3 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATGATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTAGGCAGTTATGATTGTAGTAGTGGTGATTGTTTTGTT*TTCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 161)

Ab3 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
GTGTTTATGATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTACATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 162)

Figure 4A
Ab4

Ab4 Heavy chain (chimera) Full length protein sequence.

QSLEESGGRLVTPGTPLTLTCSVSGIDLSGYYMNWVRQAPGKGLEWIGVIGINGATYYASWAKGRFTISKTSSTTVDLKMTS
LTTEDTATYFCARGDIWGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV
LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTP
EVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTIS
KAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
QGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 34)

Ab4 Variable region heavy chain (chimera) protein sequence.

QSLEESGGRLVTPGTPLTLTCSVSGIDLSGYYMNWVRQAPGKGLEWIGVIGINGATYYASWAKGRFTISKTSSTTVDLKMTS
LTTEDTATYFCARGDIWGPGTLVTVSS (SEQ ID NO: 33)

Ab4 Variable region heavy chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QSLEESGGRLVTPGTPLTLTCSVSGIDLS**GYYMN**WVRQAPGKGLEWIGVIGINGATYYASWAKGRFTISKTSSTTVDLKMTS
LTTEDTATYFCAR*GDI*WGPGTLVTVSS (SEQ ID NOS: 38, 39, 40, respectively)

Ab4 Variable region heavy chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCTGACACTCACCTGTTCCGTCTCTGGCATCG
ACCTCAGT**GGCTACTACATGAAC**TGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATTGGTATT
AATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAAAACCTCGTCGACCACGGTGGATCTG
AAAATGACCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGA*GGGGACATC*TGGGGCCCGGGCACCCTC
GTCACCGTCTCGAGC (SEQ ID NO: 173)

Figure 4B

Ab4 Heavy chain (chimera) Full length DNA sequence.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCTGACACTCACCTGTTCCGTCTCTGGCATCG
ACCTCAGTGGCTACTACATGAACTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATTGGTATTA
ATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAAAACCTCGTCGACCACGGTGGATCTGA
AAATGACCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGGGGCCCGGGCACCCTCG
TCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCAC
AGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCTGACCAG
CGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGC
AGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTTGAGCCC
AAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCTCTTCC
CCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAG
ACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAGCAG
TACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGC
AAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACCACAG
GTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATC
CCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTGGACT
CCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTC
CGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 174)

Ab4 Light chain (chimera) Full length protein sequence.

QVLTQTPSPVSAAVGSTVTINCQASQSVYHNTYLAWYQQKPGQPPKQLIYDASTLASGVPSRFSGSGSGTQFTLTISGVQCND
AAAYYCLGSYDCTNGDCFVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 32)

Ab4 Variable region light chain (chimera) protein sequence.

QVLTQTPSPVSAAVGSTVTINCQASQSVYHNTYLAWYQQKPGQPPKQLIYDASTLASGVPSRFSGSGSGTQFTLTISGVQCND
AAAYYCLGSYDCTNGDCFVFGGGTEVVVKR (SEQ ID NO: 31)

Figure 4C

Ab4 Variable region light chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQTPSPVSAAVGSTVTINC**QASQSVYHNTYLA**WYQQKPGQPPKQLIY**DASTLAS**GVPSRFSGSGSGTQFTLTISGVQCN
DAAAYYC*LGSYDCTNGDCF*TFGGGTEVVVKR (SEQ ID NOS: 35, 36, 37, respectively)

Ab4 Variable region light chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATCATAACACCTACCTGGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAACAACTGATCTATGATGC
ATCCACTCTGGCGTCTGGGGTCCCATCGCGGTTCAGCGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGGC
GTGCAGTGTAACGATGCTGCCGCTTACTACTGT*CTGGGCAGTTATGATTGTACTAATGGTGATTGTTTTGTT*TTCGGCGGAG
GGACCGAGGTGGTGGTCAAACGT (SEQ ID NO: 171)

Ab4 Light chain (chimera) Full length DNA sequence.

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGCCAGGCCAGTCAGA
GTGTTTATCATAACACCTACCTGGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAACAACTGATCTATGATGCATC
CACTCTGGCGTCTGGGGTCCCATCGCGGTTCAGCGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGGCGTG
CAGTGTAACGATGCTGCCGCTTACTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGTTTTGTTTTCGGCGGAGG
GACCGAGGTGGTGGTCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 172)

Figure 5A
Ab5

Ab5 Heavy chain (humanized) Full length protein sequence – mammalian produced.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGVIGINGATYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV
HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL
MISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPA
PIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 44)

Ab5 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGVIGINGATYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSS (SEQ ID NO: 43)

Ab5 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLS**GYYMN**WVRQAPGKGLEWVGVIGINGATYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCAR*GDI*WGQGTLVTVSS (SEQ ID NOS: 48, 49, 50, respectively)

Ab5 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACCTCAGT**GGCTACTACATGAAC**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGT
ATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTG
TATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGA*GGGGACATC*TGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGC (SEQ ID NO: 183)

Figure 5B

Ab5 Heavy chain (humanized) Full length DNA sequence – mammalian produced.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACCTCAGTGGCTACTACATGAACTGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
TTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 184)

Ab5 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIYDASTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCTNGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 42)

Ab5 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIYDASTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCTNGDCFVFGGGTKVEIKR (SEQ ID NO: 41)

Figure 5C

Ab5 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQSVYHNTYLA**WYQQKPGKVPKQLIYDASTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCTNGDCFV*FGGGTKVEIKR (SEQ ID NOS: 45, 46, 47, respectively)

Ab5 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATCATAACACCTACCTGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATGATGC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTGGGCAGTTATGATTGTACTAATGGTGATTGTTTTGTT*TTCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 181)

Ab5 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
GTGTTTATCATAACACCTACCTGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATGATGCATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 182)

Figure 6A

Ab6

Ab6 Heavy chain (humanized) Full length protein sequence – yeast produced.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGVIGINGATYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV
HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDARVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL
MISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPA
PIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 54)

Ab6 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGVIGINGATYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSS (SEQ ID NO: 53)

Ab6 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLS**GYYMN**WVRQAPGKGLEWVGVIGINGATYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCAR*GDI*WGQGTLVTVSS (SEQ ID NOS: 58, 59, 60, respectively)

Ab6 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACCTCAGT**GGCTACTACATGAAC**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGT
ATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTG
TATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGA*GGGGACATC*TGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGC (SEQ ID NO: 193)

Figure 6B

Ab6 Heavy chain (humanized) Full length DNA sequence – yeast produced.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACCTCAGTGGCTACTACATGAACTGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
TTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACGCGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 194)

Ab6 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIYDASTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCTNGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 52)

Ab6 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIYDASTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCTNGDCFVFGGGTKVEIKR (SEQ ID NO: 51)

Figure 6C

Ab6 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQSVYHNTYLA**WYQQKPGKVPKQLIYDASTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCTNGDCFV*FGGGTKVEIKR (SEQ ID NOS: 55, 56, 57, respectively)

Ab6 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATCATAACACCTACCTGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATGATGC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTGGGCAGTTATGATTGTACTAATGGTGATTGTTTTGTT*TTCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 191)

Ab6 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
GTGTTTATCATAACACCTACCTGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATGATGCATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 192)

Figure 7A

Ab7

Ab7 Heavy chain (chimera) Full length protein sequence.

QEQLKESGGRLVTPGTSLTLTCTVSGIDLSNHYMQWVRQAPGKGLEWIGVVGINGRTYYASWAKGRFTISRTSSTTVDLKM
TRLTTEDTATYFCARGDIWGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFP
AVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISR
TPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKT
ISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW
QQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 64)

Ab7 Variable region heavy chain (chimera) protein sequence.

QEQLKESGGRLVTPGTSLTLTCTVSGIDLSNHYMQWVRQAPGKGLEWIGVVGINGRTYYASWAKGRFTISRTSSTTVDLKM
TRLTTEDTATYFCARGDIWGPGTLVTVSS (SEQ ID NO: 63)

Ab7 Variable region heavy chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QEQLKESGGRLVTPGTSLTLTCTVSGIDLS**NHYMQ**WVRQAPGKGLEWIGVVGINGRTYYASWAKGRFTISRTSSTTVDLKM
TRLTTEDTATYFCAR*GDI*WGPGTLVTVSS (SEQ ID NOS: 68, 69, 70, respectively)

Ab7 Variable region heavy chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAGGAGCAGCTGAAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACATCCCTGACACTCACCTGCACCGTCTCTGGA
ATCGACCTCAGT**AACCACTACATGCAA**TGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGAGTCGTTGG
TATTAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAACCTCGTCGACCACGGTGGAT
CTGAAAATGACCAGGCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGA*GGGGACATC*TGGGGCCCAGGCACC
CTGGTCACCGTCTCGAGC (SEQ ID NO: 203)

Figure 7B

Ab7 Heavy chain (chimera) Full length DNA sequence.

CAGGAGCAGCTGAAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACATCCCTGACACTCACCTGCACCGTCTCTGGA
ATCGACCTCAGTAACCACTACATGCAATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGAGTCGTTGGT
ATTAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAACCTCGTCGACCACGGTGGAT
CTGAAAATGACCAGGCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGGGGCCCAGGCACC
CTGGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGG
GCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCTGA
CCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTC
CAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTTGA
GCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCTC
TTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACG
AAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAG
CAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAG
TGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACCA
CAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTC
TATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTG
GACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCAT
GCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 204)

Ab7 Light chain (chimera) Full length protein sequence.

QVLTQTASPVSAAVGSTVTINCQASQSVYNYNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDVQCD
DAATYYCLGSYDCSTGDCFVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGN
SQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 62)

Ab7 Variable region light chain (chimera) protein sequence.

QVLTQTASPVSAAVGSTVTINCQASQSVYNYNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDVQCD
DAATYYCLGSYDCSTGDCFVFGGGTEVVVKR (SEQ ID NO: 61)

Figure 7C

Ab7 Variable region light chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQTASPVSAAVGSTVTINC**QASQSVYNYNYLA**WYQQKPGQPPKQLIY**STSTLAS**GVSSRFKGSGSGTQFTLTISDVQCD
DAATYYC*LGSYDCSTGDCFV*FGGGTEVVVKR (SEQ ID NOS: 65, 66, 67, respectively)

Ab7 Variable region light chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTATAATTACAACTACCTTGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTACA
TCCACTCTGGCATCTGGGGTCTCATCGCGATTCAAAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGACG
TGCAGTGTGACGATGCTGCCACTTACTACTGT*CTAGGCAGTTATGACTGTAGTACTGGTGATTGTTTTGTT*TTCGGCGGAGG
GACCGAGGTGGTGGTCAAACGT (SEQ ID NO: 201)

Ab7 Light chain (chimera) Full length DNA sequence.

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGCCAGGCCAGTCAG
AGTGTTTATAATTACAACTACCTTGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTACAT
CCACTCTGGCATCTGGGGTCTCATCGCGATTCAAAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGACGT
GCAGTGTGACGATGCTGCCACTTACTACTGTCTAGGCAGTTATGACTGTAGTACTGGTGATTGTTTTGTTTTCGGCGGAG
GGACCGAGGTGGTGGTCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATC
TGGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCC
CTCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTG
ACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTC
ACAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 202)

Figure 8A

Ab8

Ab8 Heavy chain (humanized) Full length protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSNHYMQWVRQAPGKGLEWVGVVGINGRTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV
HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL
MISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPA
PIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 74)

Ab8 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSNHYMQWVRQAPGKGLEWVGVVGINGRTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSS (SEQ ID NO: 73)

Ab8 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGIDLS**NHYMQ**WVRQAPGKGLEWVGVVGINGRTYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCAR*GDI*WGQGTLVTVSS (SEQ ID NOS: 78, 79, 80, respectively)

Ab8 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACCTCAGT**AACCACTACATGCAA**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCGTTGGTA
TCAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGA*GGGGACAT*CTGGGGCCAAGGGAC
CCTCGTCACCGTCTCGAGC (SEQ ID NO: 213)

Figure 8B

Ab8 Heavy chain (humanized) Full length DNA sequence.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACCTCAGTAACCACTACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCGTTGGTA
TCAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 214)

Ab8 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYNYNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSTGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 72)

Ab8 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQSVYNYNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSTGDCFVFGGGTKVEIKR (SEQ ID NO: 71)

Figure 8C

Ab8 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQSVYNYNYLA**WYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCSTGDCFT*FGGGTKVEIKR (SEQ ID NOS: 75, 76, 77, respectively)

Ab8 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AGTGTTTACAATTACAACTACCTTGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTGGGCAGTTATGATTGTAGTACTGGTGATTGTTTTGTT*TTCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 211)

Ab8 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
GTGTTTACAATTACAACTACCTTGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTACATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTACTGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 212)

Figure 9A

Ab9

Ab9 Heavy chain (chimera) Full length protein sequence.

QSLEESGGRLVTPGTPLTLTCTVSGIGLSSYYMQWVRQSPGRGLEWIGVIGSDGKTYYATWAKGRFTISKTSSTTVDLRMAS
LTTEDTATYFCTRGDIWGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAV
LQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTP
EVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTIS
KAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
QGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 84)

Ab9 Variable region heavy chain (chimera) protein sequence.

QSLEESGGRLVTPGTPLTLTCTVSGIGLSSYYMQWVRQSPGRGLEWIGVIGSDGKTYYATWAKGRFTISKTSSTTVDLRMAS
LTTEDTATYFCTRGDIWGPGTLVTVSS (SEQ ID NO: 83)

Ab9 Variable region heavy chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QSLEESGGRLVTPGTPLTLTCTVSGIGLS**SYYMQ**WVRQSPGRGLEWIGVIGSDGKTYYATWAKGRFTISKTSSTTVDLRMAS
LTTEDTATYFCTR*GDI*WGPGTLVTVSS (SEQ ID NOS: 88, 89, 90, respectively)

Ab9 Variable region heavy chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCTGACACTCACCTGCACAGTCTCTGGAATCG
GCCTCAGT**AGCTACTACATGCAG**TGGGTCCGCCAGTCTCCAGGGAGGGGGCTGGAATGGATCGGAGTCATTGGTAGT
GATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCACCATCTCCAAGACCTCGTCGACCACGGTGGATCTG
AGAATGGCCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTACCAGA*GGGGACATC*TGGGGCCCGGGGACCCTC
GTCACCGTCTCGAGC (SEQ ID NO: 223)

Figure 9B

Ab9 Heavy chain (chimera) Full length DNA sequence.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCTGACACTCACCTGCACAGTCTCTGGAATCG
GCCTCAGTAGCTACTACATGCAGTGGGTCCGCCAGTCTCCAGGGAGGGGGCTGGAATGGATCGGAGTCATTGGTAGTG
ATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCACCATCTCCAAGACCTCGTCGACCACGGTGGATCTGA
GAATGGCCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTACCAGAGGGGACATCTGGGGCCCGGGGACCCTCG
TCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCAC
AGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCTGACCAG
CGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGC
AGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTTGAGCCC
AAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCTCTTCC
CCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAG
ACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAGCAG
TACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGC
AAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACCACAG
GTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATC
CCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTGGACT
CCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTC
CGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 224)

Ab9 Light chain (chimera) Full length protein sequence.

QVLTQTPSPVSAAVGSTVTINCQASQNVYNNNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFRGSGSGTQFTLTISDVQCD
DAATYYCLGSYDCSRGDCFVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGN
SQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 82)

Ab9 Variable region light chain (chimera) protein sequence.

QVLTQTPSPVSAAVGSTVTINCQASQNVYNNNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFRGSGSGTQFTLTISDVQCD
DAATYYCLGSYDCSRGDCFVFGGGTEVVVKR (SEQ ID NO: 81)

Figure 9C

Ab9 Variable region light chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQTPSPVSAAVGSTVTINC**QASQNVYNNNYLA**WYQQKPGQPPKQLIY**STSTLAS**GVSSRFRGSGSGTQFTLTISDVQCD
DAATYYC*LGSYDCSRGDCFV*FGGGTEVVVKR (SEQ ID NOS: 85, 86, 87, respectively)

Ab9 Variable region light chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGC**CAGGCCAGTCAG**
AATGTTTATAATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTAC
GTCCACTCTGGCATCTGGGGTCTCATCGCGATTCAGAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGAC
GTGCAGTGTGACGATGCTGCCACTTACTACTGT*CTAGGCAGTTATGATTGTAGTCGTGGTGATTGTTTTGTTT*TCGGCGGAG
GGACCGAGGTGGTGGTCAAACGT (SEQ ID NO: 221)

Ab9 Light chain (chimera) Full length DNA sequence.

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCACAGTCACCATCAATTGCCAGGCCAGTCAGA
ATGTTTATAATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTACGTC
CACTCTGGCATCTGGGGTCTCATCGCGATTCAGAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGACGTG
CAGTGTGACGATGCTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTCGTGGTGATTGTTTTGTTTTCGGCGGAGG
GACCGAGGTGGTGGTCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 222)

Figure 10A
Ab10

Ab10 Heavy chain (humanized) Full length protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGVIGSDGKTYYATWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCTRGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVH
TFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLM
ISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPI
EKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 94)

Ab10 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGVIGSDGKTYYATWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCTRGDIWGQGTLVTVSS (SEQ ID NO: 93)

Ab10 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGIGL**SSYYMQ**WVRQAPGKGLEWVGVIGSDGKTYYATWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCTR*GDI*WGQGTLVTVSS (SEQ ID NOS: 98, 99, 100, respectively)

Ab10 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGGCCTCAGT**AGCTACTACATGCAA**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
GTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGA*GGGGACATC*TGGGGCCAAGGGAC
CCTCGTCACCGTCTCGAGC (SEQ ID NO: 233)

Figure 10B

Ab10 Heavy chain (humanized) Full length DNA sequence.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGGCCTCAGTAGCTACTACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
GTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 234)

Ab10 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSRGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 92)

Ab10 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSRGDCFVFGGGTKVEIKR (SEQ ID NO: 91)

Figure 10C

Ab10 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQNVYNNNYLA**WYQQKPGKVPKQLIY**STSTLAS**GVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCSRGDCFV*FGGGTKVEIKR (SEQ ID NOS: 95, 96, 97, respectively)

Ab10 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AATGTTTACAATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTGGGCAGTTATGATTGTAGTCGTGGTGATTGTTTTGTTT*TTCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 231)

Ab10 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
ATGTTTACAATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTACATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTCGTGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 232)

Figure 11A
Ab11

Ab11 Heavy chain (chimera) Full length protein sequence.

QSLEESGGRLVTPGGSLTLTCTVSGIDVTNYYMQWVRQAPGKGLEWIGVIGVNGKRYYASWAKGRFTISKTSSTTVDLKMT
SLTTEDTATYFCARGDIWGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPA
VLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT
PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTI
SKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQ
QGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 104)

Ab11 Variable region heavy chain (chimera) protein sequence.

QSLEESGGRLVTPGGSLTLTCTVSGIDVTNYYMQWVRQAPGKGLEWIGVIGVNGKRYYASWAKGRFTISKTSSTTVDLKMT
SLTTEDTATYFCARGDIWGPGTLVTVSS (SEQ ID NO: 103)

Ab11 Variable region heavy chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined: CDR3: Italics.

QSLEESGGRLVTPGGSLTLTCTVSGIDVT**NYYMQ**WVRQAPGKGLEWIGVIGVNGKRYYASWAKGRFTISKTSSTTVDLKMT
SLTTEDTATYFCAR*GDI*WGPGTLVTVSS (SEQ ID NOS: 108, 109, 110, respectively)

Ab11 Variable region heavy chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined: CDR3: Italics.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGAGGATCCCTGACACTCACCTGCACAGTCTCTGGAATCG
ACGTCACT**AACTACTATATGCAA**ATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATTGGTGTGA
ATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAAAACCTCGTCGACCACGGTGGATCTGA
AAATGACCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGA*GGCGACATC*TGGGGCCCGGGGACCCTCGT
CACCGTCTCGAGC (SEQ ID NO: 243)

Figure 11B

Ab11 Heavy chain (chimera) Full length DNA sequence.

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGAGGATCCCTGACACTCACCTGCACAGTCTCTGGAATCG
ACGTCACTAACTACTATATGCAATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATTGGTGTGA
ATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAAAACCTCGTCGACCACGGTGGATCTGA
AAATGACCAGTCTGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGCGACATCTGGGGCCCCGGGGACCCTCG
TCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCAC
AGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCTGACCAG
CGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGC
AGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTTGAGCCC
AAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCTCTTCC
CCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAG
ACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAGCAG
TACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGC
AAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACCACAG
GTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATC
CCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCTGGACT
CCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTC
CGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 244)

Ab11 Light chain (chimera) Full length protein sequence.

QVLTQTASPVSPAVGSTVTINCRASQSVYYNNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDVQCDD
AATYYCLGSYDCSNGDCFVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 102)

Ab11 Variable region light chain (chimera) protein sequence.

QVLTQTASPVSPAVGSTVTINCRASQSVYYNNYLAWYQQKPGQPPKQLIYSTSTLASGVSSRFKGSGSGTQFTLTISDVQCDD
AATYYCLGSYDCSNGDCFVFGGGTEVVVKR (SEQ ID NO: 101)

Figure 11C

Ab11 Variable region light chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQTASPVSPAVGSTVTINC**RASQSVYYNNYLA**WYQQKPGQPPKQLIY<u>STSTLAS</u>GVSSRFKGSGSGTQFTLTISDVQCD
DAATYYC*LGSYDCSNGDCFV*FGGGTEVVVKR (SEQ ID NOS: 105, 106, 107, respectively)

Ab11 Variable region light chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAGGTGCTGACCCAGACTGCATCCCCCGTGTCTCCAGCTGTGGGAAGCACAGTCACCATCAATTGC**CGGGCCAGTCAG**
AGTGTTTATTATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGAC
GTGCAGTGTGACGATGCTGCCACTTACTACTGT*CTAGGCAGTTATGATTGTAGTAATGGTGATTGTTTTGTT*TTCGGCGGAG
GGACCGAGGTGGTGGTCAAACGT (SEQ ID NO: 241)

Ab11 Light chain (chimera) Full length DNA sequence.

CAGGTGCTGACCCAGACTGCATCCCCCGTGTCTCCAGCTGTGGGAAGCACAGTCACCATCAATTGCCGGGCCAGTCAGA
GTGTTTATTATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTATTCTACATC
CACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAGTGGATCTGGGACACAGTTCACTCTCACCATCAGCGACGTG
CAGTGTGACGATGCTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTAATGGTGATTGTTTTGTTTTCGGCGGAGG
GACCGAGGTGGTGGTCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 242)

Figure 12A

Ab12

Ab12 Heavy chain (humanized) Full length protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIDVTNYYMQWVRQAPGKGLEWVGVIGVNGKRYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV
HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL
MISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPA
PIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVD
KSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 114)

Ab12 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIDVTNYYMQWVRQAPGKGLEWVGVIGVNGKRYYASWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCARGDIWGQGTLVTVSS (SEQ ID NO: 113)

Ab12 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGIDVT**NYYMQ**WVRQAPGKGLEWVGVIGVNGKRYYASWAKGRFTISRDNSKTTVY
LQMNSLRAEDTAVYFCAR*GDI*WGQGTLVTVSS (SEQ ID NOS: 118, 119, 120, respectively)

Ab12 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACGTCACT**AACTACTACATGCAA**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTG
TGAATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCCAGA*GGGGACATC*TGGGGCCAAGGGAC
CCTCGTCACCGTCTCGAGC (SEQ ID NO: 253)

Figure 12B

Ab12 Heavy chain (humanized) Full length DNA sequence.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGACGTCACTAACTACTACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTG
TGAATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCCAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 254)

Ab12 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCRASQSVYYNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPEDV
ATYYCLGSYDCSNGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQ
ESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 112)

Ab12 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCRASQSVYYNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPEDV
ATYYCLGSYDCSNGDCFVFGGGTKVEIKR (SEQ ID NO: 111)

Figure 12C

Ab12 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**RASQSVYYNNYL**AWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED VATYYC*LGSYDCSNGDCFV*FGGGTKVEIKR (SEQ ID NOS: 115, 116, 117, respectively)

Ab12 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CGGGCCAGTCAG AGTGTTTACTATAACAACTACCTAGCC**TGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTAC ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTGGGCAGTTATGATTGTAGTAATGGTGATTGTTTTGTT*TTCGGCGGAG GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 251)

Ab12 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCGGGCCAGTCAGA GTGTTTACTATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTACATC CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG CAGCCTGAAGATGTTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTAATGGTGATTGTTTTGTTTTCGGCGGAGG AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 252)

Figure 13A

Ab13

Ab13 Heavy chain (chimera) Full length protein sequence.

QSVEESGGGLVQPEGSLTLTCTASGFDFSSNAMWWVRQAPGKGLEWIGCIYNGDGSTYYASWVNGRFSISKTSSTTVTLQL
NSLTVADTATYYCARDLDLWGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHT
FPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMI
SRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIE
KTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSR
WQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 124)

Ab13 Variable region heavy chain (chimera) protein sequence.

QSVEESGGGLVQPEGSLTLTCTASGFDFSSNAMWWVRQAPGKGLEWIGCIYNGDGSTYYASWVNGRFSISKTSSTTVTLQL
NSLTVADTATYYCARDLDLWGPGTLVTVSS (SEQ ID NO: 123)

Ab13 Variable region heavy chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QSVEESGGGLVQPEGSLTLTCTASGFDFS**SNAMW**WVRQAPGKGLEWIGCIYNGDGSTYYASWVNGRFSISKTSSTTVTLQL
NSLTVADTATYYCAR*DLDL*WGPGTLVTVSS (SEQ ID NOS: 128, 129, 130, respectively)

Ab13 Variable region heavy chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

CAGTCGGTGGAGGAGTCCGGGGGAGGCCTGGTCCAGCCTGAGGGATCCCTGACACTCACCTGCACAGCCTCTGGATTC
GACTTCAGT**AGCAATGCAATGTGG**TGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGATGCATTTACAA
TGGTGATGGCAGCACATACTACGCGAGCTGGGTGAATGGCCGATTCTCCATCTCCAAAACCTCGTCGACCACGGTGACT
CTGCAACTGAATAGTCTGACAGTCGCGGACACGGCCACGTATTATTGTGCGAGA*GATCTTGACTTG*TGGGGCCCGGGCA
CCCTCGTCACCGTCTCGAGC (SEQ ID NO: 263)

Figure 13B

Ab13 Heavy chain (chimera) Full length DNA sequence.

CAGTCGGTGGAGGAGTCCGGGGGAGGCCTGGTCCAGCCTGAGGGATCCCTGACACTCACCTGCACAGCCTCTGGATTC
GACTTCAGTAGCAATGCAATGTGGTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGATGCATTTACAAT
GGTGATGGCAGCACATACTACGCGAGCTGGGTGAATGGCCGATTCTCCATCTCCAAAACCTCGTCGACCACGGTGACTC
TGCAACTGAATAGTCTGACAGTCGCGGACACGGCCACGTATTATTGTGCGAGAGATCTTGACTTGTGGGGCCCGGGCAC
CCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGG
GGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCTG
ACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT
CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACAAGAGAGTTG
AGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCCT
CTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCAC
GAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGA
GCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAA
GTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAACC
ACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTT
CTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGCT
GGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCA
TGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 264)

Ab13 Light chain (chimera) Full length protein sequence.

AIVMTQTPSSKSVPVGDTVTINCQASESLYNNNALAWFQQKPGQPPKRLIYDASKLASGVPSRFSGGGSGTQFTLTISGVQCD
DAATYYCGGYRSDSVDGVAFAGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGN
SQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 122)

Ab13 Variable region light chain (chimera) protein sequence.

AIVMTQTPSSKSVPVGDTVTINCQASESLYNNNALAWFQQKPGQPPKRLIYDASKLASGVPSRFSGGGSGTQFTLTISGVQCD
DAATYYCGGYRSDSVDGVAFAGGTEVVVKR (SEQ ID NO: 121)

Figure 13C

Ab13 Variable region light chain (chimera) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

AIVMTQTPSSKSVPVGDTVTINC**QASESLYNNNALA**WFQQKPGQPPKRLIY<u>DASKLAS</u>GVPSRFSGGGSGTQFTLTISGVQCD
DAATYYC*GGYRSDSVDGVA*FAGGTEVVVKR (SEQ ID NOS: 125, 126, 127, respectively)

Ab13 Variable region light chain (chimera) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GCCATCGTGATGACCCAGACTCCATCTTCCAAGTCTGTCCCTGTGGGAGACACAGTCACCATCAATTGC**CAGGCCAGT**
GAGAGTCTTTATAATAACAACGCCTTGGCCTGGTTTCAGCAGAAACCAGGGCAGCCTCCCAAGCGCCTGATCTATG<u>A</u>
<u>TGCATCCAAACTGGCATCT</u>GGGGTCCCATCGCGGTTCAGTGGCGGTGGGTCTGGGACACAGTTCACTCTCACCATCAGT
GGCGTGCAGTGTGACGATGCTGCCACTTACTACTGT*GGAGGCTACAGAAGTGATAGTGTTGATGGTGTTGCT*TTCGCCGGA
GGGACCGAGGTGGTGGTCAAACGT (SEQ ID NO: 261)

Ab13 Light chain (chimera) Full length DNA sequence.

GCCATCGTGATGACCCAGACTCCATCTTCCAAGTCTGTCCCTGTGGGAGACACAGTCACCATCAATTGCCAGGCCAGTG
AGAGTCTTTATAATAACAACGCCTTGGCCTGGTTTCAGCAGAAACCAGGGCAGCCTCCCAAGCGCCTGATCTATGATGC
ATCCAAACTGGCATCTGGGGTCCCATCGCGGTTCAGTGGCGGTGGGTCTGGGACACAGTTCACTCTCACCATCAGTGGC
GTGCAGTGTGACGATGCTGCCACTTACTACTGTGGAGGCTACAGAAGTGATAGTGTTGATGGTGTTGCTTTCGCCGGAG
GGACCGAGGTGGTGGTCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATC
TGGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCC
CTCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTG
ACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTC
ACAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 262)

Figure 14A

Ab14

Ab14 Heavy chain (humanized) Full length protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGVIGSDGKTYYATWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCTRGDIWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVH
TFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDARVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLM
ISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPI
EKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKS
RWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK (SEQ ID NO: 134)

Ab14 Variable region heavy chain (humanized) protein sequence.

EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGVIGSDGKTYYATWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCTRGDIWGQGTLVTVSS (SEQ ID NO: 133)

Ab14 Variable region heavy chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

EVQLVESGGGLVQPGGSLRLSCAVSGIGL**SSYYMQ**WVRQAPGKGLEWVGVIGSDGKTYYATWAKGRFTISRDNSKTTVYL
QMNSLRAEDTAVYFCTR*GDI*WGQGTLVTVSS (SEQ ID NOS: 138, 139, 140, respectively)

Ab14 Variable region heavy chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGGCCTCAGT**AGCTACTACATGCAA**TGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
GTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGA*GGGGACATC*TGGGGCCAAGGGAC
CCTCGTCACCGTCTCGAGC (SEQ ID NO: 273)

Figure 14B

Ab14 Heavy chain (humanized) Full length DNA sequence.

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCTGAGACTCTCCTGTGCAGTCTCTGGAA
TCGGCCTCAGTAGCTACTACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTCATTGGTA
GTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGT
ATCTTCAAATGAACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGAGGGGACATCTGGGGCCAAGGGA
CCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGG
GGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCCCT
GACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCC
TCCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACCAAGGTGGACGCGAGAGTT
GAGCCCAAATCTTGTGACAAAACTCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCA
CGAAGACCCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGG
AGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGCAGCCCCGAGAAC
CACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCT
TCTATCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACAAGACCACGCCTCCCGTGC
TGGACTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC
ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO: 274)

Ab14 Light chain (humanized) Full length protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSRGDCFVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNS
QESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO: 132)

Ab14 Variable region Light chain (humanized) protein sequence.

QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIYSTSTLASGVPSRFSGSGSGTDFTLTISSLQPED
VATYYCLGSYDCSRGDCFVFGGGTKVEIKR (SEQ ID NO: 131)

Figure 14C

Ab14 Variable region Light chain (humanized) protein sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

QVLTQSPSSLSASVGDRVTINC**QASQNVYNNNYLA**WYQQKPGKVPKQLIY**STSTLAS**GVPSRFSGSGSGTDFTLTISSLQPED
VATYYC*LGSYDCSRGDCFV*FGGGTKVEIKR (SEQ ID NOS: 135, 136, 137, respectively)

Ab14 Variable region Light chain (humanized) DNA sequence. CDR1: Bold; CDR2: Underlined; CDR3: Italics.

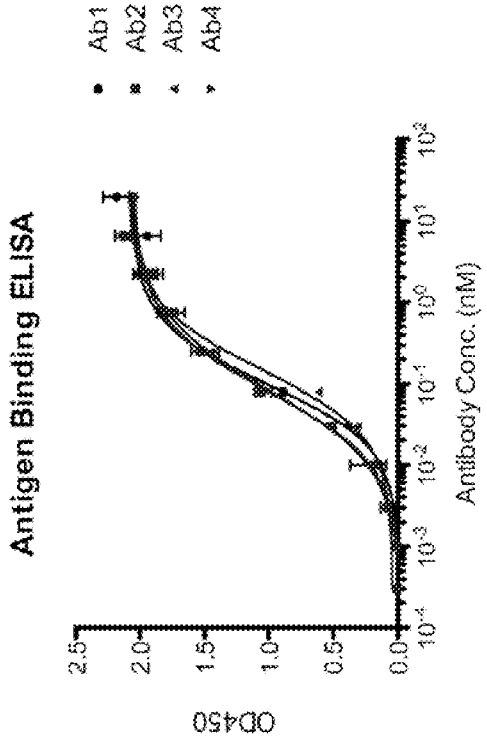
CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGC**CAGGCCAGTCAG**
AATGTTTACAATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTAC
ATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGC
CTGCAGCCTGAAGATGTTGCAACTTATTACTGT*CTGGGCAGTTATGATTGTAGTCGTGGTGATTGTTTTGTTT*TCGGCGGAG
GAACCAAGGTGGAAATCAAACGT (SEQ ID NO: 271)

Ab14 Light chain (humanized) Full length DNA sequence.

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCAATTGCCAGGCCAGTCAGA
ATGTTTACAATAACAACTACCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTATTCTACATC
CACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTG
CAGCCTGAAGATGTTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTCGTGGTGATTGTTTTGTTTTCGGCGGAGG
AACCAAGGTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCC
TCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGA
CGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCA
CAAAGAGCTTCAACAGGGGAGAGTGTTAG (SEQ ID NO: 272)

Figure 15

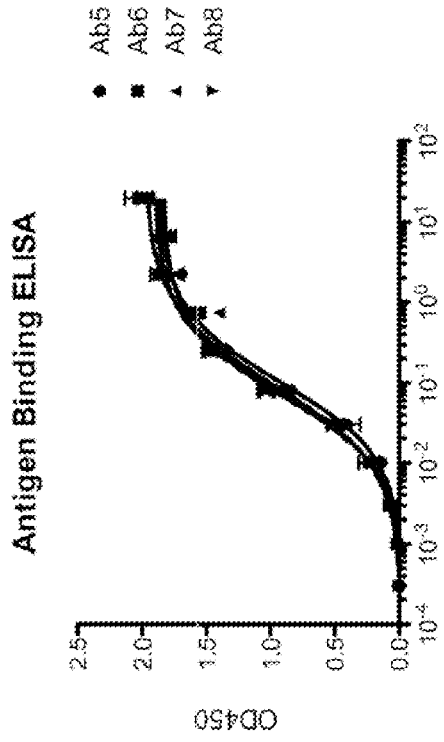
Human CGRPα ELISA



	EC50 (pM)
Ab1	103
Ab2	83
Ab3	154
Ab4	88

Figure 16

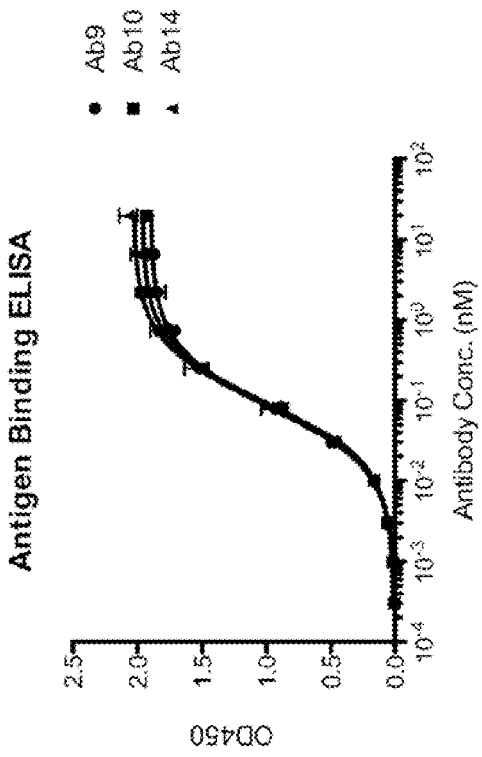
Human CGRPα ELISA



	EC50 (pM)
Ab5	103
Ab6	95
Ab7	70
Ab8	74

Figure 17

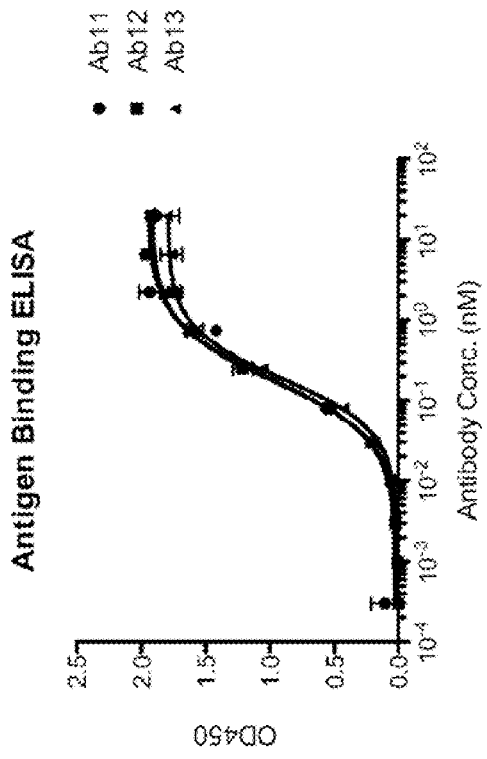
Human CGRPα ELISA



	EC50 (pM)
Ab9	79
Ab10	92
Ab14	89

Figure 18

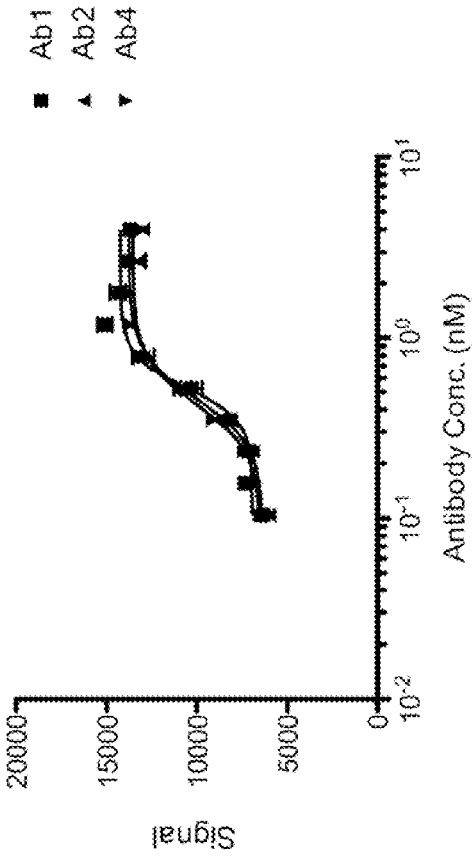
Human CGRPα ELISA



	EC50 (pM)
Ab11	184
Ab12	171
Ab13	188

Figure 19

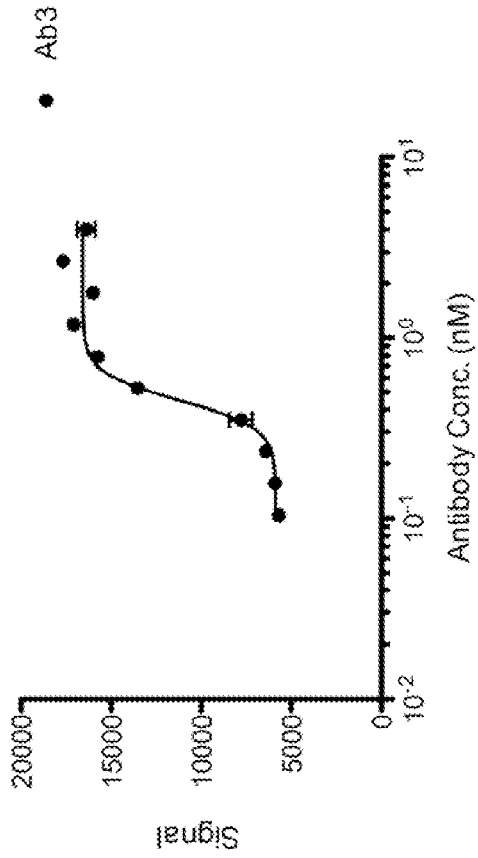
Human CGRPα cAMP



	IC50 (pM)
Ab1	531
Ab2	452
Ab4	429

Figure 20

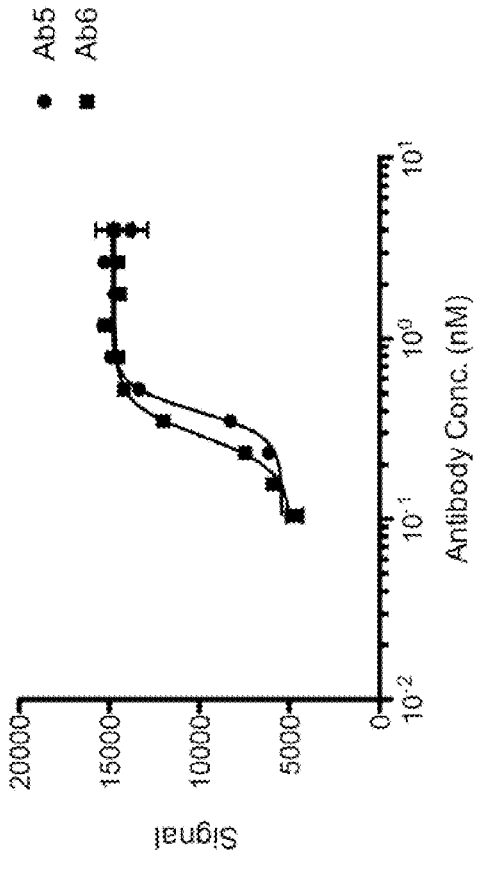
Human CGRPα cAMP



	IC50 (pM)
Ab3	452

Figure 21

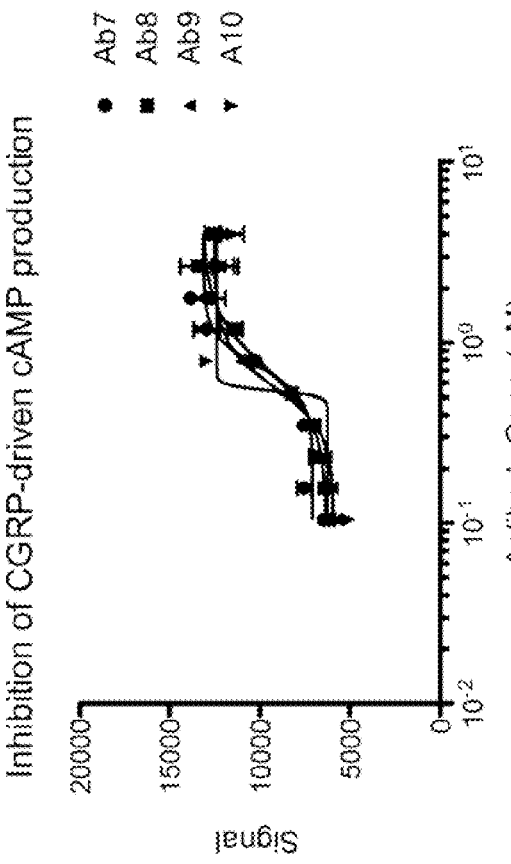
Human CGRPα cAMP



	IC50 (pM)
Ab5	400
Ab6	288

Figure 22

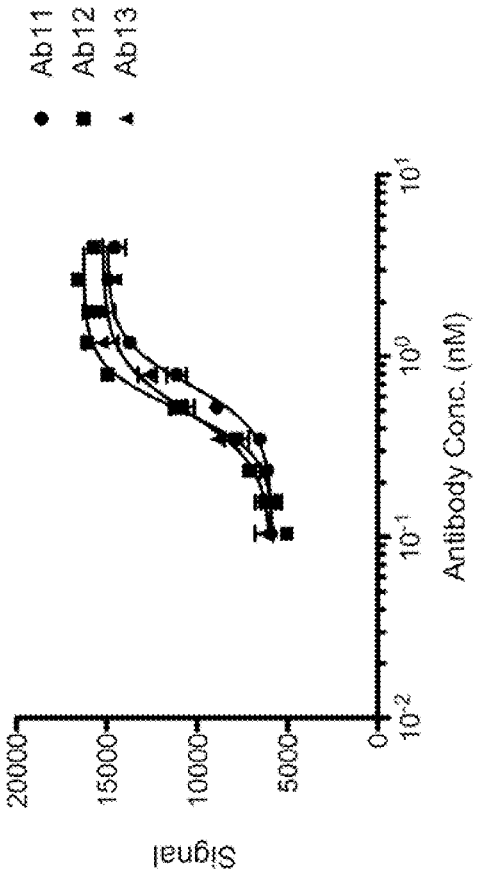
Human CGRPα cAMP



	IC50 (pM)
Ab7	743
Ab8	734
Ab9	568
A10	542

Figure 23

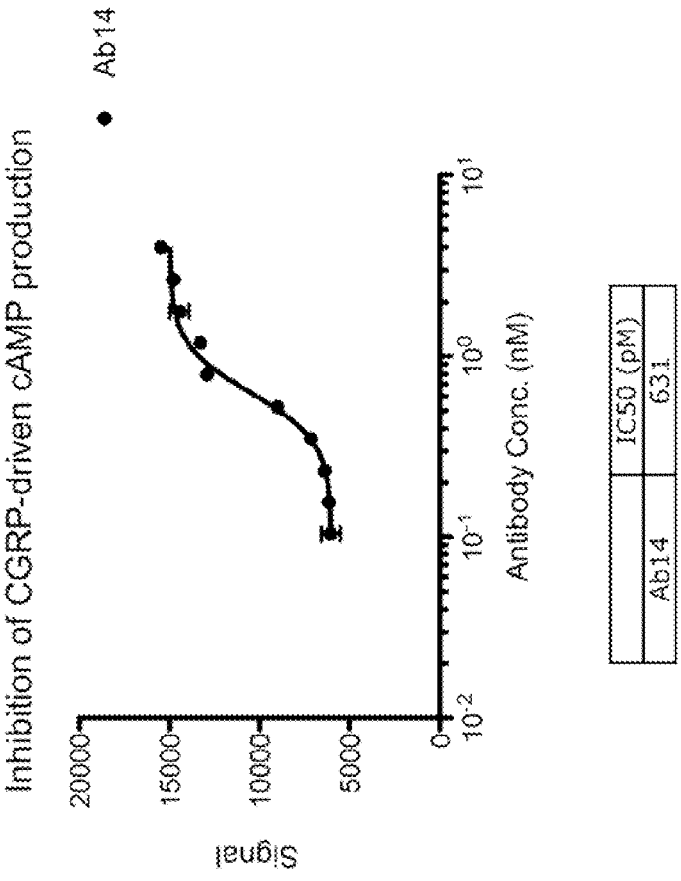
Human CGRPα cAMP



	IC50 (pM)
Ab11	698
Ab12	511
Ab13	498

Figure 24

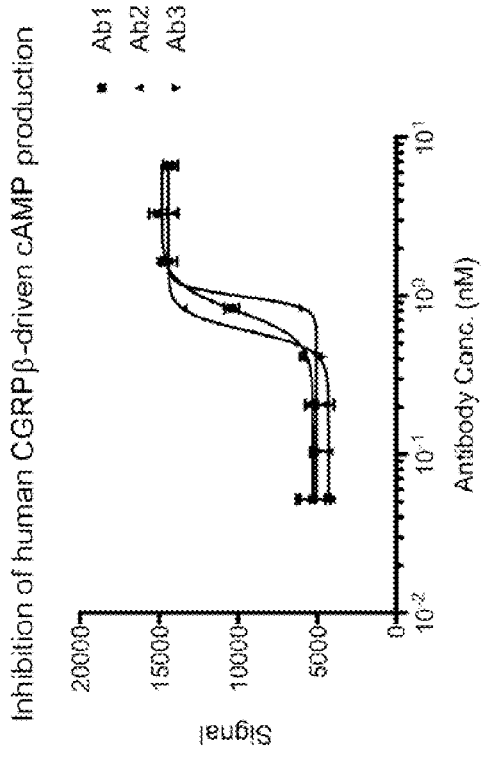
Human CGRPα cAMP



	IC50 (pM)
Ab14	631

Figure 25

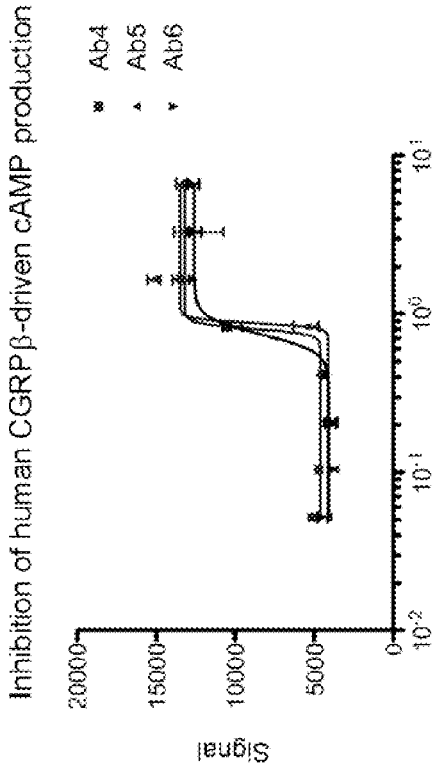
Human CGRPβ cAMP



	IC50 (pM)
Ab1	801
Ab2	601
Ab3	989

Figure 26

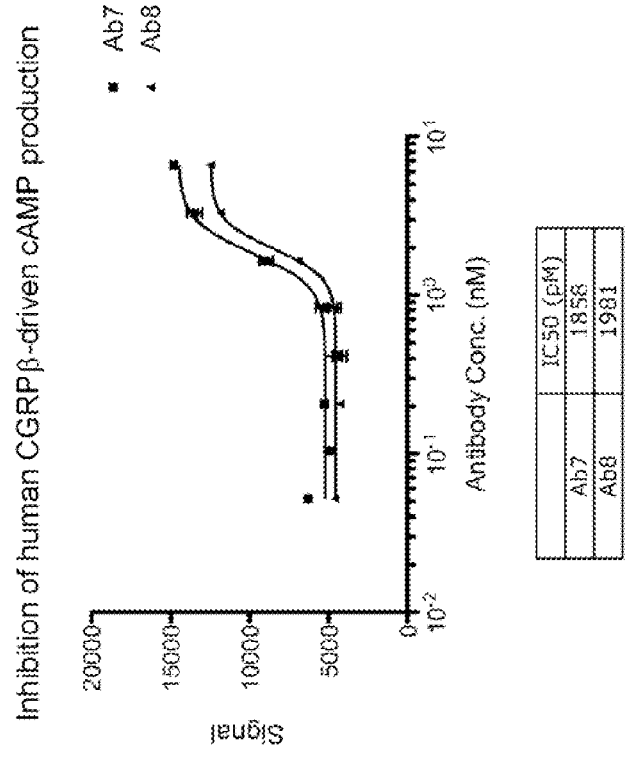
Human CGRPβ cAMP



	IC50 (pM)
Ab4	805
Ab5	875
Ab6	740

Figure 27

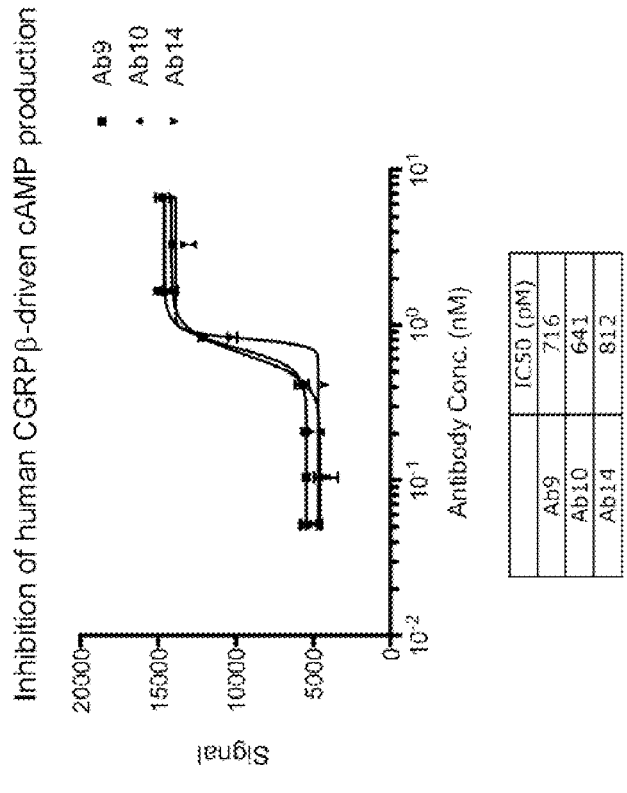
Human CGRPβ cAMP



	IC50 (pM)
Ab7	1858
Ab8	1981

Figure 28

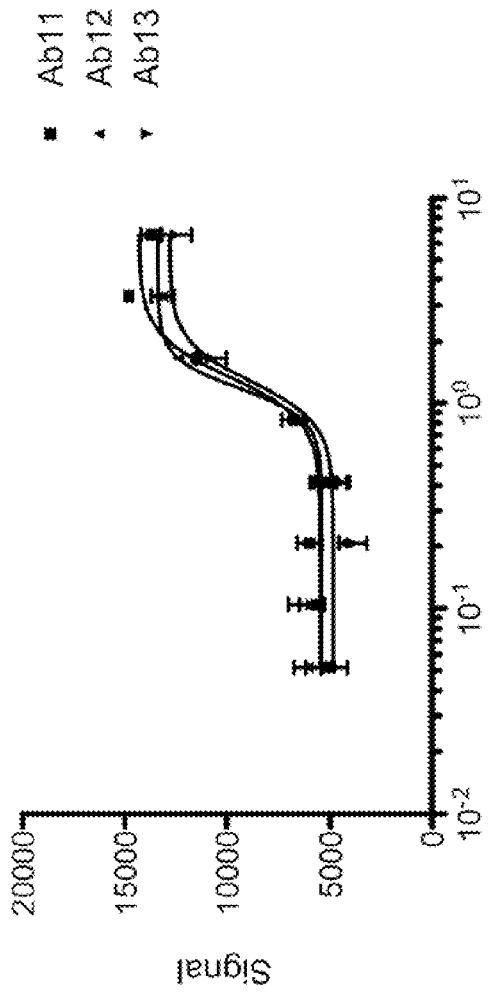
Human CGRPβ cAMP



	IC50 (pM)
Ab9	716
Ab10	641
Ab14	812

Figure 29

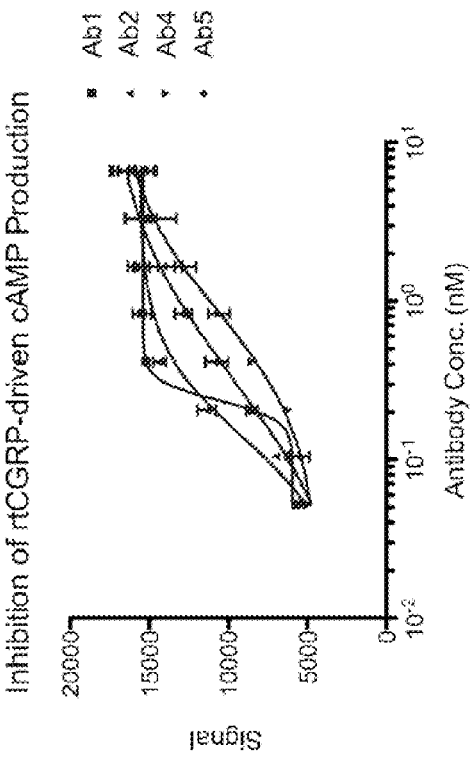
Human CGRPβ cAMP



	IC50 (pM)
Ab11	1344
Ab12	1181
Ab13	1276

Figure 30

Rat CGRP cAMP



	IC50 (pM)
Ab1	239
Ab2	142
Ab4	868
Ab5	334

Figure 31

Rat CGRP cAMP

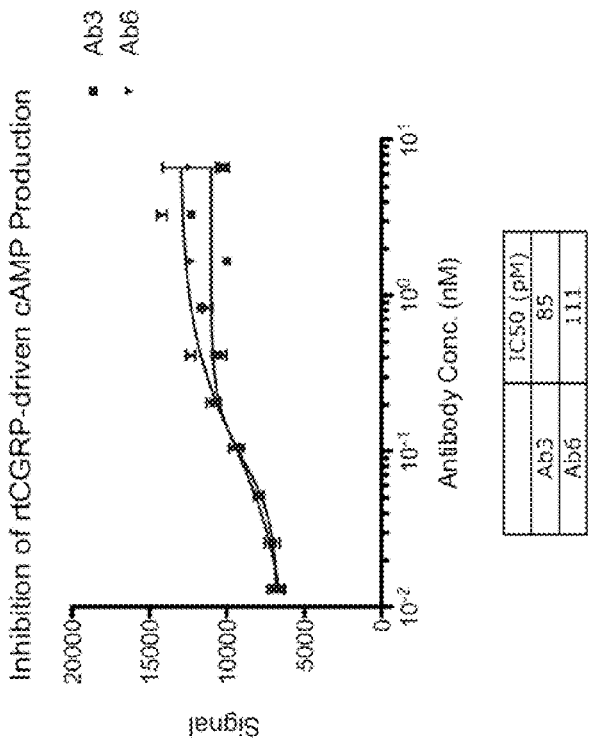
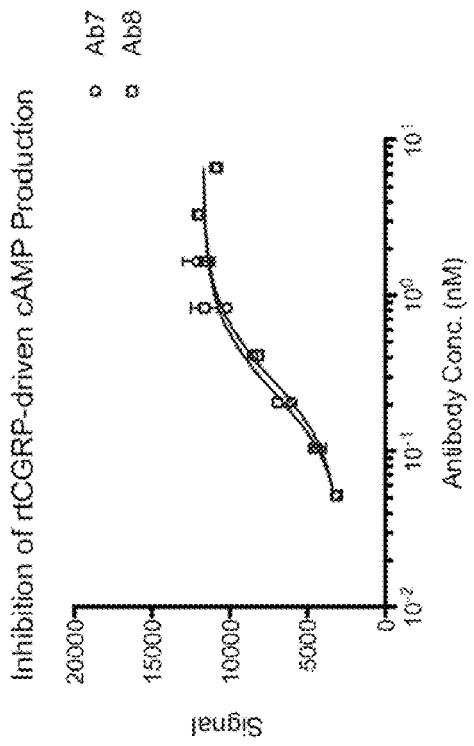


Figure 32

Rat CGRP cAMP



	IC50 (pM)
Ab7	297
Ab8	243

Figure 33

Rat CGRP cAMP

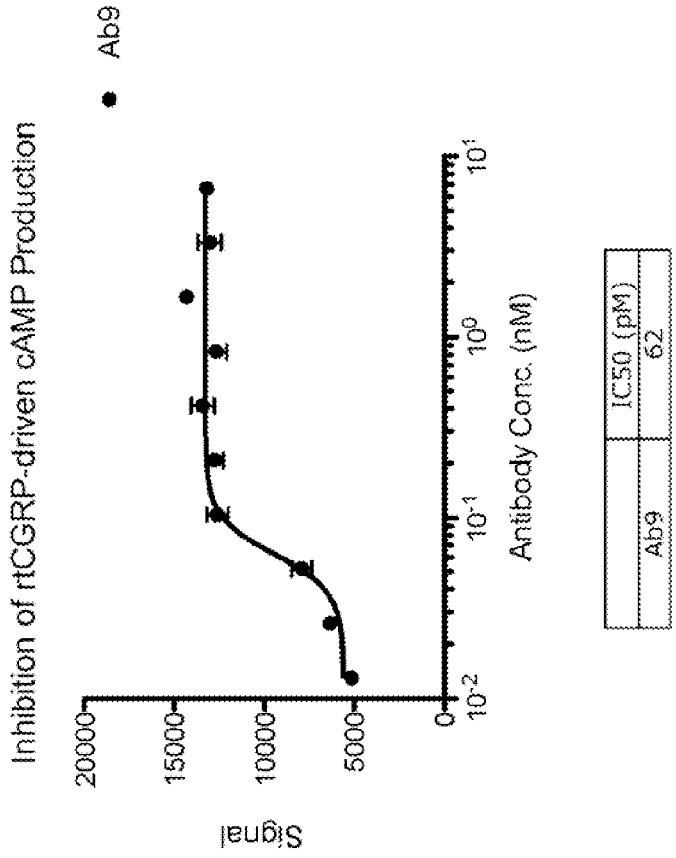


Figure 34

Rat CGRP cAMP

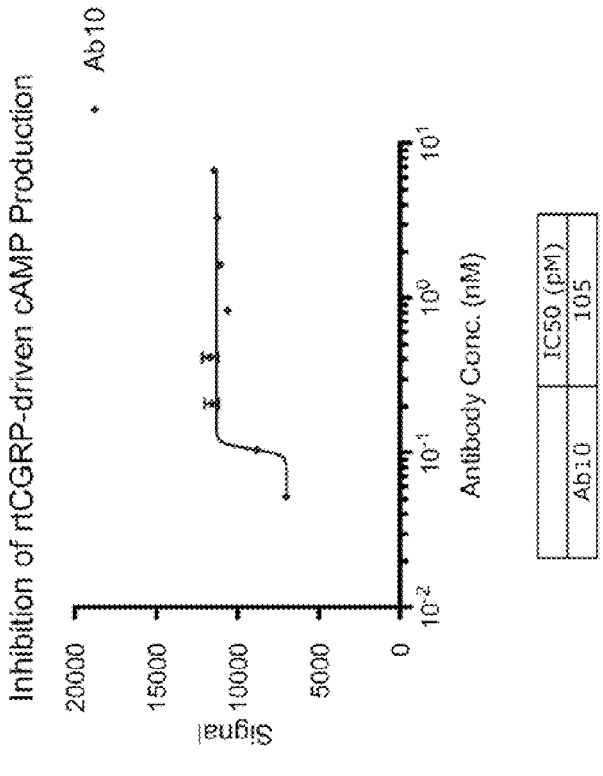


Figure 35

Rat CGRP cAMP

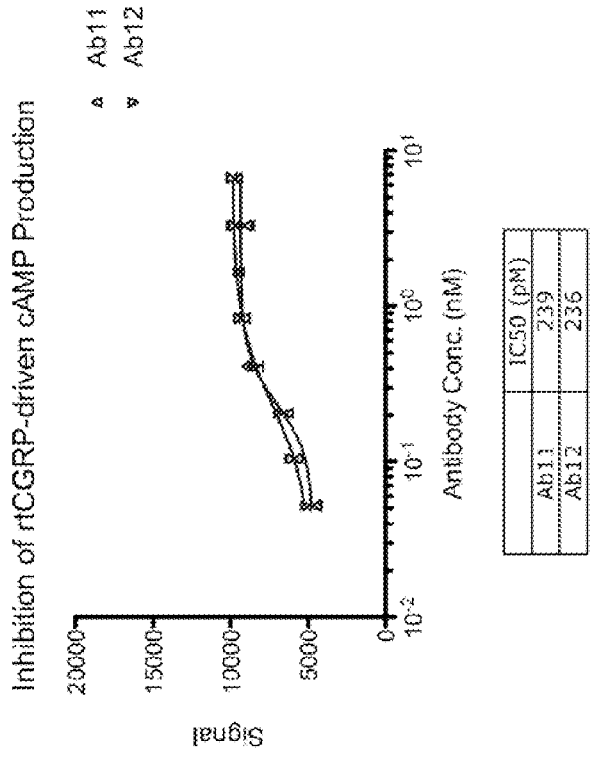


Figure 36

Rat CGRP cAMP

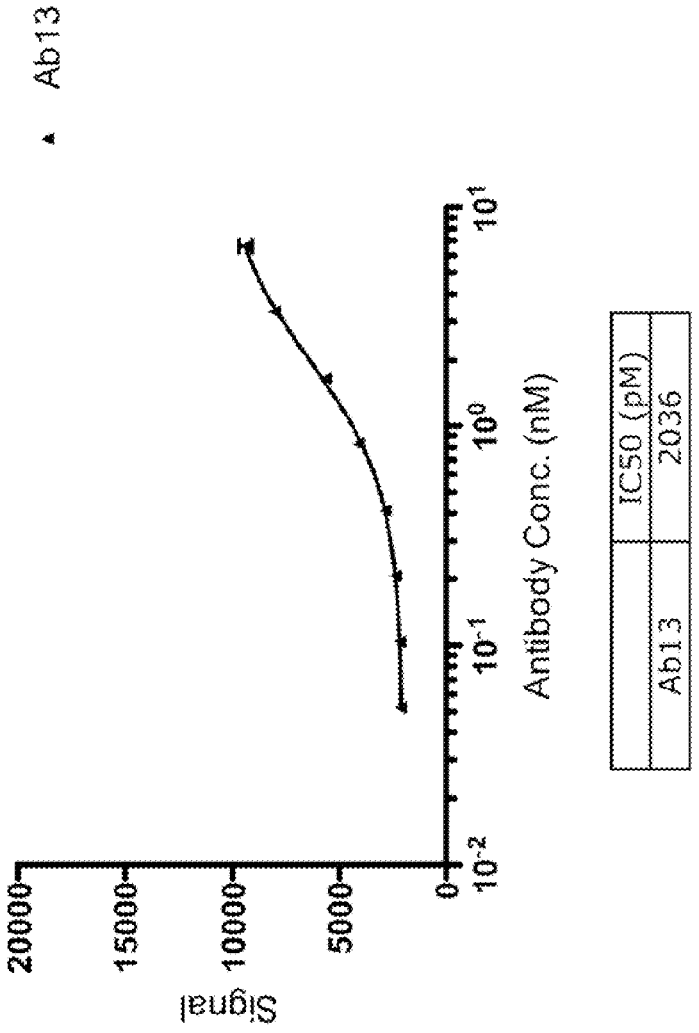


Figure 37

Rat CGRP cAMP

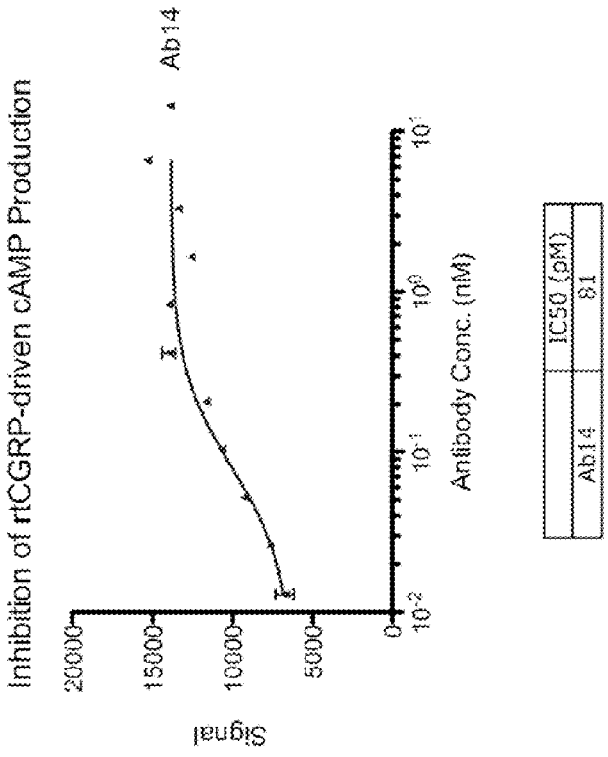


Figure 38

Inhibition of Radioligand Binding

	IC_{50} (nM)	K_I (nM)
Ab1	0.585	0.46
Ab2	0.482	0.378
Ab3	2.49	10.96
Ab4	0.579	0.455
Ab5	0.586	0.461
Ab6	2.46	1.94
Ab7	4.53	3.56
Ab8	0.936	0.736
Ab9	2.03	1.6
Ab10	0.28	0.22
Ab11	2.26	1.78
Ab12	0.315	0.248
Ab13	0.335	0.264

Figure 39

Reduction in Vasodilatation Following Capsaicin Administration

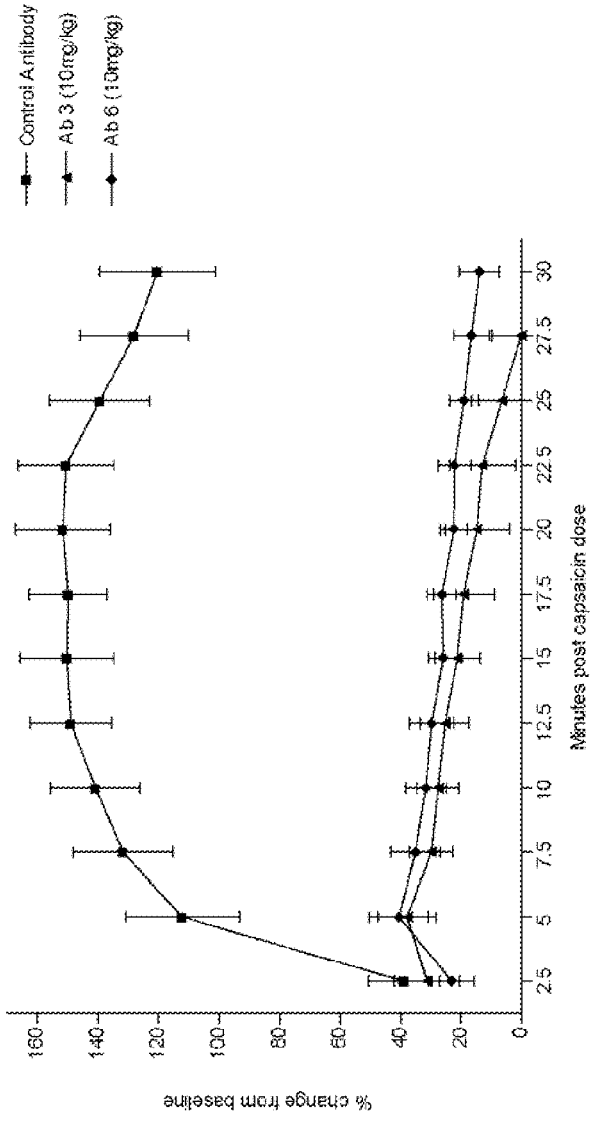
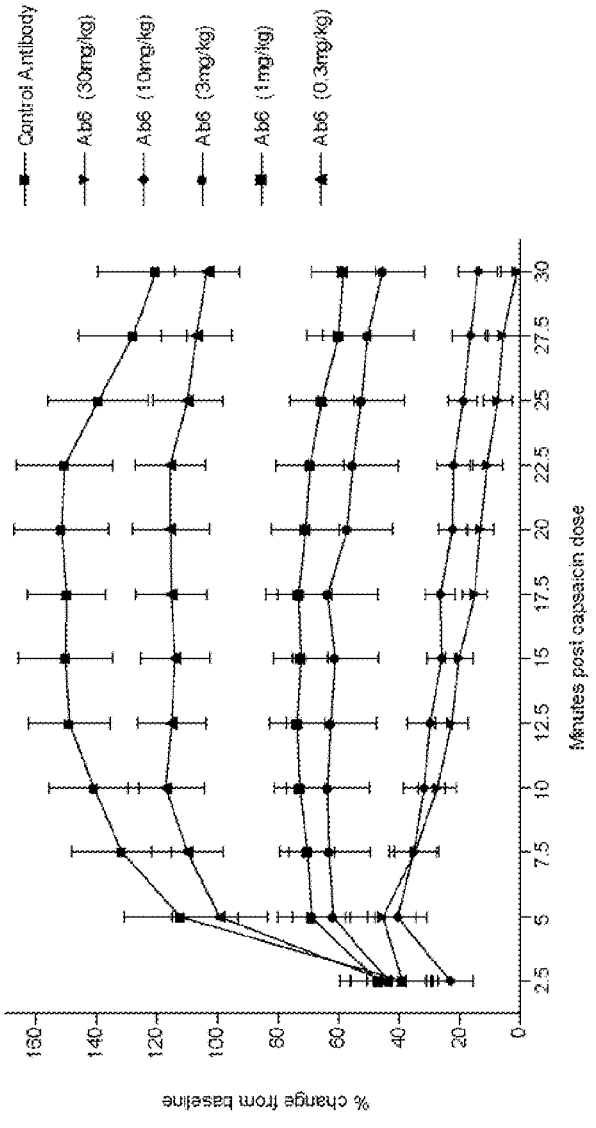


Figure 40

Reduction in Vasodilatation Following Capsaicin Administration



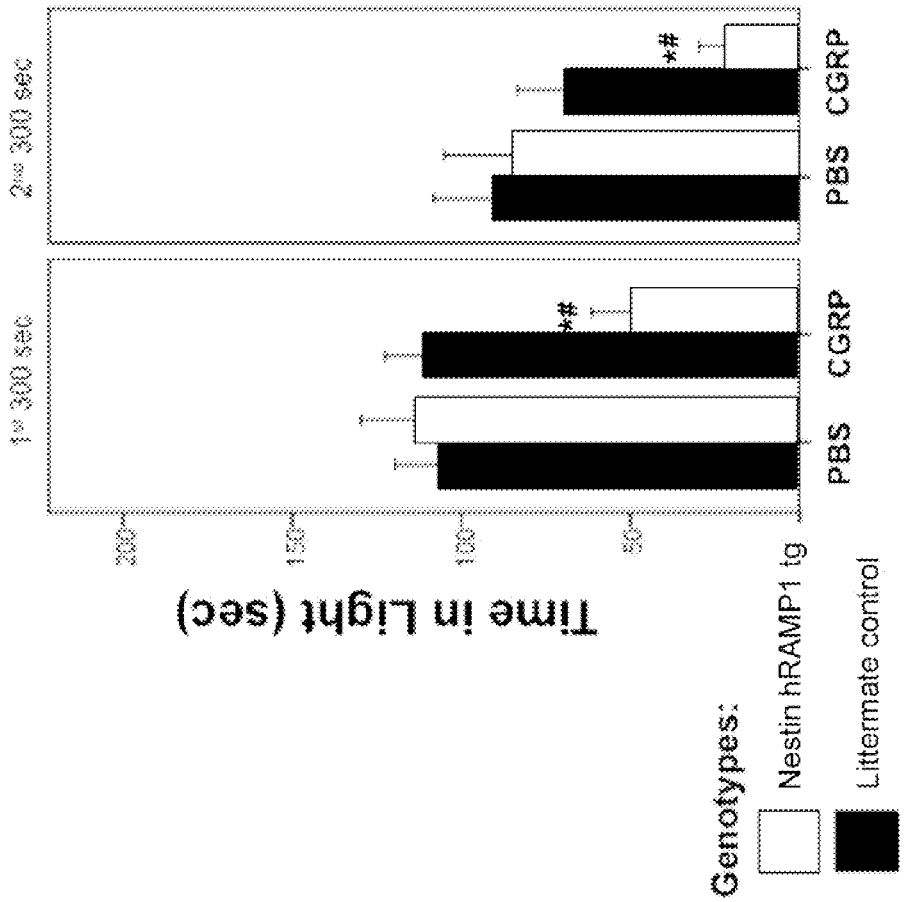
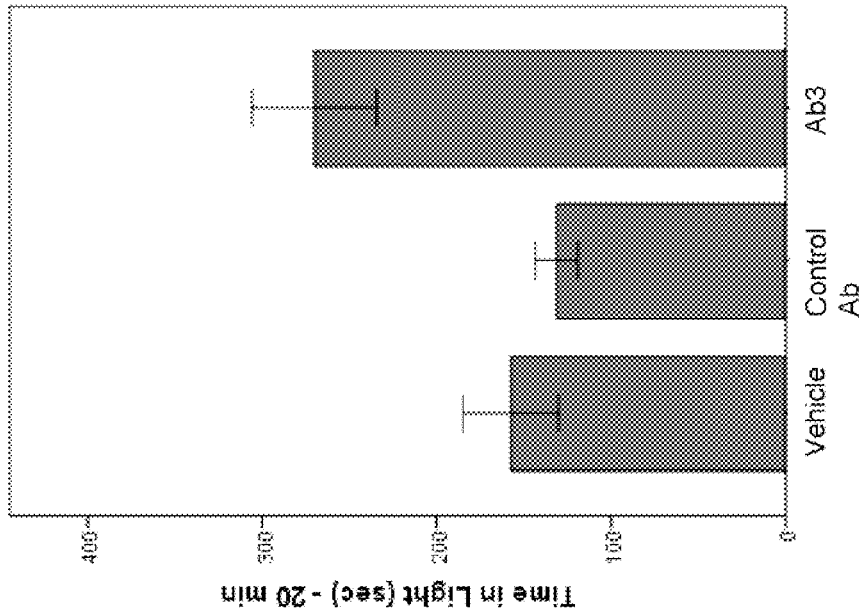

Time in Light (sec)
200
150
100
50
1st 300 sec
2nd 300 sec
PBS
CGRP
PBS
CGRP
*#
*#
Genotypes:
Nestin hRAMP1 tg
Littermate control

FIG. 41


Time in Light (sec) - 20 min
400
300
200
100
0
Vehicle
Control Ab
Ab3

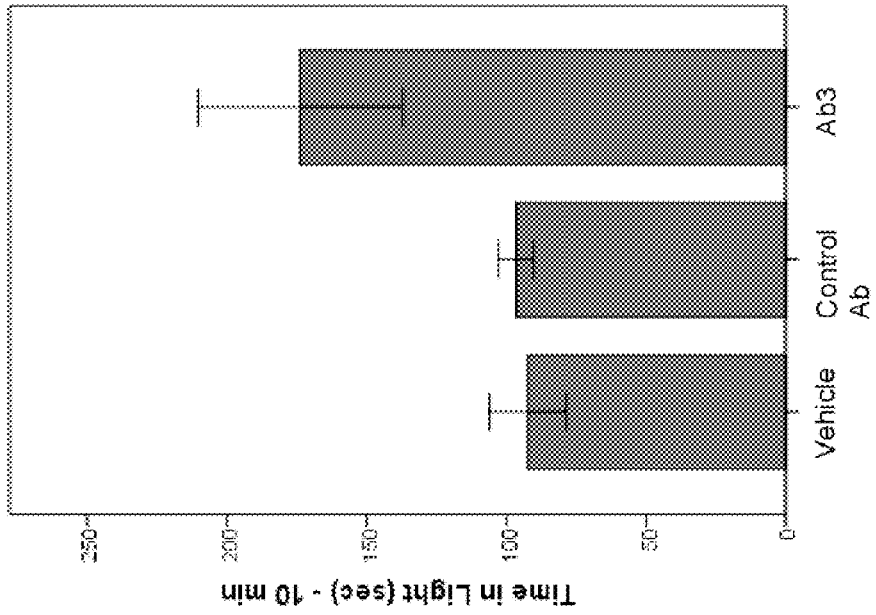

Time in Light (sec) - 10 min
250
200
150
100
50
0
Vehicle
Control Ab
Ab3

FIG. 42

USE OF ANTI-CGRP ANTIBODIES AND ANTIBODY FRAGMENTS TO PREVENT OR INHIBIT PHOTOPHOBIA OR LIGHT A VERSION IN SUBJECTS IN NEED THEREOF, ESPECIALLY MIGRAINE SUFFERERS

CROSS-REFERENCE TO RELATED APPLICATIONS

This application is a Divisional of U.S. application Ser. No. 15/621,105 filed Jun. 13, 2017, which is a Divisional of U.S. application Ser. No. 13/476,632, filed May 21, 2012 (now U.S. Pat. No. 9,708,393), which claims benefit of U.S. Provisional Appl. No. 61/496,860, filed Jun. 14, 2011, and U.S. Provisional Appl. No. 61/488,600, filed May 20, 2011, each of which is hereby incorporated by reference in its entirety.

This application includes as part of its disclosure a biological sequence listing which is being concurrently submitted through EFS-Web. Said biological sequence listing is contained in a file named "4325702604.txt" which was created on Apr. 18, 2019, and has a size of 204,115 bytes, and is hereby incorporated by reference in its entirety.

BACKGROUND OF THE INVENTION

Field of the Invention

This invention pertains to the discovery that polypeptides that inhibit the CGRP/CGRP receptor interaction and/or antibodies and antibody fragments that specifically bind CGRP or to a CGRP receptor may be used to inhibit CGRP-induced photophobia when administered to a subject in need thereof. Polypeptides that inhibit the CGRP/CGRP receptor interaction for use in the invention include by way of example antibodies and antibody fragments specific to CGRP or the CGRP receptor and fragments or variants of CGRP or the CGRP receptor that inhibit CGRP from interacting with CGRP receptors. As photophobia is an adverse side-effect often associated with many disorders including by way of example migraine with and without aura and other headache conditions (as well as other indications disclosed infra) t these CGRP-receptor inhibitors, e.g., antibodies and antibody fragments specific to CGRP or the CGRP receptor should be well suited for inhibiting the photophobia often associated with migraine and other headache conditions as well as for treating other conditions associated with photophobia. The results also suggest that these antibodies and antibody fragments may be used to prevent the onset of photophobia in subjects in need thereof such as individuals with a chronic history of photophobia, e.g., as a result of migraine (with or without aura), other headache condition, depression, agoraphobia or other conditions prone to photophobia if the antibodies are administered prophylactically. The invention contemplates the use of these anti-CGRP antibodies and antibody fragments as a monotherapy or in therapeutic regimens with other active agents, e.g., analgesics, opioids, antidepressants or other actives dependent on the condition and the individual treated.

The invention further provides methods of screening CGRP-receptor inhibitors, e.g., anti-CGRP or anti-CGRP receptor antibodies and fragments thereof (including Fab fragments) having binding specificity to human Calcitonin Gene Related Peptide (hereinafter "CGRP") or the CGRP receptor in specific animal models to determine the in vivo effects thereof, most especially their ability to antagonize the photophobic side effects of CGRP and to treat conditions involving photophobia including e.g., migraine.

Description of Related Art

Calcitonin Gene Related Peptide (CGRP) is produced as a multifunctional neuropeptide of 37 amino acids in length. Two forms of CGRP, the CGRP-alpha and CGRP-beta forms, exist in humans and have similar activities. CGRP-alpha and CGRP-beta differ by three amino acids in humans, and are derived from different genes. The CGRP family of peptides includes amylin, adrenomedullin, and calcitonin, although each has distinct receptors and biological activities. Doods, H., *Curr. Op. Invest. Drugs,* 2(9):1261-78 (2001).

CGRP is released from numerous tissues such as trigeminal nerves, which when activated release neuropeptides within the meninges, mediating neurogenic inflammation that is characterized by vasodilation, vessel leakage, and mast-cell degradation. Durham, P. L., New Eng. J. Med., 350 (11):1073-75 (2004). The biological effects of CGRP are mediated via the CGRP receptor (CGRP-R), which consists of a seven-transmembrane component, in conjunction with receptor-associated membrane protein (RAMP). CGRP-R further requires the activity of the receptor component protein (RCP), which is essential for an efficient coupling to adenylate cyclase through G proteins and the production of cAMP. Doods, H., Curr. Op. Invest. Drugs, 2(9): 1261-78 (2001).

Migraines constitute a neurovascular disorder affecting approximately 10% of the adult population in the U.S., and are typically accompanied by intense headaches. Approximately 20-30% of migraine sufferers experience aura, comprising focal neurological phenomena that precede and/or accompany the event. CGRP is believe to play a prominent role in the development of migraines. For example, plasma concentrations of CGRP were identified elevated in jugular venous blood during the headache phase of migraines, to the exclusion of other neuropeptides. Moreover, according to Arulmozhi et al, the following has been identified in migraine sufferers: (1) a strong correlation between plasma CGRP concentrations and migraines; (2) the infusion of CGRP produced a migraine-like headache; (3) baseline CGRP levels were elevated; and (4) changes in plasma CGRP levels during migraine attacks significantly correlated with headache intensity. Arulmozhi, D. K., et al., Vas. Pharma., 43: 176-187 (2005). In addition, in the Journal of the International Association for the Study of Pain PII: S0304-3959(11)00313-7; doi:10.1016/j.pain.2011.04.033, published online 6 Jun. 2011, Hou et al., reported that keratinocyte expression of calcitonin gene-related peptide β has implications for neuropathic and inflammatory pain mechanisms.

One effective treatment for migraines is the administration of triptans, which are a family of tryptamine-based drugs, including sumatriptan and rizatriptan. Members of this family have an affinity for multiple serotonin receptors, including 5-HT_{1B}, 5-HT_{1D}, and 5-HT_{1F}. Members of this family of drugs selectively constrict cerebral vessels, but also cause vasoconstrictive effects on coronary vessels. Durham, P. L., *New Eng. J. Med.,* 350 (11):1073-75 (2004). There is a theoretical risk of coronary spasm in patients with established heart disease following administration, and cardiac events after taking triptans may rarely occur. Noted to be contraindicated for patients with coronary vascular disease.

Similarly, pain may often be addressed through the administration of certain narcotics or non-steroidal anti-

inflammatory drugs (NSAIDs). However, the administration of these treatments may occur at the cost of certain negative consequences. NSAIDs have the potential to cause kidney failure, intestinal bleeding, and liver dysfunction. Narcotics have the potential to cause nausea, vomiting, impaired mental functioning, and addiction. Therefore, it is desirable to identify alternative treatments for pain in order to avoid certain of these negative consequences.

Aside from migraine, CGRP is believed to play a role in a multitude of diseases and disorders, including but not limited to other headache conditions, and pain. Due to the perceived involvement of CGRP in these diseases and disorders, there remains a need in the art for compositions and methods useful for preventing or treating diseases and disorders associated with CGRP, while avoiding adverse side effects. There in particular remains a need in the art for compositions or methods that reduce or inhibit photophobia in diseases or disorders associated with CGRP, such as migraines, headaches, and pain.

Migraineurs typically develop worsening pain and migraine symptoms when exposed to light, a phenomenon known as photophobia. Photophobia is also common in ocular disorders, such as iritis and uveitis, and intracranial disorders, such as meningitis. In the classic visual pathway, light activates rods and cones in the retina, which activate retinal ganglion cells that project via the optic nerve, to the lateral geniculate nucleus, superior colliculus, and then the visual cortex. This pathway includes image-forming and non-image-forming data. A new pathway (non-image-forming information) allows maintenance of normal circadian rhythms via the suprachiasmatic nucleus and is regulated by intrinsically photosensitive retinal ganglion cells (ipRGCs). These ipRGCs are independent of the rods and cones and contain melanopsin, a photopigment.

Noseda et al. (Noseda, R. et al. A neural mechanism for exacerbation of headache by light. Nat. Neurosci. 13, 239-245 (2010)) studied blind individuals who had migraine and correlated these findings with rat models involving tracing of ipRGC projections to areas in perception of pain from the dura. Of the blind patients with migraine, 6 had no light perception due to severe optic nerve damage or bilateral enucleation. These subjects experienced abnormal sleep patterns and poor pupillary light responses. Their migraines did not worsen with light exposure. In contrast, 14 blind subjects who were able to detect light despite minimal perception of images had normal sleep patterns and a normal pupillary light reflex. Despite widespread rod and cone degeneration, these patients had worsening migraine symptoms with light exposure during migraine attacks, suggesting that ipRGCs, and not rods and cones, are important in photophobia.

These retinal projections of non-image-forming brain areas project to the contralateral dorsocaudal region of the posterior thalamus, as demonstrated by anterograde tracing in the rat. ipRGC input to this area modulates dura-sensitive pain neurons, which also project to this region. Thalamic neurons, dually sensitive to dural pain and light input, project widely to multiple cortical regions, including the primary somatosensory cortex, the primary and secondary motor cortices, the parietal association cortex, and the primary and secondary visual cortices. These cortical projections may help explain other common migraine symptoms, in addition to photophobia, such as motor weakness or incoordination, visual disturbances, and poor concentration.

Photophobia also accompanies other less frequent but likewise disabling conditions, such as cluster headache and other trigeminal autonomic cephalalgias and blepharospasm. The mechanisms underlying photophobia involve the trigeminal system. Photophobia in blind patients suggests contributions from a nonvisual pathway. In addition, trigeminal autonomic cephalalgias, a less common group of primary headache disorders, are characterized by unilateral trigeminal-mediated pain frequently associated with ipsilateral photophobia.

Stimulation of trigeminal sensory neurons results in the release of neuropeptides (including substance P and calcitonin gene-related peptide, producing blood vessel dilation and mast cell, endothelial, and platelet activation (neurogenic inflammation), which leads to migraine. (Buzzi M G, Dimitriadou V, Theoharides T C, Moskowitz M A. 5-Hydroxytryptamine receptor agonists for the abortive treatment of vascular headaches block mast cell, endothelial and platelet activation within the rat dura mater after trigeminal stimulation. Brain Res 1992; 583:137-149). CGRP is elevated in external jugular venous blood during acute migraine pain, (Goadsby P J, Edvinsson L, Ekman R. Vasoactive peptide release in the extracerebral circulation of humans during migraine headache. Ann Neurol 1990; 28:183-187) and triptans reduce elevated CGRP levels. In animal models, mice sensitized to CGRP demonstrate more light-aversive behavior when exposed to exogenous CGRP. The administration of olcegepant, a CGRP receptor antagonist, prevented photophobia in these mice. (See Recober A, Kaiser E A, Kuburas A, Russo A F. Induction of multiple photophobic behaviors in a transgenic mouse sensitized to CGRP. Neuropharmacology 2010; 58:156-165).

However, while the use of anti-CGRP or anti-CGRP receptor antibodies and fragments to treat migraine has been suggested, to the best of Applicant's knowledge there has been no report of any polypeptide CGRP antagonist or in particular an anti-CGRP or anti-CGRP receptor antibody or antibody fragment able to alleviate or prevent the photophobic side effects of CGRP in vivo. The development of novel polypeptides that act as inhibitors of the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies or anti-CGRP or anti-CGRP receptor antibody fragments would be beneficial for patients who either do not respond to current migraine therapeutics such as triptans or who cannot take or tolerate them because of their potential vasoconstrictive effects.

BRIEF SUMMARY OF THE INVENTION

This invention relates to the discovery that polypeptides which inhibit the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies and anti-CGRP or anti-CGRP receptor antibody fragments (including Fab fragments) having binding specificity to human Calcitonin Gene Related Peptide (hereinafter "CGRP") as well as fragments of CGRP and the CGRP receptor that inhibit the CGRP/CGRP receptor interaction may be used to prevent or inhibit photophobia, especially CGR associated photophobia. Herein we particularly exemplify an anti-CGRP antibody identified as Ab3 infra, that very effectively alleviates or prevents photophobia, especially the photophobic effects of CGRP. Other preferred examples for use in the claimed therapies are Ab6 and Ab10 among others.

Based thereon the invention relates to the use of polypeptides which inhibit the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies and anti-CGRP or anti-CGRP receptor antibody fragments (including Fab fragments) having binding specificity to human Calcitonin Gene Related Peptide (hereinafter "CGRP") as well as fragments of CGRP and the CGRP receptor that

inhibit the CGRP/CGRP receptor interaction, preferably anti-CGRP antibodies and anti-CGRP antibody fragments for treating or preventing photophobia. The invention embraces the treatment or prevention of any photophobia, and in particular includes treatment or prevention of photophobia associated with migraine, and other disorders associated with photophobia such as cluster headache and other trigeminal autonomic cephalalgias and blepharospasm, depression, bipolar disorders, agoraphobia, meningitis, and photophobias associated with eye related conditions, autism, chronic fatigue syndrome, menstrual migraines, and other photopbia-associated conditions.

This invention also pertains to methods of screening polypeptides which inhibit the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies and anti-CGRP or anti-CGRP receptor binding antibody fragments (including Fab fragments) having binding specificity to human CGRP as well as fragments of CGRP and the CGRP receptor that inhibit the CGRP/CGRP receptor interaction, in specific photophobia animal models, e.g., the nestin/hRAMP1 rodent model disclosed infra, to determine the in vivo effects thereof, especially the ability of these polypeptides to inhibit the CGRP/CGRP receptor interaction in vivo and thereby antagonize the adverse in vivo side effects of CGRP including photophobia and to treat CGRP conditions involving CGRP associated photophobia including migraine and other disorders associated with photophobia such as cluster headache and other trigeminal autonomic cephalalgias and blepharospasm, depression, bipolar disorders, and other photophobia-associated conditions identified herein.

Also the invention specifically involves a method of assessing the potential in vivo efficacy of a candidate polypeptide which inhibit the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies and anti-CGRP or anti-CGRP receptor antibody fragments (including Fab fragments) having binding specificity to CGRP as well as fragments of CGRP and the CGRP receptor that inhibit the CGRP/CGRP receptor interaction, preferably an anti-CGRP or anti-CGRP receptor antibody or antibody fragment comprising determining whether the polypeptide, e.g., an antibody, inhibits light aversive behavior in a transgenic rodent which exhibits photoaversion when administered CGRP compared to the photoaversive behavior of the rodent administered CGRP in the absence of the candidate CGRP/CGRP receptor inhibitor polypeptide.

Also, the invention involves a method of assessing the potential in vivo efficacy of a candidate polypeptide which inhibit the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies and anti-CGRP or anti-CGRP receptor antibody fragments (including Fab fragments) as well as fragments of CGRP and the CGRP receptor that inhibit the CGRP/CGRP receptor interaction, preferably an anti-CGRP antibody or anti-CGRP receptor antibody or antibody fragment to treat a neurological condition or other condition characterized by increased CGRP levels that result in photophobia.

Further, the invention specifically involves a method of assessing the potential in vivo efficacy of a candidate polypeptide which inhibits the CGRP/CGRP receptor interaction such as anti-CGRP or anti-CGRP receptor antibodies and anti-CGRP or anti-CGRP receptor antibody fragments as well as fragments or variants of CGRP species and CGRP receptors that inhibit the CGRP/CGRP receptor interaction, preferably anti-CGRP or anti-CGRP receptor antibodies or antibody fragments to treat or prevent photophobia in migraine or chronic migraine, menstrual or menopausal or other hormonal associated migraines, cluster headaches or pain disorder associated with headache.

Still further, the invention involves a method of determining a suitable therapeutic dosage or dosage regimen of the candidate polypeptide CGRP/CGRP receptor inhibitor, e.g., anti-CGRP or anti-CGRP receptor antibody or antibody fragment in humans based on the effects of said polypeptide, e.g., an antibody or antibody fragment in a light aversive behavioral Nestin/hRAMP1 rodent animal model described in detail infra.

Further the invention relates to methods of assessing based on results in a rodent CGRP (Nestin/hRAMP1 animal model) a suitable therapeutic dosage or dosage regimen of the candidate polypeptide, e.g., an anti-CGRP or anti-CGRP receptor antibody or antibody fragment in humans.

In preferred embodiments the present invention is directed to therapeutic usage of specific antibodies and fragments thereof having binding specificity for CGRP, in particular antibodies having desired epitopic specificity, high affinity or avidity and/or functional properties. In preferred embodiments this invention relates to assays and usage of the antibodies described herein, comprising the sequences of the V_H, V_L and CDR polypeptides described herein, and the polynucleotides encoding them. A preferred embodiment of the invention is directed to chimeric or humanized antibodies and fragments thereof (including Fab fragments) capable of binding to CGRP or the CGRP receptor and/or inhibiting the biological activities mediated by the binding of CGRP to the CGRP receptor ("CGRP-R").

In another preferred embodiment of the invention, the assays and therapies use full length antibodies and Fab fragments thereof that inhibit the CGRP-alpha-, CGRP-beta-, and rat CGRP-driven production of cAMP. In a further preferred embodiment of the invention, full length and Fab fragments thereof are contemplated that reduce vasodilation and inhibit or prevent photophobia in a recipient following administration.

In another embodiment of the invention, chimeric or humanized antibodies and fragments thereof (including Fab fragments) capable of binding to CGRP or the CGRP receptor are useful in methods directed to reducing, treating, or preventing photophobia associated with one or more of the following conditions: migraines (with or without aura), cancer or tumors, angiogenesis associated with cancer or tumor growth, angiogenesis associated with cancer or tumor survival, weight loss, pain, hemiplagic migraines, cluster headaches, menstrual migranes, migrainous neuralgia, chronic headaches, tension headaches, general headaches, hot flashes, chronic paroxysomal hemicrania, secondary headaches due to an underlying structural problem in the head or neck, cranial neuralgia, sinus headaches (such as for example associated with sinusitis), headache-free migraine, abdominal migraine, and allergy-induced headaches or migraines.

Common causes of photophobia include migraine headaches, cataracts, or severe ophthalmologic diseases such as uveitis or corneal abrasion. A more extensive list of disorders associated with photophobia includes eye related causes such as Achromatopsia, Aniridia, Anticholinergic drugs may cause photophobia by paralyzing the iris sphincter muscle, Aphakia (absence of the lens of the eye), Buphthalmos (abnormally narrow angle between the cornea and iris), Cataracts, Cone dystrophy, Congenital abnormalities of the eye, Viral conjunctivitis ("pink eye") Corneal abrasion, Corneal dystrophy, Corneal ulcer, disruption of the corneal epithelium, such as that caused by a corneal foreign body or keratitis, Ectopia lentis, Endophthalmitis, Eye trauma

caused by disease, injury, or infection such as chalazion, episcleritis, glaucoma, keratoconus, or optic nerve hypoplasia, Hydrophthalmos, or congenital glaucoma Iritis, Optic neuritis, Pigment dispersion syndrom, Pupillary dilation (naturally or chemically induced), Retinal detachment, Scarring of the cornea or sclera and Uveitis.

In addition photophobia has nervous-system-related or urological causes including: Autism spectrum disorders, Chiari malformation, Dyslexia, Encephalitis including Myalgic encephalomyelitis aka Chronic fatigue syndrome, Meningitis, Subarachnoid haemorrhage, Tumor of the posterior cranial fossa, as well as other causes such as Ankylosing spondylitis, Albinism, Ariboflavinosis, Benzodiazepines (long term use of or withdrawal from benzodiazepines), Chemotherapy, Chikungunya, Cystinosis, Ehlers-Danlos syndrome, Hangover, Influenza, Infectious Mononucleosis, Magnesium deficiency, Mercury poisoning, Migraine, Rabies, and Tyrosinemia type II, also known as "Richner-Hanhart syndrome". Additionally it is known that photophobia is elevated in depression, bipolar disorder and agoraphobia.

In another embodiment of the invention these antibodies and humanized versions for treatment or prevention of photophobia may be derived from rabbit immune cells (B lymphocytes) and may be selected based on their homology (sequence identity) to human germ line sequences. These antibodies may require minimal or no sequence modifications, thereby facilitating retention of functional properties after humanization. A further embodiment of the invention is directed to fragments from anti-CGRP or anti-CGRP receptor antibodies encompassing V_H, V_L and CDR polypeptides, e.g., derived from rabbit immune cells and the polynucleotides encoding the same, as well as the use of these antibody fragments and the polynucleotides encoding them in the creation of novel antibodies and polypeptide compositions capable of binding to CGRP and/or CGRP/CGRP-R complexes.

The invention also contemplates conjugates of anti-CGRP or anti-CGRP receptor antibodies and binding fragments thereof for treatment or prevention of photophobia conjugated to one or more functional or detectable moieties. The invention also contemplates methods of making said chimeric or humanized anti-CGRP or anti-CGRP-R antibodies or anti-CGRP/CGRP-R complex antibodies and binding fragments thereof for treatment or prevention of photophobia. In one embodiment, binding fragments include, but are not limited to, Fab, Fab', $F(ab')_2$, Fv, scFv fragments, SMIPs (small molecule immunopharmaceuticals), camelbodies, nanobodies, and IgNAR.

Embodiments of the invention pertain to the use of polypeptide CGRP/CGRP receptor inhibitors, e.g., anti-CGRP or anti-CGRP-R antibodies or antibody fragments and CGRP or CGRP-R fragments, preferably anti-CGRP or anti-CGRP-R antibodies and binding fragments thereof for the diagnosis, assessment and treatment of diseases and disorders associated with CGRP or aberrant expression thereof especially for the treatment or prevention of photophobia. The invention also contemplates the use of polypeptide CGRP/CGRP receptor inhibitors, e.g., anti-CGRP or anti-CGRP receptor antibodies or CGRP or CGRP receptor fragments, especially fragments of anti-CGRP antibodies for the diagnosis, assessment and treatment of diseases and disorders associated with CGRP or aberrant expression thereof especially for treatment or prevention of photophobia. Other embodiments of the invention relate to the production of anti-CGRP or anti-CGRP receptor antibodies or fragments thereof in recombinant host cells, for example mammalian cells such as CHO, NSO or HEK 293 cells, or yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains.

BRIEF DESCRIPTION OF THE SEVERAL VIEWS OF THE DRAWINGS

FIG. **1**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab1.

FIG. **2**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab2.

FIG. **3**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab3.

FIG. **4**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab4.

FIG. **5**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab5.

FIG. **6**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab6.

FIG. **7**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab7.

FIG. **8**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab8.

FIG. **9**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab9.

FIG. **10**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab10.

FIG. **11**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab11.

FIG. **12**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab12.

FIG. **13**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab13.

FIG. **14**A-C provides polynucleotide and polypeptide sequences corresponding to the full-length Antibody Ab14.

FIG. **15** provides the CGRP-alpha ELISA binding data obtained following the protocol in Example 1 infra for antibodies Ab1, Ab2, Ab3, and Ab4.

FIG. **16** provides the CGRP-alpha ELISA binding data obtained following the protocol in Example 1 infra for antibodies Ab5, Ab6, Ab7, and Ab8.

FIG. **17** provides the CGRP-alpha ELISA binding data obtained following the protocol in Example 1 infra for antibodies Ab9, Ab10, and Ab14.

FIG. **18** provides the CGRP-alpha ELISA binding data obtained following the protocol in Example 1 infra for antibodies Ab11, Ab12, and Ab13.

FIG. **19** demonstrates the inhibition of CGRP-alpha-driven cAMP production by antibodies Ab1, Ab2, and Ab4, obtained following the protocol in Example 1 infra.

FIG. **20** demonstrates the inhibition of CGRP-alpha-driven cAMP production by antibody Ab3, obtained following the protocol in Example 1 infra.

FIG. **21** demonstrates the inhibition of CGRP-alpha-driven cAMP production by antibodies Ab5 and Ab6, obtained following the protocol in Example 1 infra.

FIG. **22** demonstrates the inhibition of CGRP-alpha-driven cAMP production by antibodies Ab7, Ab8, Ab9, and Ab10, obtained following the protocol in Example 1 infra.

FIG. **23** demonstrates the inhibition of CGRP-alpha-driven cAMP production by antibodies Ab11, Ab12, and Ab13, obtained following the protocol in Example 1 infra.

FIG. **24** demonstrates the inhibition of CGRP-alpha-driven cAMP production by antibody Ab14, obtained following the protocol in Example 1 infra.

FIG. **25** demonstrates the inhibition of CGRP-beta-driven cAMP production by antibodies Ab1, Ab2, and Ab3, obtained following the protocol in Example 1 infra.

FIG. **26** demonstrates the inhibition of CGRP-beta-driven cAMP production by antibodies Ab4, Ab5, and Ab6, obtained following the protocol in Example 1 infra.

FIG. **27** demonstrates the inhibition of CGRP-beta-driven cAMP production by antibodies Ab7 and Ab8, obtained following the protocol in Example 1 infra.

FIG. **28** demonstrates the inhibition of CGRP-beta-driven cAMP production by antibodies Ab9, Ab10, and Ab14, obtained following the protocol in Example 1 infra.

FIG. **29** demonstrates the inhibition of CGRP-beta-driven cAMP production by antibodies Ab11, Ab12, and Ab13, obtained following the protocol in Example 1 infra.

FIG. **30** demonstrates the inhibition of rat CGRP-driven cAMP production by antibodies Ab1, Ab2, Ab4, and Ab5, obtained following the protocol in Example 1 infra.

FIG. **31** demonstrates the inhibition of rat CGRP-driven cAMP production by antibodies Ab3 and Ab6, obtained following the protocol in Example 1 infra.

FIG. **32** demonstrates the inhibition of rat CGRP-driven cAMP production by antibodies Ab7 and Ab8, obtained following the protocol in Example 1 infra.

FIG. **33** demonstrates the inhibition of rat CGRP-driven cAMP production by antibody Ab9, obtained following the protocol in Example 1 infra.

FIG. **34** demonstrates the inhibition of rat CGRP-driven cAMP production by antibody Ab10, obtained following the protocol in Example 1 infra.

FIG. **35** demonstrates the inhibition of rat CGRP-driven cAMP production by antibodies Ab11 and Ab12, obtained following the protocol in Example 1 infra.

FIG. **36** demonstrates the inhibition of rat CGRP-driven cAMP production by antibody Ab13, obtained following the protocol in Example 1 infra.

FIG. **37** demonstrates the inhibition of rat CGRP-driven cAMP production by antibody Ab14, obtained following the protocol in Example 1 infra.

FIG. **38** demonstrates the inhibition of binding of radio-labeled CGRP to CGRP-R by antibodies Ab1-Ab13, obtained following the protocol in Example 6 infra.

FIG. **39** demonstrates a reduction in vasodilation obtained by administering antibodies Ab3 and Ab6 following capsaicin administration in a rat model, relative to a control antibody, obtained following the protocol in Example 7 infra.

FIG. **40** demonstrates a reduction in vasodilation obtained by administering antibody Ab6 at differing concentrations following capsaicin administration in a rat model, relative to a control antibody, obtained following the protocol in Example 7 infra.

FIG. **41** shows the effect of ICV injection of CGRP in hRAMP1 tg mice and control littermate mice and in particular contains data that shows that CGRP administration decreases time in light behavior in the hRAMP1 tg mice relative to their control littermates. Mice were injected hCGRP (2 ug) via ICV under anesthesia and allowed to recover for 30 minutes. Mice were placed individually in the two chamber light/dark boxes and movement was recorded for 30 minutes. Six mice were run in parallel at a time in six different boxes. Each group consisted of seven to nine mice.

FIG. **42** contains data that compares the effect of systemic (IP) injection of anti-CGRP antibody (Ab3) on CGRP driven light aversion. Ab3 in in vehicle, vehicle, and control antibody in vehicle were administered at a dosage of 30 mg/kg in Nestin/RAMP1 mice and thereafter mice were administered CGRP via ICV administration. The data in the left side of the graph is the total time in light (seconds) for the first 10 minutes post-CGRP administration, and the data on the right side of the graph is the total time in light (seconds) for the first 20 minutes measured post-CGRP injection. The data reveal that the mice who received the anti-CGRP antibody Ab3 (disclosed infra) had a statistically significant increase in the amount of time spent in the light relative to the mice who received the controls.

DETAILED DESCRIPTION OF PREFERRED EMBODIMENTS

Definitions

It is to be understood that this invention is not limited to the particular methodology, protocols, cell lines, animal species or genera, and reagents described, as such may vary. It is also to be understood that the terminology used herein is for the purpose of describing particular embodiments only, and is not intended to limit the scope of the present invention which will be limited only by the appended claims. As used herein the singular forms "a", "and", and "the" include plural referents unless the context clearly dictates otherwise. Thus, for example, reference to "a cell" includes a plurality of such cells and reference to "the protein" includes reference to one or more proteins and equivalents thereof known to those skilled in the art, and so forth. All technical and scientific terms used herein have the same meaning as commonly understood to one of ordinary skill in the art to which this invention belongs unless clearly indicated otherwise.

Calcitonin Gene Related Peptide (CGRP): As used herein, CGRP encompasses not only the following *Homo sapiens* CGRP-alpha and *Homo sapiens* CGRP-beta amino acid sequences available from American Peptides (Sunnyvale Calif.) and Bachem (Torrance, Calif.):

CGRP-alpha: ACDTATCVTHRLAGLLSRSGGVVKNNFVPTNVGSKAF-NH_2 (SEQ ID NO: 281), wherein the N-terminal phenylalanine is amidated;

CGRP-beta: ACNTATCVTHRLAGLLSRSGGMVKSNFVPTNVGSKAF-NH_2 (SEQ ID NO: 282), wherein the N-terminal phenylalanine is amidated; but also any membrane-bound forms of these CGRP amino acid sequences, as well as mutants (mutiens), splice variants, isoforms, orthologues, homologues and variants of this sequence. In particular CGRP herein encompasses rodent (rat or mouse) CGRP as well as CGRP from other mammals.

"CGRP receptor" or "CGRP-R" refers to the receptor binding partner of CGRP, preferably the human CGRP receptor, but encompassing other species CGRP-R's, especially rodent (rat or mouse), non-human primate and other mammalian CGRP-R's.

"CGRP/CGRP receptor inhibitor" herein refers to any polypeptide that inhibits the interaction of CGRP and CGRP receptors, e.g., anti-CGRP or anti-CGRP-R antibodies or antibody fragments and fragments of CGRP or CGRP-R polypeptides. Preferably these inhibitors will inhibit this interaction in vitro and in vivo and will inhibit the adverse side effects of CGRP including photoaversion or photophobia.

"Photophobia" herein refers to a symptom of abnormal intolerance to visual perception of light, sometimes additionally defined by abnormal or irrational fear of light, or by presence of actual physical photosensitivity of the eyes. In

the present invention photophobia includes in particular light aversion associated with migraine, cluster headaches and other neurological causes of light aversive behavior that may trigger a migraine or cluster headache. Patients may develop photophobia as a result of several different medical conditions, related to the eye or the nervous system. Photophobia can be caused by an increased response to light starting at any step in the visual system such as: (i) too much light entering the eye, (ii) too much light can enter the eye if it is damaged, such as with corneal abrasion and retinal damage, or if a pupil(s) is unable to normally constrict (seen with damage to the oculomotor nerve, (iii) overstimulation of the photoreceptors in the retina, (iv) excessive electric impulses to the optic nerve, and (v) excessive response in the central nervous system.

Common causes of photophobia include migraine headaches, cataracts, or severe ophthalmologic diseases such as uveitis or corneal abrasion. A more extensive list of disorders associated with photophobia includes eye related causes such as Achromatopsia, Aniridia, Anticholinergic drugs may cause photophobia by paralyzing the iris sphincter muscle, Aphakia (absence of the lens of the eye), Buphthalmos (abnormally narrow angle between the cornea and iris), Cataracts, Cone dystrophy, Congenital abnormalities of the eye, Viral conjunctivitis ("pink eye") Corneal abrasion, Corneal dystrophy, Corneal ulcer, disruption of the corneal epithelium, such as that caused by a corneal foreign body or keratitis, Ectopia lentis, Endophthalmitis, Eye trauma caused by disease, injury, or infection such as chalazion, episcleritis, glaucoma, keratoconus, or optic nerve hypoplasia, Hydrophthalmos, or congenital glaucoma Iritis, Optic neuritis, Pigment dispersion syndrom, Pupillary dilation (naturally or chemically induced), Retinal detachment, Scarring of the cornea or sclera and Uveitis.

In addition photophobia has nervous-system-related or urological causes including: Autism spectrum disorders, Chiari malformation, Dyslexia, Encephalitis including Myalgic encephalomyelitis aka Chronic fatigue syndrome, Meningitis, Subarachnoid haemorrhage, Tumor of the posterior cranial fossa, as well as other causes such as Ankylosing spondylitis, Albinism, Ariboflavinosis, Benzodiazepines (long term use of or withdrawal from benzodiazepines), Chemotherapy, Chikungunya, Cystinosis, Ehlers-Danlos syndrome, Hangover, Influenza, Infectious Mononucleosis, Magnesium deficiency, Mercury poisoning, Migraine, Rabies, and Tyrosinemia type II, also known as "Richner-Hanhart syndrome". Additionally it is known that photophobia is elevated in depression, bipolar disorder and agoraphobia.

"Migraine" from the Greek words hemi, meaning half, and kranion, meaning skull) is a debilitating condition characterized by moderate to severe headaches, and nausea. It is about three times more common in women than in men. The typical migraine headache is unilateral (affecting one half of the head) and pulsating in nature and lasting from 4 to 72 hours; symptoms include nausea, vomiting, photophobia (increased sensitivity to light), phonophobia (increased sensitivity to sound); the symptoms are generally aggravated by routine activity. Approximately one-third of people who suffer from migraine headaches perceive an aura-unusual visual, olfactory, or other sensory experiences that are a sign that the migraine will soon occur. Initial treatment of migraine headaches typically is with analgesics for the headache, an antiemetic for the nausea, and the avoidance of triggering conditions. Studies of twins indicate a 60- to 65-percent genetic influence upon their propensity to develop migraine headaches. Moreover, fluctuating hormone levels indicate a migraine relation: 75 percent of adult patients are women, although migraine affects approximately equal numbers of prepubescent boys and girls; propensity to migraine headache is known to disappear during pregnancy, although in some women migraines may become more frequent during pregnancy.

"Effective treatment or prevention of photophobia" herein refers to inhibiting light aversive behavior or photophobia or inhibiting the onset of light aversive behavior or photophobia in a subject in need thereof, e.g., a subject having an active migraine attack or cluster headache or a subject prone to migraine or cluster headaches, or one of the other photophobia-associated disorders identified herein after administration of an effective amount of an CGRP/CGRP receptor inhibitor polypeptide according to the invention, e.g., an anti-CGRP antibody or antibody fragment according to the invention. The treatment may be effected as a monotherapy or in association with another active agent such as Topirimate or dihydroergotamine by way of example.

Mating Competent Yeast Species: In the present invention this is intended to broadly encompass any diploid or tetraploid yeast which can be grown in culture. Such species of yeast may exist in a haploid, diploid, or other polyploid form. The cells of a given ploidy may, under appropriate conditions, proliferate for an indefinite number of generations in that form. Diploid cells can also sporulate to form haploid cells. Sequential mating can result in tetraploid strains through further mating or fusion of diploid strains. The present invention contemplates the use of haploid yeast, as well as diploid or other polyploid yeast cells produced, for example, by mating or spheroplast fusion.

In one embodiment of the invention, the mating competent yeast is a member of the Saccharomycetaceae family, which includes the genera *Arxiozyma; Ascobotryozyma; Citeromyces; Debaryomyces; Dekkera; Eremothecium; Issatchenkia; Kazachstania; Kluyveromyces; Kodamaea; Lodderomyces; Pachysolen; Pichia; Saccharomyces; Saturnispora; Tetrapisispora; Torulaspora; Williopsis*; and *Zygosaccharomyces*. Other types of yeast potentially useful in the invention include *Yarrowia; Rhodosporidium; Candida; Hansenula; Filobasium; Sporidiobolus; Bullera; Leucosporidium* and *Filobasidella.*

In a preferred embodiment of the invention, the mating competent yeast is a member of the genus *Pichia*. In a further preferred embodiment of the invention, the mating competent yeast of the genus *Pichia* is one of the following species: *Pichia pastoris, Pichia methanolica*, and *Hansenula polymorpha* (*Pichia angusta*). In a particularly preferred embodiment of the invention, the mating competent yeast of the genus *Pichia* is the species *Pichia pastoris.*

Haploid Yeast Cell: A cell having a single copy of each gene of its normal genomic (chromosomal) complement.

Polyploid Yeast Cell: A cell having more than one copy of its normal genomic (chromosomal) complement.

Diploid Yeast Cell: A cell having two copies (alleles) of essentially every gene of its normal genomic complement, typically formed by the process of fusion (mating) of two haploid cells.

Tetraploid Yeast Cell: A cell having four copies (alleles) of essentially every gene of its normal genomic complement, typically formed by the process of fusion (mating) of two haploid cells. Tetraploids may carry two, three, four or more different expression cassettes. Such tetraploids might be obtained in *S. cerevisiae* by selective mating homozygotic heterothallic a/a and alpha/alpha diploids and in *Pichia* by sequential mating of haploids to obtain auxotrophic diploids. For example, a [met his] haploid can be mated with

[ade his] haploid to obtain diploid [his]; and a [met arg] haploid can be mated with [ade arg] haploid to obtain diploid [arg]; then the diploid [his]×diploid [arg] to obtain a tetraploid prototroph. It will be understood by those of skill in the art that reference to the benefits and uses of diploid cells may also apply to tetraploid cells.

Yeast Mating: The process by which two haploid yeast cells naturally fuse to form one diploid yeast cell.

Meiosis: The process by which a diploid yeast cell undergoes reductive division to form four haploid spore products. Each spore may then germinate and form a haploid vegetatively growing cell line.

Selectable Marker: A selectable marker is a gene or gene fragment that confers a growth phenotype (physical growth characteristic) on a cell receiving that gene as, for example through a transformation event. The selectable marker allows that cell to survive and grow in a selective growth medium under conditions in which cells that do not receive that selectable marker gene cannot grow. Selectable marker genes generally fall into several types, including positive selectable marker genes such as a gene that confers on a cell resistance to an antibiotic or other drug, temperature when two temperature sensitive ("ts") mutants are crossed or a ts mutant is transformed; negative selectable marker genes such as a biosynthetic gene that confers on a cell the ability to grow in a medium without a specific nutrient needed by all cells that do not have that biosynthetic gene, or a mutagenized biosynthetic gene that confers on a cell inability to grow by cells that do not have the wild type gene; and the like. Suitable markers include but are not limited to: ZEO; G418; LYS3; MET1; MET3a; ADE1; ADE3; URA3; and the like.

Expression Vector: These DNA vectors contain elements that facilitate manipulation for the expression of a foreign protein within the target host cell. Conveniently, manipulation of sequences and production of DNA for transformation is first performed in a bacterial host, e.g. *E. coli*, and usually vectors will include sequences to facilitate such manipulations, including a bacterial origin of replication and appropriate bacterial selection marker. Selection markers encode proteins necessary for the survival or growth of transformed host cells grown in a selective culture medium. Host cells not transformed with the vector containing the selection gene will not survive in the culture medium. Typical selection genes encode proteins that (a) confer resistance to antibiotics or other toxins, (b) complement auxotrophic deficiencies, or (c) supply critical nutrients not available from complex media. Exemplary vectors and methods for transformation of yeast are described, for example, in Burke, D., Dawson, D., & Steams, T. (2000). Methods in yeast genetics: a Cold Spring Harbor Laboratory course manual. Plainview, N.Y.: Cold Spring Harbor Laboratory Press.

Expression vectors for use in the methods of the invention will further include yeast specific sequences, including a selectable auxotrophic or drug marker for identifying transformed yeast strains. A drug marker may further be used to amplify copy number of the vector in a yeast host cell.

The polypeptide coding sequence of interest is operably linked to transcriptional and translational regulatory sequences that provide for expression of the polypeptide in yeast cells. These vector components may include, but are not limited to, one or more of the following: an enhancer element, a promoter, and a transcription termination sequence. Sequences for the secretion of the polypeptide may also be included, e.g. a signal sequence, and the like. A yeast origin of replication is optional, as expression vectors are often integrated into the yeast genome. In one embodiment of the invention, the polypeptide of interest is operably linked, or fused, to sequences providing for optimized secretion of the polypeptide from yeast diploid cells.

Nucleic acids are "operably linked" when placed into a functional relationship with another nucleic acid sequence. For example, DNA for a signal sequence is operably linked to DNA for a polypeptide if it is expressed as a preprotein that participates in the secretion of the polypeptide; a promoter or enhancer is operably linked to a coding sequence if it affects the transcription of the sequence. Generally, "operably linked" means that the DNA sequences being linked are contiguous, and, in the case of a secretory leader, contiguous and in reading frame. However, enhancers do not have to be contiguous. Linking is accomplished by ligation at convenient restriction sites or alternatively via a PCR/recombination method familiar to those skilled in the art (Gateway® Technology; Invitrogen, Carlsbad Calif.). If such sites do not exist, the synthetic oligonucleotide adapters or linkers are used in accordance with conventional practice.

Promoters are untranslated sequences located upstream (5') to the start codon of a structural gene (generally within about 100 to 1000 bp) that control the transcription and translation of particular nucleic acid sequences to which they are operably linked. Such promoters fall into several classes: inducible, constitutive, and repressible promoters (that increase levels of transcription in response to absence of a repressor). Inducible promoters may initiate increased levels of transcription from DNA under their control in response to some change in culture conditions, e.g., the presence or absence of a nutrient or a change in temperature.

The yeast promoter fragment may also serve as the site for homologous recombination and integration of the expression vector into the same site in the yeast genome; alternatively a selectable marker is used as the site for homologous recombination. *Pichia* transformation is described in Cregg et al. (1985) *Mol. Cell. Biol.* 5:3376-3385.

Examples of suitable promoters from *Pichia* include the AOX1 and promoter (Cregg et al. (1989) *Mol. Cell. Biol.* 9:1316-1323); ICL1 promoter (Menendez et al. (2003) *Yeast* 20(13):1097-108); glyceraldehyde-3-phosphate dehydrogenase promoter (GAP) (Waterham et al. (1997) *Gene* 186(1): 37-44); and FLD1 promoter (Shen et al. (1998) *Gene* 216(1):93-102). The GAP promoter is a strong constitutive promoter and the AOX and FLD1 promoters are inducible.

Other yeast promoters include ADH1, alcohol dehydrogenase II, GAL4, PHO3, PHO5, Pyk, and chimeric promoters derived therefrom. Additionally, non-yeast promoters may be used in the invention such as mammalian, insect, plant, reptile, amphibian, viral, and avian promoters. Most typically the promoter will comprise a mammalian promoter (potentially endogenous to the expressed genes) or will comprise a yeast or viral promoter that provides for efficient transcription in yeast systems.

The polypeptides of interest may be produced recombinantly not only directly, but also as a fusion polypeptide with a heterologous polypeptide, e.g. a signal sequence or other polypeptide having a specific cleavage site at the N-terminus of the mature protein or polypeptide. In general, the signal sequence may be a component of the vector, or it may be a part of the polypeptide coding sequence that is inserted into the vector. The heterologous signal sequence selected preferably is one that is recognized and processed through one of the standard pathways available within the host cell. The *S. cerevisiae* alpha factor pre-pro signal has proven effective in the secretion of a variety of recombinant proteins from *P. pastoris*. Other yeast signal sequences include the alpha

mating factor signal sequence, the invertase signal sequence, and signal sequences derived from other secreted yeast polypeptides. Additionally, these signal peptide sequences may be engineered to provide for enhanced secretion in diploid yeast expression systems. Other secretion signals of interest also include mammalian signal sequences, which may be heterologous to the protein being secreted, or may be a native sequence for the protein being secreted. Signal sequences include pre-peptide sequences, and in some instances may include propeptide sequences. Many such signal sequences are known in the art, including the signal sequences found on immunoglobulin chains, e.g., K28 pre-protoxin sequence, PHA-E, FACE, human MCP-1, human serum albumin signal sequences, human Ig heavy chain, human Ig light chain, and the like. For example, see Hashimoto et. al. Protein Eng 11(2) 75 (1998); and Kobayashi et. al. Therapeutic Apheresis 2(4) 257 (1998).

Transcription may be increased by inserting a transcriptional activator sequence into the vector. These activators are cis-acting elements of DNA, usually about from 10 to 300 bp, which act on a promoter to increase its transcription. Transcriptional enhancers are relatively orientation and position independent, having been found 5' and 3' to the transcription unit, within an intron, as well as within the coding sequence itself. The enhancer may be spliced into the expression vector at a position 5' or 3' to the coding sequence, but is preferably located at a site 5' from the promoter.

Expression vectors used in eukaryotic host cells may also contain sequences necessary for the termination of transcription and for stabilizing the mRNA. Such sequences are commonly available from 3' to the translation termination codon, in untranslated regions of eukaryotic or viral DNAs or cDNAs. These regions contain nucleotide segments transcribed as polyadenylated fragments in the untranslated portion of the mRNA.

Construction of suitable vectors containing one or more of the above-listed components employs standard ligation techniques or PCR/recombination methods. Isolated plasmids or DNA fragments are cleaved, tailored, and re-ligated in the form desired to generate the plasmids required or via recombination methods. For analysis to confirm correct sequences in plasmids constructed, the ligation mixtures are used to transform host cells, and successful transformants selected by antibiotic resistance (e.g. ampicillin or Zeocin) where appropriate. Plasmids from the transformants are prepared, analyzed by restriction endonuclease digestion and/or sequenced.

As an alternative to restriction and ligation of fragments, recombination methods based on att sites and recombination enzymes may be used to insert DNA sequences into a vector. Such methods are described, for example, by Landy (1989) Ann. Rev. Biochem. 58:913-949; and are known to those of skill in the art. Such methods utilize intermolecular DNA recombination that is mediated by a mixture of lambda and *E. coli*-encoded recombination proteins. Recombination occurs between specific attachment (att) sites on the interacting DNA molecules. For a description of att sites see Weisberg and Landy (1983) Site-Specific Recombination in Phage Lambda, in Lambda II, Weisberg, ed. (Cold Spring Harbor, N.Y.: Cold Spring Harbor Press), pp. 211-250. The DNA segments flanking the recombination sites are switched, such that after recombination, the att sites are hybrid sequences comprised of sequences donated by each parental vector. The recombination can occur between DNAs of any topology.

Att sites may be introduced into a sequence of interest by ligating the sequence of interest into an appropriate vector; generating a PCR product containing att B sites through the use of specific primers; generating a cDNA library cloned into an appropriate vector containing att sites; and the like.

Folding, as used herein, refers to the three-dimensional structure of polypeptides and proteins, where interactions between amino acid residues act to stabilize the structure. While non-covalent interactions are important in determining structure, usually the proteins of interest will have intra- and/or intermolecular covalent disulfide bonds formed by two cysteine residues. For naturally occurring proteins and polypeptides or derivatives and variants thereof, the proper folding is typically the arrangement that results in optimal biological activity, and can conveniently be monitored by assays for activity, e.g. ligand binding, enzymatic activity, etc.

In some instances, for example where the desired product is of synthetic origin, assays based on biological activity will be less meaningful. The proper folding of such molecules may be determined on the basis of physical properties, energetic considerations, modeling studies, and the like.

The expression host may be further modified by the introduction of sequences encoding one or more enzymes that enhance folding and disulfide bond formation, i.e. foldases, chaperonins, etc. Such sequences may be constitutively or inducibly expressed in the yeast host cell, using vectors, markers, etc. as known in the art. Preferably the sequences, including transcriptional regulatory elements sufficient for the desired pattern of expression, are stably integrated in the yeast genome through a targeted methodology.

For example, the eukaryotic PDI is not only an efficient catalyst of protein cysteine oxidation and disulfide bond isomerization, but also exhibits chaperone activity. Co-expression of PDI can facilitate the production of active proteins having multiple disulfide bonds. Also of interest is the expression of BIP (immunoglobulin heavy chain binding protein); cyclophilin; and the like. In one embodiment of the invention, each of the haploid parental strains expresses a distinct folding enzyme, e.g. one strain may express BIP, and the other strain may express PDI or combinations thereof.

The terms "desired protein" or "desired antibody" are used interchangeably and refer generally to a parent antibody specific to a target, i.e., CGRP or CGRP receptor or a chimeric or humanized antibody or a binding portion thereof derived therefrom as described herein. The term "antibody" is intended to include any polypeptide chain-containing molecular structure with a specific shape that fits to and recognizes an epitope, where one or more non-covalent binding interactions stabilize the complex between the molecular structure and the epitope. The archetypal antibody molecule is the immunoglobulin, and all types of immunoglobulins, IgG, IgM, IgA, IgE, IgD, etc., from all sources, e.g. human, rodent, rabbit, cow, sheep, pig, dog, other mammals, chicken, other avians, etc., are considered to be "antibodies." A preferred source for producing antibodies useful as starting material according to the invention is rabbits. Numerous antibody coding sequences have been described; and others may be raised by methods well-known in the art. Examples thereof include chimeric antibodies, human antibodies and other non-human mammalian antibodies, humanized antibodies, single chain antibodies (such as scFvs), camelbodies, nanobodies, IgNAR (single-chain antibodies derived from sharks), small-modular immunopharmaceuticals (SMIPs), and antibody fragments such as Fabs, Fab', $F(ab')_2$ and the like. See Streltsov V A, et al.,

Structure of a shark IgNAR antibody variable domain and modeling of an early-developmental isotype, Protein Sci. 2005 November; 14(11):2901-9. Epub 2005 Sep. 30; Greenberg A S, et al., A new antigen receptor gene family that undergoes rearrangement and extensive somatic diversification in sharks, Nature. 1995 Mar. 9; 374(6518):168-73; Nuttall S D, et al., Isolation of the new antigen receptor from wobbegong sharks, and use as a scaffold for the display of protein loop libraries, Mol Immunol. 2001 August; 38(4): 313-26; Hamers-Casterman C, et al., Naturally occurring antibodies devoid of light chains, Nature. 1993 Jun. 3; 363(6428):446-8; Gill D S, et al., Biopharmaceutical drug discovery using novel protein scaffolds, Curr Opin Biotechnol. 2006 December; 17(6):653-8. Epub 2006 Oct. 19.

For example, antibodies or antigen binding fragments may be produced by genetic engineering. In this technique, as with other methods, antibody-producing cells are sensitized to the desired antigen or immunogen. The messenger RNA isolated from antibody producing cells is used as a template to make cDNA using PCR amplification. A library of vectors, each containing one heavy chain gene and one light chain gene retaining the initial antigen specificity, is produced by insertion of appropriate sections of the amplified immunoglobulin cDNA into the expression vectors. A combinatorial library is constructed by combining the heavy chain gene library with the light chain gene library. This results in a library of clones which co-express a heavy and light chain (resembling the Fab fragment or antigen binding fragment of an antibody molecule). The vectors that carry these genes are co-transfected into a host cell. When antibody gene synthesis is induced in the transfected host, the heavy and light chain proteins self-assemble to produce active antibodies that can be detected by screening with the antigen or immunogen.

Antibody coding sequences of interest include those encoded by native sequences, as well as nucleic acids that, by virtue of the degeneracy of the genetic code, are not identical in sequence to the disclosed nucleic acids, and variants thereof. Variant polypeptides can include amino acid (aa) substitutions, additions or deletions. The amino acid substitutions can be conservative amino acid substitutions or substitutions to eliminate non-essential amino acids, such as to alter a glycosylation site, or to minimize misfolding by substitution or deletion of one or more cysteine residues that are not necessary for function. Variants can be designed so as to retain or have enhanced biological activity of a particular region of the protein (e.g., a functional domain, catalytic amino acid residues, etc). Variants also include fragments of the polypeptides disclosed herein, particularly biologically active fragments and/or fragments corresponding to functional domains. Techniques for in vitro mutagenesis of cloned genes are known. Also included in the subject invention are polypeptides that have been modified using ordinary molecular biological techniques so as to improve their resistance to proteolytic degradation or to optimize solubility properties or to render them more suitable as a therapeutic agent.

Chimeric antibodies may be made by recombinant means by combining the variable light and heavy chain regions (V_L and V_H), obtained from antibody producing cells of one species with the constant light and heavy chain regions from another. Typically chimeric antibodies utilize rodent or rabbit variable regions and human constant regions, in order to produce an antibody with predominantly human domains. The production of such chimeric antibodies is well known in the art, and may be achieved by standard means (as described, e.g., in U.S. Pat. No. 5,624,659, incorporated herein by reference in its entirety). It is further contemplated that the human constant regions of chimeric antibodies of the invention may be selected from IgG1, IgG2, IgG3, and IgG4 constant regions.

Humanized antibodies are engineered to contain even more human-like immunoglobulin domains, and incorporate only the complementarity-determining regions of the animal-derived antibody. This is accomplished by carefully examining the sequence of the hyper-variable loops of the variable regions of the monoclonal antibody, and fitting them to the structure of the human antibody chains. Although facially complex, the process is straightforward in practice. See, e.g., U.S. Pat. No. 6,187,287, incorporated fully herein by reference.

In addition to entire immunoglobulins (or their recombinant counterparts), immunoglobulin fragments comprising the epitope binding site (e.g., Fab', $F(ab')_2$, or other fragments) may be synthesized. "Fragment," or minimal immunoglobulins may be designed utilizing recombinant immunoglobulin techniques. For instance "Fv" immunoglobulins for use in the present invention may be produced by synthesizing a fused variable light chain region and a variable heavy chain region. Combinations of antibodies are also of interest, e.g. diabodies, which comprise two distinct Fv specificities. In another embodiment of the invention, SMIPs (small molecule immunopharmaceuticals), camelbodies, nanobodies, and IgNAR are encompassed by immunoglobulin fragments.

Immunoglobulins and fragments thereof may be modified post-translationally, e.g. to add effector moieties such as chemical linkers, detectable moieties, such as fluorescent dyes, enzymes, toxins, substrates, bioluminescent materials, radioactive materials, chemiluminescent moieties and the like, or specific binding moieties, such as streptavidin, avidin, or biotin, and the like may be utilized in the methods and compositions of the present invention. Examples of additional effector molecules are provided infra.

A polynucleotide sequence "corresponds" to a polypeptide sequence if translation of the polynucleotide sequence in accordance with the genetic code yields the polypeptide sequence (i.e., the polynucleotide sequence "encodes" the polypeptide sequence), one polynucleotide sequence "corresponds" to another polynucleotide sequence if the two sequences encode the same polypeptide sequence.

A "heterologous" region or domain of a DNA construct is an identifiable segment of DNA within a larger DNA molecule that is not found in association with the larger molecule in nature. Thus, when the heterologous region encodes a mammalian gene, the gene will usually be flanked by DNA that does not flank the mammalian genomic DNA in the genome of the source organism. Another example of a heterologous region is a construct where the coding sequence itself is not found in nature (e.g., a cDNA where the genomic coding sequence contains introns, or synthetic sequences having codons different than the native gene). Allelic variations or naturally-occurring mutational events do not give rise to a heterologous region of DNA as defined herein.

A "coding sequence" is an in-frame sequence of codons that (in view of the genetic code) correspond to or encode a protein or peptide sequence. Two coding sequences correspond to each other if the sequences or their complementary sequences encode the same amino acid sequences. A coding sequence in association with appropriate regulatory sequences may be transcribed and translated into a polypeptide. A polyadenylation signal and transcription termination sequence will usually be located 3' to the coding sequence.

A "promoter sequence" is a DNA regulatory region capable of binding RNA polymerase in a cell and initiating transcription of a downstream (3' direction) coding sequence. Promoter sequences typically contain additional sites for binding of regulatory molecules (e.g., transcription factors) which affect the transcription of the coding sequence. A coding sequence is "under the control" of the promoter sequence or "operatively linked" to the promoter when RNA polymerase binds the promoter sequence in a cell and transcribes the coding sequence into mRNA, which is then in turn translated into the protein encoded by the coding sequence.

Vectors are used to introduce a foreign substance, such as DNA, RNA or protein, into an organism or host cell. Typical vectors include recombinant viruses (for polynucleotides) and liposomes (for polypeptides). A "DNA vector" is a replicon, such as plasmid, phage or cosmid, to which another polynucleotide segment may be attached so as to bring about the replication of the attached segment. An "expression vector" is a DNA vector which contains regulatory sequences which will direct polypeptide synthesis by an appropriate host cell. This usually means a promoter to bind RNA polymerase and initiate transcription of mRNA, as well as ribosome binding sites and initiation signals to direct translation of the mRNA into a polypeptide(s). Incorporation of a polynucleotide sequence into an expression vector at the proper site and in correct reading frame, followed by transformation of an appropriate host cell by the vector, enables the production of a polypeptide encoded by said polynucleotide sequence.

"Amplification" of polynucleotide sequences is the in vitro production of multiple copies of a particular nucleic acid sequence. The amplified sequence is usually in the form of DNA. A variety of techniques for carrying out such amplification are described in a review article by Van Brunt (1990, Bio/Technol., 8(4):291-294). Polymerase chain reaction or PCR is a prototype of nucleic acid amplification, and use of PCR herein should be considered exemplary of other suitable amplification techniques.

The general structure of antibodies in vertebrates now is well understood (Edelman, G. M., Ann. N.Y. Acad. Sci., 190: 5 (1971)). Antibodies consist of two identical light polypeptide chains of molecular weight approximately 23,000 daltons (the "light chain"), and two identical heavy chains of molecular weight 53,000-70,000 (the "heavy chain"). The four chains are joined by disulfide bonds in a "Y" configuration wherein the light chains bracket the heavy chains starting at the mouth of the "Y" configuration. The "branch" portion of the "Y" configuration is designated the F_{ab} region; the stem portion of the "Y" configuration is designated the F_C region. The amino acid sequence orientation runs from the N-terminal end at the top of the "Y" configuration to the C-terminal end at the bottom of each chain. The N-terminal end possesses the variable region having specificity for the antigen that elicited it, and is approximately 100 amino acids in length, there being slight variations between light and heavy chain and from antibody to antibody.

The variable region is linked in each chain to a constant region that extends the remaining length of the chain and that within a particular class of antibody does not vary with the specificity of the antibody (i.e., the antigen eliciting it). There are five known major classes of constant regions that determine the class of the immunoglobulin molecule (IgG, IgM, IgA, IgD, and IgE corresponding to γ, μ, α, δ, and ε (gamma, mu, alpha, delta, or epsilon) heavy chain constant regions). The constant region or class determines subsequent effector function of the antibody, including activation of complement (Kabat, E. A., Structural Concepts in Immunology and Immunochemistry, 2nd Ed., p. 413-436, Holt, Rinehart, Winston (1976)), and other cellular responses (Andrews, D. W., et al., Clinical Immunobiology, pp 1-18, W. B. Sanders (1980); Kohl, S., et al., Immunology, 48: 187 (1983)); while the variable region determines the antigen with which it will react. Light chains are classified as either κ (kappa) or λ (lambda). Each heavy chain class can be prepared with either kappa or lambda light chain. The light and heavy chains are covalently bonded to each other, and the "tail" portions of the two heavy chains are bonded to each other by covalent disulfide linkages when the immunoglobulins are generated either by hybridomas or by B cells.

The expression "variable region" or "VR" refers to the domains within each pair of light and heavy chains in an antibody that are involved directly in binding the antibody to the antigen. Each heavy chain has at one end a variable domain (V_H) followed by a number of constant domains. Each light chain has a variable domain (V_L) at one end and a constant domain at its other end; the constant domain of the light chain is aligned with the first constant domain of the heavy chain, and the light chain variable domain is aligned with the variable domain of the heavy chain.

The expressions "complementarity determining region," "hypervariable region," or "CDR" refer to one or more of the hyper-variable or complementarity determining regions (CDRs) found in the variable regions of light or heavy chains of an antibody (See Kabat, E. A. et al., Sequences of Proteins of Immunological Interest, National Institutes of Health, Bethesda, Md., (1987)). These expressions include the hypervariable regions as defined by Kabat et al. ("Sequences of Proteins of Immunological Interest," Kabat E., et al., US Dept. of Health and Human Services, 1983) or the hypervariable loops in 3-dimensional structures of antibodies (Chothia and Lesk, J Mol. Biol. 196 901-917 (1987)). The CDRs in each chain are held in close proximity by framework regions and, with the CDRs from the other chain, contribute to the formation of the antigen binding site. Within the CDRs there are select amino acids that have been described as the selectivity determining regions (SDRs) which represent the critical contact residues used by the CDR in the antibody-antigen interaction (Kashmiri, S., Methods, 36:25-34 (2005)).

The expressions "framework region" or "FR" refer to one or more of the framework regions within the variable regions of the light and heavy chains of an antibody (See Kabat, E. A. et al., Sequences of Proteins of Immunological Interest, National Institutes of Health, Bethesda, Md., (1987)). These expressions include those amino acid sequence regions interposed between the CDRs within the variable regions of the light and heavy chains of an antibody.

Anti-CGRP Antibodies and Binding Fragments Thereof Having Binding Activity for CGRP

Antibody Ab1

The present invention broadly contemplates inhibition or prevention of photophobia in a subject in need thereof, e.g., a migraine sufferer or another photophobia associated disorder by administering an effective amount of a CGRP/CGRP receptor inhibitor polypeptide, e.g., an anti-CGRP or an anti-CGRP receptor antibody or fragment thereof or a fragment of CGRP or a CGRP receptor which is capable of effective treatment or prevention of photophobia. This may be determined e.g., using appropriate in vivo models such as the transgenic mice model disclosed in Example 8.

In one exemplary embodiment, the invention includes chimeric antibodies derived from Ab1 having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 1)
QVLTQTASPVSAAVGSTVTINCQASQSVYDNNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFKGSGSGTQFTLTISDLECADAATYYCLGSYDCSSGDC

FVFGGGTEVVVKR.

The invention also includes chimeric antibodies having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 2)
QVLTQTASPVSAAVGSTVTINCQASQSVYDNNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFKGSGSGTQFTLTISDLECADAATYYCLGSYDCSSGDC

FVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes chimeric antibodies having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 3)
QSLEESGGRLVTPGTPLTLTCTVSGLDLSSYYMQWVRQAPGKGLEWIGVI

GINDNTYYASWAKGRFTISRASSTTVDLKMTSLTTEDTATYFCARGDIWG

PGTLVTVSS.

The invention also includes chimeric antibodies having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 4)
QSLEESGGRLVTPGTPLTLTCTVSGLDLSSYYMQWVRQAPGKGLEWIGVI

GINDNTYYASWAKGRFTISRASSTTVDLKMTSLTTEDTATYFCARGDIWG

PGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSW

NSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSN

TKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPE

VTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTV

LHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEM

TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS

KLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies comprising one or more of the polypeptide sequences of SEQ ID NO: 5; SEQ ID NO: 6; and SEQ ID NO: 7 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 1 or the light chain sequence of SEQ ID NO: 2, and/or one or more of the polypeptide sequences of SEQ ID NO: 8; SEQ ID NO: 9; and SEQ ID NO: 10 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 3 or the heavy chain sequence of SEQ ID NO: 4, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 1 or SEQ ID NO: 2. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 3 or SEQ ID NO: 4.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 5; SEQ ID NO: 6; and SEQ ID NO: 7 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 1 or the light chain sequence of SEQ ID NO: 2.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 8; SEQ ID NO: 9; and SEQ ID NO: 10 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 3 or the heavy chain sequence of SEQ ID NO: 4.

The invention also contemplates antibody fragments which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 1; the variable heavy chain region of SEQ ID NO: 3; the complementarity-determining regions (SEQ ID NO: 5; SEQ ID NO: 6; and SEQ ID NO: 7) of the variable light chain region of SEQ ID NO: 1; and the complementarity-determining regions (SEQ ID NO: 8; SEQ ID NO: 9; and SEQ ID NO: 10) of the variable heavy chain region of SEQ ID NO: 3.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody is Ab1, comprising, or alternatively consisting of, SEQ ID NO: 2 and SEQ ID NO: 4, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab1, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 1 and the variable heavy chain sequence of SEQ ID NO: 3. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 1 and/or SEQ ID NO: 3 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments may for potential treatment or prevention of photophobia be produced by enzymatic digestion (e.g., papain) of Ab1. In another embodiment of the invention, anti-CGRP antibodies such as Ab1 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid

yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab2

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 11)
QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSSGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 12)
QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSSGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 13)
EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGV

IGINDNTYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 14)
EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGV

IGINDNTYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 15; SEQ ID NO: 16; and SEQ ID NO: 17 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 11 or the light chain sequence of SEQ ID NO: 12, and/or one or more of the polypeptide sequences of SEQ ID NO: 18; SEQ ID NO: 19; and SEQ ID NO: 20 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 13 or the heavy chain sequence of SEQ ID NO: 14, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 11 or SEQ ID NO: 12. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 13 or SEQ ID NO: 14.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 15; SEQ ID NO: 16; and SEQ ID NO: 17 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 11 or the light chain sequence of SEQ ID NO: 12.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 18; SEQ ID NO: 19; and SEQ ID NO: 20 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 13 or the heavy chain sequence of SEQ ID NO: 14.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 11; the variable heavy chain region of SEQ ID NO: 13; the complementarity-determining regions (SEQ ID NO: 15; SEQ ID NO: 16; and SEQ ID NO: 17) of the variable light chain region of SEQ ID NO: 11; and the complementarity-determining regions (SEQ ID NO: 18; SEQ ID NO: 19; and SEQ ID NO: 20) of the variable heavy chain region of SEQ ID NO: 13.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab2, comprising, or alternatively consisting of, SEQ ID NO: 12 and SEQ ID NO: 14, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist

of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab2, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 11 and the variable heavy chain sequence of SEQ ID NO: 13. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 11 and/or SEQ ID NO: 13 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab2. In another embodiment of the invention, anti-CGRP antibodies such as Ab2 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab3

In a preferred embodiment, the invention includes humanized antibodies having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below. As disclosed in Example 8 this antibody has been demonstrated in a transgenic mouse light aversion behavioral model to effectively inhibit CGRP-associated photophobia:

(SEQ ID NO: 21)
VLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSSGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 22)
QVLTQSPSSLSASVGDRVTINCQASQSVYDNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSSGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 23)
EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGV

IGINDNTYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 24)
EVQLVESGGGLVQPGGSLRLSCAVSGLDLSSYYMQWVRQAPGKGLEWVGV

IGINDNTYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDARVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 25; SEQ ID NO: 26; and SEQ ID NO: 27 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 21 or the light chain sequence of SEQ ID NO: 22, and/or one or more of the polypeptide sequences of SEQ ID NO: 28; SEQ ID NO: 29; and SEQ ID NO: 30 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 23 or the heavy chain sequence of SEQ ID NO: 24, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 21 or SEQ ID NO: 22. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 23 or SEQ ID NO: 24.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 25; SEQ ID NO: 26; and SEQ ID NO: 27 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 21 or the light chain sequence of SEQ ID NO: 22.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 28; SEQ ID NO: 29; and SEQ ID NO: 30 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 23 or the heavy chain sequence of SEQ ID NO: 24.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or

alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 21; the variable heavy chain region of SEQ ID NO: 23; the complementarity-determining regions (SEQ ID NO: 25; SEQ ID NO: 26; and SEQ ID NO: 27) of the variable light chain region of SEQ ID NO: 21; and the complementarity-determining regions (SEQ ID NO: 28; SEQ ID NO: 29; and SEQ ID NO: 30) of the variable heavy chain region of SEQ ID NO: 23.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody for treatment or prevention of photophobia is Ab3, comprising, or alternatively consisting of, SEQ ID NO: 22 and SEQ ID NO: 24, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab3, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 21 and the variable heavy chain sequence of SEQ ID NO: 23. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 21 and/or SEQ ID NO: 23 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab3. In another embodiment of the invention, anti-CGRP antibodies such as Ab3 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab4

In one embodiment, the invention includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 31)
QVLTQTPSPVSAAVGSTVTINCQASQSVYHNTYLAWYQQKPGQPPKQLIY

DASTLASGVPSRFSGSGSGTQFTLTISGVQCNDAAAYYCLGSYDCTNGDC

FVFGGGTEVVVKR.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 32)
QVLTQTPSPVSAAVGSTVTINCQASQSVYHNTYLAWYQQKPGQPPKQLIY

DASTLASGVPSRFSGSGSGTQFTLTISGVQCNDAAAYYCLGSYDCTNGDC

FVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 33)
QSLEESGGRLVTPGTPLTLTCSVSGIDLSGYYMNWVRQAPGKGLEWIGVI

GINGATYYASWAKGRFTISKTSSTTVDLKMTSLTTEDTATYFCARGDIWG

PGTLVTVSS.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 34)
QSLEESGGRLVTPGTPLTLTCSVSGIDLSGYYMNWVRQAPGKGLEWIGVI

GINGATYYASWAKGRFTISKTSSTTVDLKMTSLTTEDTATYFCARGDIWG

PGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSW

NSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSN

TKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPE

VTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTV

LHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEM

TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS

KLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 35; SEQ ID NO: 36; and SEQ ID NO: 37 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 31 or the light chain sequence of SEQ ID NO: 32, and/or one or more of the polypeptide sequences of SEQ ID NO: 38; SEQ ID NO: 39; and SEQ ID NO: 40 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 33 or the heavy chain sequence of SEQ ID NO: 34, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 31 or SEQ ID NO: 32. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 33 or SEQ ID NO: 34.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 35; SEQ ID NO: 36; and SEQ ID NO: 37 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable

light chain sequence of SEQ ID NO: 31 or the light chain sequence of SEQ ID NO: 32.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 38; SEQ ID NO: 39; and SEQ ID NO: 40 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 33 or the heavy chain sequence of SEQ ID NO: 34.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 31; the variable heavy chain region of SEQ ID NO: 33; the complementarity-determining regions (SEQ ID NO: 35; SEQ ID NO: 36; and SEQ ID NO: 37) of the variable light chain region of SEQ ID NO: 31; and the complementarity-determining regions (SEQ ID NO: 38; SEQ ID NO: 39; and SEQ ID NO: 40) of the variable heavy chain region of SEQ ID NO: 33.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab4, comprising, or alternatively consisting of, SEQ ID NO: 32 and SEQ ID NO: 34, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab4, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 31 and the variable heavy chain sequence of SEQ ID NO: 33. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 31 and/or SEQ ID NO: 33 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab4. In another embodiment of the invention, anti-CGRP antibodies such as Ab4 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab5

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 41)
QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIY

DASTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCTNGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 42)
QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIY

DASTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCTNGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 43)
EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGV

IGINGATYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 44)
EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGV

IGINGATYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 45; SEQ ID NO: 46; and SEQ ID NO: 47 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 41 or the light chain sequence of SEQ ID NO: 42, and/or one or more of the polypeptide sequences of SEQ ID NO: 48; SEQ ID NO: 49; and SEQ ID NO: 50 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 43 or the heavy chain sequence of SEQ ID NO: 44, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 41 or SEQ ID NO: 42. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 43 or SEQ ID NO: 44.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 45; SEQ ID NO: 46; and SEQ ID NO: 47 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 41 or the light chain sequence of SEQ ID NO: 42.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 48; SEQ ID NO: 49; and SEQ ID NO: 50 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 43 or the heavy chain sequence of SEQ ID NO: 44.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 41; the variable heavy chain region of SEQ ID NO: 43; the complementarity-determining regions (SEQ ID NO: 45; SEQ ID NO: 46; and SEQ ID NO: 47) of the variable light chain region of SEQ ID NO: 41; and the complementarity-determining regions (SEQ ID NO: 48; SEQ ID NO: 49; and SEQ ID NO: 50) of the variable heavy chain region of SEQ ID NO: 43.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody for potential treatment or prevention of photophobia is Ab5, comprising, or alternatively consisting of, SEQ ID NO: 42 and SEQ ID NO: 44, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab5, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 41 and the variable heavy chain sequence of SEQ ID NO: 43. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 41 and/or SEQ ID NO: 43 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab5. In another embodiment of the invention, anti-CGRP antibodies such as Ab5 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab6

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 51)
QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIY
DASTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCTNGDC
FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 52)
QVLTQSPSSLSASVGDRVTINCQASQSVYHNTYLAWYQQKPGKVPKQLIY
DASTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCTNGDC
FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK
VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE
VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 53)
EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGV
IGINGATYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI
WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 54)
EVQLVESGGGLVQPGGSLRLSCAVSGIDLSGYYMNWVRQAPGKGLEWVGV
IGINGATYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI
WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV
SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP
SNTKVDARVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT
PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL
TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE
EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL
YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 55;

SEQ ID NO: 56; and SEQ ID NO: 57 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 51 or the light chain sequence of SEQ ID NO: 52, and/or one or more of the polypeptide sequences of SEQ ID NO: 58; SEQ ID NO: 59; and SEQ ID NO: 60 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 53 or the heavy chain sequence of SEQ ID NO: 54, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 51 or SEQ ID NO: 52. In another embodiment of the invention, antibody fragments of the invention for potential treatment or prevention of photophobia comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 53 or SEQ ID NO: 54.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 55; SEQ ID NO: 56; and SEQ ID NO: 57 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 51 or the light chain sequence of SEQ ID NO: 52.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 58; SEQ ID NO: 59; and SEQ ID NO: 60 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 53 or the heavy chain sequence of SEQ ID NO: 54.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 51; the variable heavy chain region of SEQ ID NO: 53; the complementarity-determining regions (SEQ ID NO: 55; SEQ ID NO: 56; and SEQ ID NO: 57) of the variable light chain region of SEQ ID NO: 51; and the complementarity-determining regions (SEQ ID NO: 58; SEQ ID NO: 59; and SEQ ID NO: 60) of the variable heavy chain region of SEQ ID NO: 53.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab6, comprising, or alternatively consisting of, SEQ ID NO: 52 and SEQ ID NO: 54, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab6, the Fab fragment for potential treatment or prevention of photophobia includes the variable light chain sequence of SEQ ID NO: 51 and the variable heavy chain sequence of SEQ ID NO: 53. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 51 and/or SEQ ID NO: 53 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab6. In another embodiment of the invention, anti-CGRP antibodies such as Ab6 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab7

In one embodiment, the invention includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 61)
QVLTQTASPVSAAVGSTVTINCQASQSVYNYNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFKGSGSGTQFTLTISDVQCDDAATYYCLGSYDCSTGDC

FVFGGGTEVVVKR.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 62)
QVLTQTASPVSAAVGSTVTINCQASQSVYNYNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFKGSGSGTQFTLTISDVQCDDAATYYCLGSYDCSTGDC

FVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 63)
QEQLKESGGRLVTPGTSLTLTCTVSGIDLSNHYMQWVRQAPGKGLEWIGV

VGINGRTYYASWAKGRFTISRTSSTTVDLKMTRLTTEDTATYFCARGDIW

GPGTLVTVSS.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 64)
QEQLKESGGRLVTPGTSLTLTCTVSGIDLSNHYMQWVRQAPGKGLEWIGV

VGINGRTYYASWAKGRFTISRTSSTTVDLKMTRLTTEDTATYFCARGDIW

GPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVS

WNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPS

NTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTP

EVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLT

VLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREE

MTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLY

SKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 65; SEQ ID NO: 66; and SEQ ID NO: 67 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 61 or the light chain sequence of SEQ ID NO: 62, and/or one or more of the polypeptide sequences of SEQ ID NO: 68; SEQ ID NO: 69; and SEQ ID NO: 70 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 63 or the heavy chain sequence of SEQ ID NO: 64, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention for potential treatment or prevention of photophobia comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 61 or SEQ ID NO: 62. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 63 or SEQ ID NO: 64.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 65; SEQ ID NO: 66; and SEQ ID NO: 67 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 61 or the light chain sequence of SEQ ID NO: 62.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 68; SEQ ID NO: 69; and SEQ ID NO: 70 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 63 or the heavy chain sequence of SEQ ID NO: 64.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 61; the variable heavy chain region of SEQ ID NO: 63; the complementarity-determining regions (SEQ ID NO: 65; SEQ ID NO: 66; and SEQ ID NO: 67) of the variable light chain region of SEQ ID NO: 61; and the complementarity-determining regions (SEQ ID NO: 68; SEQ ID NO: 69; and SEQ ID NO: 70) of the variable heavy chain region of SEQ ID NO: 63.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody for potential treatment or prevention of photophobia is Ab7, comprising, or alternatively consisting of, SEQ ID NO: 62 and SEQ ID NO: 64, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab7, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 61 and the variable heavy chain sequence of SEQ ID NO: 63. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 61 and/or SEQ ID NO: 63 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab7. In another embodiment of the invention, anti-CGRP antibodies such as Ab7 or Fab fragments thereof for potential treatment or prevention of photophobia may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab8

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 71)
QVLTQSPSSLSASVGDRVTINCQASQSVYNYNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSTGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies having binding specificity to CGRP for potential treatment or prevention of photophobia and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 72)
QVLTQSPSSLSASVGDRVTINCQASQSVYNYNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSTGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

-continued

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 73)

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSNHYMQWVRQAPGKGLEWVGV

VGINGRTYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 74)

EVQLVESGGGLVQPGGSLRLSCAVSGIDLSNHYMQWVRQAPGKGLEWVGV

VGINGRTYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 75; SEQ ID NO: 76; and SEQ ID NO: 77 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 71 or the light chain sequence of SEQ ID NO: 72, and/or one or more of the polypeptide sequences of SEQ ID NO: 78; SEQ ID NO: 79; and SEQ ID NO: 80 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 73 or the heavy chain sequence of SEQ ID NO: 74, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 71 or SEQ ID NO: 72. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 73 or SEQ ID NO: 74.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 75; SEQ ID NO: 76; and SEQ ID NO: 77 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 71 or the light chain sequence of SEQ ID NO: 72.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 78; SEQ ID NO: 79; and SEQ ID NO: 80 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 73 or the heavy chain sequence of SEQ ID NO: 74.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 71; the variable heavy chain region of SEQ ID NO: 73; the complementarity-determining regions (SEQ ID NO: 75; SEQ ID NO: 76; and SEQ ID NO: 77) of the variable light chain region of SEQ ID NO: 71; and the complementarity-determining regions (SEQ ID NO: 78; SEQ ID NO: 79; and SEQ ID NO: 80) of the variable heavy chain region of SEQ ID NO: 73.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab8, comprising, or alternatively consisting of, SEQ ID NO: 72 and SEQ ID NO: 74, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab8, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 71 and the variable heavy chain sequence of SEQ ID NO: 73. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 71 and/or SEQ ID NO: 73 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab8. In another embodiment of the invention, anti-CGRP antibodies such as Ab8 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab9

In one embodiment, the invention includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 81)
QVLTQTPSPVSAAVGSTVTINCQASQNVYNNNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFRGSGSGTQFTLTISDVQCDDAATYYCLGSYDCSRGDC

FVFGGGTEVVVKR.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 82)
QVLTQTPSPVSAAVGSTVTINCQASQNVYNNNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFRGSGSGTQFTLTISDVQCDDAATYYCLGSYDCSRGDC

FVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 83)
QSLEESGGRLVTPGTPLTLTCTVSGIGLSSYYMQWVRQSPGRGLEWIGVI

GSDGKTYYATWAKGRFTISKTSSTTVDLRMASLTTEDTATYFCTRGDIWG

PGTLVTVSS.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 84)
QSLEESGGRLVTPGTPLTLTCTVSGIGLSSYYMQWVRQSPGRGLEWIGVI

GSDGKTYYATWAKGRFTISKTSSTTVDLRMASLTTEDTATYFCTRGDIWG

PGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSW

NSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSN

TKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPE

VTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTV

LHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEM

TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS

KLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 85; SEQ ID NO: 86; and SEQ ID NO: 87 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 81 or the light chain sequence of SEQ ID NO: 82, and/or one or more of the polypeptide sequences of SEQ ID NO: 88; SEQ ID NO: 89; and SEQ ID NO: 90 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 83 or the heavy chain sequence of SEQ ID NO: 84, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 81 or SEQ ID NO: 82. In another embodiment of the invention, antibody fragments of the invention for potential treatment or prevention of photophobia comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 83 or SEQ ID NO: 84.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 85; SEQ ID NO: 86; and SEQ ID NO: 87 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 81 or the light chain sequence of SEQ ID NO: 82.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 88; SEQ ID NO: 89; and SEQ ID NO: 90 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 83 or the heavy chain sequence of SEQ ID NO: 84.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 81; the variable heavy chain region of SEQ ID NO: 83; the complementarity-determining regions (SEQ ID NO: 85; SEQ ID NO: 86; and SEQ ID NO: 87) of the variable light chain region of SEQ ID NO: 81; and the complementarity-determining regions (SEQ ID NO: 88; SEQ ID NO: 89; and SEQ ID NO: 90) of the variable heavy chain region of SEQ ID NO: 83.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody for potential treatment or prevention of photophobia is Ab9, comprising, or alternatively consisting of, SEQ ID NO: 82 and SEQ ID NO: 84, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab9, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 81 and the variable heavy chain sequence of SEQ ID NO: 83. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 81 and/or SEQ ID NO: 83 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab9. In another embodiment of the invention, anti-CGRP antibodies for potential treatment or prevention of photophobia such as Ab9 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab10

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 91)
QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSRGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 92)
QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSRGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 93)
EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGV

IGSDGKTYYATWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCTRGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 94)
EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGV

IGSDGKTYYATWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCTRGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

-continued

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 95; SEQ ID NO: 96; and SEQ ID NO: 97 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 91 or the light chain sequence of SEQ ID NO: 92, and/or one or more of the polypeptide sequences of SEQ ID NO: 98; SEQ ID NO: 99; and SEQ ID NO: 100 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 93 or the heavy chain sequence of SEQ ID NO: 94, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 91 or SEQ ID NO: 92. In another embodiment of the invention, antibody fragments of the invention for potential treatment or prevention of photophobia comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 93 or SEQ ID NO: 94.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 95; SEQ ID NO: 96; and SEQ ID NO: 97 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 91 or the light chain sequence of SEQ ID NO: 92.

In a further embodiment of the invention, fragments of the antibody having binding specificity to CGRP for potential treatment or prevention of photophobia comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 98; SEQ ID NO: 99; and SEQ ID NO: 100 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 93 or the heavy chain sequence of SEQ ID NO: 94.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 91; the variable heavy chain region of SEQ ID NO: 93; the complementarity-determining regions (SEQ ID NO: 95; SEQ ID NO: 96; and SEQ ID NO: 97) of the variable light chain region of SEQ ID NO: 91; and

the complementarity-determining regions (SEQ ID NO: 98; SEQ ID NO: 99; and SEQ ID NO: 100) of the variable heavy chain region of SEQ ID NO: 93.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab10, comprising, or alternatively consisting of, SEQ ID NO: 92 and SEQ ID NO: 94, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab10, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 91 and the variable heavy chain sequence of SEQ ID NO: 93. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 91 and/or SEQ ID NO: 93 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab10. In another embodiment of the invention, anti-CGRP antibodies such as Ab10 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab11

In one embodiment, the invention includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 101)
QVLTQTASPVSPAVGSTVTINCRASQSVYYNNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFKGSGSGTQFTLTISDVQCDDAATYYCLGSYDCSNGDC

FVFGGGTEVVVKR.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 102)
QVLTQTASPVSPAVGSTVTINCRASQSVYYNNYLAWYQQKPGQPPKQLIY

STSTLASGVSSRFKGSGSGTQFTLTISDVQCDDAATYYCLGSYDCSNGDC

FVFGGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 103)
QSLEESGGRLVTPGGSLTLTCTVSGIDVTNYYMQWVRQAPGKGLEWIGVI

GVNGKRYYASWAKGRFTISKTSSTTVDLKMTSLTTEDTATYFCARGDIWG

PGTLVTVSS.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 104)
QSLEESGGRLVTPGGSLTLTCTVSGIDVTNYYMQWVRQAPGKGLEWIGVI

GVNGKRYYASWAKGRFTISKTSSTTVDLKMTSLTTEDTATYFCARGDIWG

PGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSW

NSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSN

TKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPE

VTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTV

LHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEM

TKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS

KLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 105; SEQ ID NO: 106; and SEQ ID NO: 107 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 101 or the light chain sequence of SEQ ID NO: 102, and/or one or more of the polypeptide sequences of SEQ ID NO: 108; SEQ ID NO: 109; and SEQ ID NO: 110 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 103 or the heavy chain sequence of SEQ ID NO: 104, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 101 or SEQ ID NO: 102. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 103 or SEQ ID NO: 104.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 105; SEQ ID NO: 106; and SEQ ID NO: 107 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 101 or the light chain sequence of SEQ ID NO: 102.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 108; SEQ ID NO: 109; and SEQ ID NO: 110 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 103 or the heavy chain sequence of SEQ ID NO: 104.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 101; the variable heavy chain region of SEQ ID NO: 103; the complementarity-determining regions (SEQ ID NO: 105; SEQ ID NO: 106; and SEQ ID NO: 107) of the variable light chain region of SEQ ID NO: 101; and the complementarity-determining regions (SEQ ID NO: 108; SEQ ID NO: 109; and SEQ ID NO: 110) of the variable heavy chain region of SEQ ID NO: 103.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody for potential treatment or prevention of photophobia is Ab11, comprising, or alternatively consisting of, SEQ ID NO: 102 and SEQ ID NO: 104, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab1, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 101 and the variable heavy chain sequence of SEQ ID NO: 103. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 101 and/or SEQ ID NO: 103 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab11. In another embodiment of the invention, anti-CGRP antibodies such as Ab11 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab12

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 111)
QVLTQSPSSLSASVGDRVTINCRASQSVYYNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSNGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 112)
QVLTQSPSSLSASVGDRVTINCRASQSVYYNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSNGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 113)
EVQLVESGGGLVQPGGSLRLSCAVSGIDVTNYYMQWVRQAPGKGLEWVGV

IGVNGKRYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 114)
EVQLVESGGGLVQPGGSLRLSCAVSGIDVTNYYMQWVRQAPGKGLEWVGV

IGVNGKRYYASWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCARGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 115; SEQ ID NO: 116; and SEQ ID NO: 117 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 111 or the light chain sequence of SEQ ID NO: 112, and/or one or more of the polypeptide sequences of SEQ ID NO: 118; SEQ ID NO: 119; and SEQ ID NO: 120 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 113 or the heavy chain sequence of SEQ ID NO: 114, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia

having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 111 or SEQ ID NO: 112. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 113 or SEQ ID NO: 114.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 115; SEQ ID NO: 116; and SEQ ID NO: 117 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 111 or the light chain sequence of SEQ ID NO: 112.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 118; SEQ ID NO: 119; and SEQ ID NO: 120 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 113 or the heavy chain sequence of SEQ ID NO: 114.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 111; the variable heavy chain region of SEQ ID NO: 113; the complementarity-determining regions (SEQ ID NO: 115; SEQ ID NO: 116; and SEQ ID NO: 117) of the variable light chain region of SEQ ID NO: 111; and the complementarity-determining regions (SEQ ID NO: 118; SEQ ID NO: 119; and SEQ ID NO: 120) of the variable heavy chain region of SEQ ID NO: 113.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab12, comprising, or alternatively consisting of, SEQ ID NO: 112 and SEQ ID NO: 114, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab12, the Fab fragment includes the variable light chain sequence of SEQ ID NO: 111 and the variable heavy chain sequence of SEQ ID NO: 113. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 111 and/or SEQ ID NO: 113 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab12. In another embodiment of the invention, anti-CGRP antibodies for potential treatment or prevention of photophobia such as Ab12 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab13

In one embodiment, the invention includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 121)
AIVMTQTPSSKSVPVGDTVTINCQASESLYNNNALAWFQQKPGQPPKRLI
YDASKLASGVPSRFSGGGSGTQFTLTISGVQCDDAATYYCGGYRSDSVDG
VAFAGGTEVVVKR.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 122)
AIVMTQTPSSKSVPVGDTVTINCQASESLYNNNALAWFQQKPGQPPKRLI
YDASKLASGVPSRFSGGGSGTQFTLTISGVQCDDAATYYCGGYRSDSVDG
VAFAGGTEVVVKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK
VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE
VTHQGLSSPVTKSFNRGEC.

The invention further includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 123)
QSVEESGGGLVQPEGSLTLTCTASGFDFSSNAMWWVRQAPGKGLEWIGII
YNGDGSTYYASWVNGRFSISKTSSTTVTLQLNSLTVADTATYYCARDLDL
WGPGTLVTVSS.

The invention also includes chimeric antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 124)
QSVEESGGGLVQPEGSLTLTCTASGFDFSSNAMWWVRQAPGKGLEWIGCI
YNGDGSTYYASWVNGRFSISKTSSTTVTLQLNSLTVADTATYYCARDLDL
WGPGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV
SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP
SNTKVDKRVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT
PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL
TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE
EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL
YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 125;

SEQ ID NO: 126; and SEQ ID NO: 127 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 121 or the light chain sequence of SEQ ID NO: 122, and/or one or more of the polypeptide sequences of SEQ ID NO: 128; SEQ ID NO: 129; and SEQ ID NO: 130 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 123 or the heavy chain sequence of SEQ ID NO: 124, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 121 or SEQ ID NO: 122. In another embodiment of the invention, antibody fragments of the invention comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 123 or SEQ ID NO: 124.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 125; SEQ ID NO: 126; and SEQ ID NO: 127 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 121 or the light chain sequence of SEQ ID NO: 122.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 128; SEQ ID NO: 129; and SEQ ID NO: 130 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 123 or the heavy chain sequence of SEQ ID NO: 124.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 121; the variable heavy chain region of SEQ ID NO: 123; the complementarity-determining regions (SEQ ID NO: 125; SEQ ID NO: 126; and SEQ ID NO: 127) of the variable light chain region of SEQ ID NO: 121; and the complementarity-determining regions (SEQ ID NO: 128; SEQ ID NO: 129; and SEQ ID NO: 130) of the variable heavy chain region of SEQ ID NO: 123.

In a particularly preferred embodiment of the invention, the chimeric anti-CGRP antibody for potential treatment or prevention of photophobia is Ab13, comprising, or alternatively consisting of, SEQ ID NO: 122 and SEQ ID NO: 124, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab13, the Fab fragment for potential treatment or prevention of photophobia includes the variable light chain sequence of SEQ ID NO: 121 and the variable heavy chain sequence of SEQ ID NO: 123. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 121 and/or SEQ ID NO: 123 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab13. In another embodiment of the invention, anti-CGRP antibodies for potential treatment or prevention of photophobia such as Ab13 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab14

In one embodiment, the invention includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 131)
QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSRGDC

FVFGGGTKVEIKR.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 132)
QVLTQSPSSLSASVGDRVTINCQASQNVYNNNYLAWYQQKPGKVPKQLIY

STSTLASGVPSRFSGSGSGTDFTLTISSLQPEDVATYYCLGSYDCSRGDC

FVFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAK

VQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACE

VTHQGLSSPVTKSFNRGEC.

The invention further includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a variable heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 133)
EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGV

IGSDGKTYYATWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCTRGDI

WGQGTLVTVSS.

The invention also includes humanized antibodies for potential treatment or prevention of photophobia having binding specificity to CGRP and possessing a heavy chain sequence comprising the sequence set forth below:

(SEQ ID NO: 134)
EVQLVESGGGLVQPGGSLRLSCAVSGIGLSSYYMQWVRQAPGKGLEWVGV

IGSDGKTYYATWAKGRFTISRDNSKTTVYLQMNSLRAEDTAVYFCTRGDI

WGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTV

SWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKP

SNTKVDARVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRT

PEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVL

TVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRE

EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFL

YSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

The invention further contemplates antibodies for potential treatment or prevention of photophobia comprising one or more of the polypeptide sequences of SEQ ID NO: 135; SEQ ID NO: 136; and SEQ ID NO: 137 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 131 or the light chain sequence of SEQ ID NO: 132, and/or one or more of the polypeptide sequences of SEQ ID NO: 138; SEQ ID NO: 139; and SEQ ID NO: 140 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 133 or the heavy chain sequence of SEQ ID NO: 134, or combinations of these polypeptide sequences. In another embodiment of the invention, the antibodies of the invention or fragments thereof for potential treatment or prevention of photophobia comprise, or alternatively consist of, combinations of one or more of the CDRs, the variable heavy and variable light chain sequences, and the heavy and light chain sequences set forth above, including all of them.

The invention also contemplates fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP. In one embodiment of the invention, antibody fragments of the invention for potential treatment or prevention of photophobia comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 131 or SEQ ID NO: 132. In another embodiment of the invention, antibody fragments of the invention for potential treatment or prevention of photophobia comprise, or alternatively consist of, the polypeptide sequence of SEQ ID NO: 133 or SEQ ID NO: 134.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 135; SEQ ID NO: 136; and SEQ ID NO: 137 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable light chain sequence of SEQ ID NO: 131 or the light chain sequence of SEQ ID NO: 132.

In a further embodiment of the invention, fragments of the antibody for potential treatment or prevention of photophobia having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polypeptide sequences of SEQ ID NO: 138; SEQ ID NO: 139; and SEQ ID NO: 140 which correspond to the complementarity-determining regions (CDRs, or hypervariable regions) of the variable heavy chain sequence of SEQ ID NO: 133 or the heavy chain sequence of SEQ ID NO: 134.

The invention also contemplates antibody fragments for potential treatment or prevention of photophobia which include one or more of the antibody fragments described herein. In one embodiment of the invention, fragments of the antibodies having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following antibody fragments: the variable light chain region of SEQ ID NO: 131; the variable heavy chain region of SEQ ID NO: 133; the complementarity-determining regions (SEQ ID NO: 135; SEQ ID NO: 136; and SEQ ID NO: 137) of the variable light chain region of SEQ ID NO: 131; and the complementarity-determining regions (SEQ ID NO: 138; SEQ ID NO: 139; and SEQ ID NO: 140) of the variable heavy chain region of SEQ ID NO: 133.

In a particularly preferred embodiment of the invention, the humanized anti-CGRP antibody for potential treatment or prevention of photophobia is Ab14, comprising, or alternatively consisting of, SEQ ID NO: 132 and SEQ ID NO: 134, and having at least one of the biological activities set forth herein.

In a further particularly preferred embodiment of the invention, antibody fragments for potential treatment or prevention of photophobia comprise, or alternatively consist of, Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab14, the Fab fragment for potential treatment or prevention of photophobia includes the variable light chain sequence of SEQ ID NO: 131 and the variable heavy chain sequence of SEQ ID NO: 133. This embodiment of the invention further contemplates additions, deletions, and variants of SEQ ID NO: 131 and/or SEQ ID NO: 133 in said Fab while retaining binding specificity for CGRP.

In one embodiment of the invention described herein (infra), Fab fragments for potential treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab14. In another embodiment of the invention, anti-CGRP antibodies for potential treatment or prevention of photophobia such as Ab14 or Fab fragments thereof may be produced via expression in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

In another embodiment, antibody fragments for potential treatment or prevention of photophobia may be present in one or more of the following non-limiting forms: Fab, Fab', $F(ab')_2$, Fv and single chain Fv antibody forms. In a preferred embodiment, the anti-CGRP antibodies for potential treatment or prevention of photophobia described herein further comprises the kappa constant light chain sequence comprising the sequence set forth below:

(SEQ ID NO: 283)
VAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGN

SQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKS

FNRGEC.

In another preferred embodiment, the anti-CGRP antibodies described herein for potential treatment or prevention of photophobia further comprises the gamma-1 constant heavy chain polypeptide sequence comprising the sequence set forth below:

(SEQ ID NO: 284)
ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGV

HTFPAVLQSSGLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEP

KSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVS

HEDPEVKFNWYVDGVEVHNAKTKPREEQYASTYRVVSVLTVLHQDWLNGK

EYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTC

LVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPGK.

In another embodiment, the invention contemplates an isolated anti-CGRP antibody for potential treatment or prevention of photophobia comprising a V_H polypeptide sequence selected from: SEQ ID NO: 3, 13, 23, 33, 43, 53, 63, 73, 83, 93, 103, 113, 123, or 133, or a variant thereof, and further comprising a V_L polypeptide sequence selected from: SEQ ID NO: 1, 11, 21, 31, 41, 51, 61, 71, 81, 91, 101, 111, 121, or 131, or a variant thereof, wherein one or more of the framework residues (FR residues) in said V_H or V_L polypeptide has been substituted with another amino acid residue resulting in an anti-CGRP antibody that specifically binds CGRP for potential treatment or prevention of photophobia. The invention contemplates humanized and chimeric forms of these antibodies. The chimeric antibodies for potential treatment or prevention of photophobia may include an Fc derived from IgG1, IgG2, IgG3, IgG4, IgG5, IgG6, IgG7, IgG8, IgG9, IgG10, IgG11, IgG12, IgG13, IgG14, IgG5, IgG16, IgG17, IgG18 or IgG19 constant regions.

In one embodiment of the invention, the antibodies or V_H or V_L polypeptides originate or are selected from one or more rabbit B cell populations prior to initiation of the humanization process referenced herein.

In another embodiment of the invention, the anti-CGRP antibodies and fragments thereof do not have binding specificity for CGRP-R. In a further embodiment of the invention, the anti-CGRP antibodies and fragments thereof inhibit the association of CGRP with CGRP-R. In another embodiment of the invention, the anti-CGRP antibodies and fragments thereof inhibit the association of CGRP with CGRP-R and/or additional proteins and/or multimers thereof, and/or antagonizes the biological effects thereof.

As stated herein, antibodies and fragments thereof may be modified post-translationally to add effector moieties such as chemical linkers, detectable moieties such as for example fluorescent dyes, enzymes, substrates, bioluminescent materials, radioactive materials, and chemiluminescent moieties, or functional moieties such as for example streptavidin, avidin, biotin, a cytotoxin, a cytotoxic agent, and radioactive materials.

Antibodies or fragments thereof may also be chemically modified to provide additional advantages such as increased solubility, stability and circulating time (in vivo half-life) of the polypeptide, or decreased immunogenicity (See U.S. Pat. No. 4,179,337). The chemical moieties for derivitization may be selected from water soluble polymers such as polyethylene glycol, ethylene glycol/propylene glycol copolymers, carboxymethylcellulose, dextran, polyvinyl alcohol and the like. The antibodies and fragments thereof may be modified at random positions within the molecule, or at predetermined positions within the molecule and may include one, two, three or more attached chemical moieties.

The polymer may be of any molecular weight, and may be branched or unbranched. For polyethylene glycol, the preferred molecular weight is between about 1 kDa and about 100 kDa (the term "about" indicating that in preparations of polyethylene glycol, some molecules will weigh more, some less, than the stated molecular weight) for ease in handling and manufacturing. Other sizes may be used, depending on the desired therapeutic profile (e.g., the duration of sustained release desired, the effects, if any on biological activity, the ease in handling, the degree or lack of antigenicity and other known effects of the polyethylene glycol to a therapeutic protein or analog). For example, the polyethylene glycol may have an average molecular weight of about 200, 500, 1000, 1500, 2000, 2500, 3000, 3500, 4000, 4500, 5000, 5500, 6000, 6500, 7000, 7500, 8000, 8500, 9000, 9500, 10,000, 10,500, 11,000, 11,500, 12,000, 12,500, 13,000, 13,500, 14,000, 14,500, 15,000, 15,500, 16,000, 16,500, 17,000, 17,500, 18,000, 18,500, 19,000, 19,500, 20,000, 25,000, 30,000, 35,000, 40,000, 50,000, 55,000, 60,000, 65,000, 70,000, 75,000, 80,000, 85,000, 90,000, 95,000, or 100,000 kDa. Branched polyethylene glycols are described, for example, in U.S. Pat. No. 5,643,575; Morpurgo et al., Appl. Biochem. Biotechnol. 56:59-72 (1996); Vorobjev et al., Nucleosides Nucleotides 18:2745-2750 (1999); and Caliceti et al., Bioconjug. Chem. 10:638-646 (1999), the disclosures of each of which are incorporated herein by reference.

There are a number of attachment methods available to those skilled in the art, See e.g., EP 0 401 384, herein incorporated by reference (coupling PEG to G-CSF), See also Malik et al., Exp. Hematol. 20:1028-1035 (1992) (reporting pegylation of GM-CSF using tresyl chloride). For example, polyethylene glycol may be covalently bound through amino acid residues via a reactive group, such as, a free amino or carboxyl group. Reactive groups are those to which an activated polyethylene glycol molecule may be bound. The amino acid residues having a free amino group may include lysine residues and the N-terminal amino acid residues; those having a free carboxyl group may include aspartic acid residues glutamic acid residues and the C-terminal amino acid residue. Sulfhydryl groups may also be used as a reactive group for attaching the polyethylene glycol molecules. Preferred for therapeutic purposes is attachment at an amino group, such as attachment at the N-terminus or lysine group.

As suggested above, polyethylene glycol may be attached to proteins via linkage to any of a number of amino acid residues. For example, polyethylene glycol can be linked to polypeptides via covalent bonds to lysine, histidine, aspartic acid, glutamic acid, or cysteine residues. One or more reaction chemistries may be employed to attach polyethylene glycol to specific amino acid residues (e.g., lysine, histidine, aspartic acid, glutamic acid, or cysteine) or to more than one type of amino acid residue (e.g., lysine, histidine, aspartic acid, glutamic acid, cysteine and combinations thereof).

Alternatively, antibodies or fragments thereof may have increased in vivo half lives via fusion with albumin (including but not limited to recombinant human serum albumin or fragments or variants thereof (See, e.g., U.S. Pat. No. 5,876,969, issued Mar. 2, 1999, EP Patent 0 413 622, and U.S. Pat. No. 5,766,883, issued Jun. 16, 1998, herein incorporated by reference in their entirety)) or other circulating blood proteins such as transferrin or ferritin. In a preferred embodiment, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) are fused with the mature form of human serum albumin (i.e., amino acids 1-585 of human serum albumin as shown in FIGS. 1 and 2 of EP Patent 0 322 094) which is herein incorporated

by reference in its entirety. Polynucleotides encoding fusion proteins of the invention are also encompassed by the invention.

Regarding detectable moieties, further exemplary enzymes include, but are not limited to, horseradish peroxidase, acetylcholinesterase, alkaline phosphatase, beta-galactosidase and luciferase. Further exemplary fluorescent materials include, but are not limited to, rhodamine, fluorescein, fluorescein isothiocyanate, umbelliferone, dichlorotriazinylamine, phycoerythrin and dansyl chloride. Further exemplary chemiluminescent moieties include, but are not limited to, luminol. Further exemplary bioluminescent materials include, but are not limited to, luciferin and aequorin. Further exemplary radioactive materials include, but are not limited to, Iodine 125 (^{125}I), Carbon 14 (^{14}C), Sulfur 35 (^{35}S), Tritium (^{3}H) and Phosphorus 32 (^{32}P).

Regarding functional moieties, exemplary cytotoxic agents include, but are not limited to, methotrexate, aminopterin, 6-mercaptopurine, 6-thioguanine, cytarabine, 5-fluorouracil decarbazine; alkylating agents such as mechlorethamine, thioepa chlorambucil, melphalan, carmustine (BSNU), mitomycin C, lomustine (CCNU), 1-methylnitrosourea, cyclothosphamide, mechlorethamine, busulfan, dibromomannitol, streptozotocin, mitomycin C, cis-dichlorodiamine platinum (II) (DDP) cisplatin and carboplatin (paraplatin); anthracyclines include daunorubicin (formerly daunomycin), doxorubicin (adriamycin), detorubicin, carminomycin, idarubicin, epirubicin, mitoxantrone and bisantrene; antibiotics include dactinomycin (actinomycin D), bleomycin, calicheamicin, mithramycin, and anthramycin (AMC); and antimytotic agents such as the vinca alkaloids, vincristine and vinblastine. Other cytotoxic agents include paclitaxel (taxol), ricin, *pseudomonas* exotoxin, gemcitabine, cytochalasin B, gramicidin D, ethidium bromide, emetine, etoposide, tenoposide, colchicin, dihydroxy anthracin dione, 1-dehydrotestosterone, glucocorticoids, procaine, tetracaine, lidocaine, propranolol, puromycin, procarbazine, hydroxyurea, asparaginase, corticosteroids, mytotane (O,P'-(DDD)), interferons, and mixtures of these cytotoxic agents.

Further cytotoxic agents include, but are not limited to, chemotherapeutic agents such as carboplatin, cisplatin, paclitaxel, gemcitabine, calicheamicin, doxorubicin, 5-fluorouracil, mitomycin C, actinomycin D, cyclophosphamide, vincristine and bleomycin. Toxic enzymes from plants and bacteria such as ricin, diphtheria toxin and *Pseudomonas* toxin may be conjugated to the humanized or chimeric antibodies, or binding fragments thereof, to generate cell-type-specific-killing reagents (Youle, et al., Proc. Nat'l Acad. Sci. USA 77:5483 (1980); Gilliland, et al., Proc. Nat'l Acad. Sci. USA 77:4539 (1980); Krolick, et al., Proc. Nat'l Acad. Sci. USA 77:5419 (1980)).

Other cytotoxic agents include cytotoxic ribonucleases as described by Goldenberg in U.S. Pat. No. 6,653,104. Embodiments of the invention also relate to radioimmunoconjugates where a radionuclide that emits alpha or beta particles is stably coupled to the antibody, or binding fragments thereof, with or without the use of a complex-forming agent. Such radionuclides include beta-emitters such as Phosphorus-32 (^{32}P), Scandium-47 (^{47}Sc), Copper-67 (^{67}Cu), Gallium-67 (^{67}Ga), Yttrium-88 (^{88}Y), Yttrium-90 (^{90}Y), Iodine-125 (^{125}I), Iodine-131 (^{131}I), Samarium-153 (^{153}Sm), Lutetium-177 (^{177}Lu), Rhenium-186 (^{186}Re) or Rhenium-188 (^{188}Re), and alpha-emitters such as Astatine-211 (^{211}At), Lead-212 (^{212}Pb), Bismuth-212 (^{212}Bi) or -213 (^{213}Bi) or Actinium-225 (^{225}Ac).

Methods are known in the art for conjugating an antibody or binding fragment thereof to a detectable moiety and the like, such as for example those methods described by Hunter et al, Nature 144:945 (1962); David et al, Biochemistry 13:1014 (1974); Pain et al, J. Immunol. Meth. 40:219 (1981); and Nygren, J., Histochem. and Cytochem. 30:407 (1982).

Embodiments described herein further include variants and equivalents that are substantially homologous to the antibodies, antibody fragments, diabodies, SMIPs, camelbodies, nanobodies, IgNAR, polypeptides, variable regions and CDRs set forth herein. These may contain, e.g., conservative substitution mutations, (i.e., the substitution of one or more amino acids by similar amino acids). For example, conservative substitution refers to the substitution of an amino acid with another within the same general class, e.g., one acidic amino acid with another acidic amino acid, one basic amino acid with another basic amino acid, or one neutral amino acid by another neutral amino acid. What is intended by a conservative amino acid substitution is well known in the art.

In another embodiment, the invention contemplates polypeptide sequences having at least 90% or greater sequence homology to any one or more of the polypeptide sequences of antibody fragments, variable regions and CDRs set forth herein. More preferably, the invention contemplates polypeptide sequences having at least 95% or greater sequence homology, even more preferably at least 98% or greater sequence homology, and still more preferably at least 99% or greater sequence homology to any one or more of the polypeptide sequences of antibody fragments, variable regions and CDRs set forth herein. Methods for determining homology between nucleic acid and amino acid sequences are well known to those of ordinary skill in the art.

In another embodiment, the invention further contemplates the above-recited polypeptide homologs of the antibody fragments, variable regions and CDRs set forth herein further having anti-CGRP activity. Non-limiting examples of anti-CGRP activity are set forth herein, for example, in paragraphs [0329]-[0350] infra.

In another embodiment, the invention further contemplates the generation and use of anti-idiotypic antibodies that bind any of the foregoing sequences. In an exemplary embodiment, such an anti-idiotypic antibody could be administered to a subject who has received an anti-CGRP antibody to modulate, reduce, or neutralize, the effect of the anti-CGRP antibody. Such anti-idiotypic antibodies could also be useful for treatment of an autoimmune disease characterized by the presence of anti-CGRP antibodies. A further exemplary use of such anti-idiotypic antibodies is for detection of the anti-CGRP antibodies of the present invention, for example to monitor the levels of the anti-CGRP antibodies present in a subject's blood or other bodily fluids.

The present invention also contemplates anti-CGRP antibodies for potential treatment or prevention of photophobia comprising any of the polypeptide or polynucleotide sequences described herein substituted for any of the other polynucleotide sequences described herein. For example, without limitation thereto, the present invention contemplates antibodies comprising the combination of any of the variable light chain and variable heavy chain sequences described herein, and further contemplates antibodies resulting from substitution of any of the CDR sequences described herein for any of the other CDR sequences described herein.

Additional Exemplary Embodiments of the Invention

In another embodiment, the invention contemplates one or more anti-human anti-CGRP antibodies or antibody frag-

ments thereof for potential treatment or prevention of photophobia which specifically bind to the same or overlapping linear or conformational epitope(s) and/or competes for binding to the same overlapping linear or conformational epitope(s) on an intact human CGRP polypeptide or fragment thereof as an anti-human CGRP antibody selected from Ab1, Ab2, Ab3, Ab4, Ab5, Ab6, Ab7, Ab8, Ab9, Ab10, Ab11, Ab12, Ab13, or Ab14. In a preferred embodiment, the anti-human CGRP antibody or fragment thereof specifically binds to the same overlapping linear or conformational epitope(s) and/or competes for binding to the same overlapping linear or conformational epitope(s) on an intact human CGRP polypeptide or a fragment thereof as Ab3, Ab6, Ab13, or Ab14, and most preferably Ab3.

A preferred embodiment of the invention is directed to chimeric or humanized antibodies and fragments thereof (including Fab fragments) having binding specificity for CGRP and inhibiting biological activities mediated by the binding of CGRP to the CGRP receptor especially for treatment or prevention of photophobia. In a particularly preferred embodiment of the invention, the chimeric or humanized anti-CGRP antibodies are selected from Ab3, Ab6, Ab13, or Ab14, or more preferably Ab3.

A preferred embodiment of the invention is directed to methods of screening antibodies and fragments thereof (including Fab fragments) having binding specificity to human Calcitonin Gene Related Peptide (hereinafter "CGRP") in animal models to determine the in vivo effects thereof, especially their ability to antagonize the adverse side effects of CGRP and to treat conditions involving excess CGRP especially their ability to treat or prevent photophobia, e.g., in migraine.

A more specific preferred embodiment of the invention involves a method of assessing the potential in vivo efficacy of a candidate CGRP/CGRP receptor inhibitor polypeptide, e.g., an anti-CGRP or anti-CGRP antibody or antibody fragment comprising determining whether the antibody inhibits light aversive behavior in a rodent administered CGRP compared to a rodent administered CGRP in the absence of the candidate anti-CGRP antibody or antibody fragment.

A more specific preferred embodiment of the invention involves a method of assessing the potential in vivo efficacy of a candidate anti-CGRP antibody or antibody fragment to treat a neurological condition characterized by increased CGRP levels and photophobia.

Another more specific preferred embodiment of the invention involves a method of assessing the potential in vivo efficacy of a candidate anti-CGRP antibody or antibody fragment to treat a CGRP associated disorder associated with photophobia such as migraine or chronic migraine, (with or without aura), or conditions such as weight loss, cancer or tumors, angiogenesis associated with cancer or tumor growth, angiogenesis associated with cancer or tumor survival, diarrhea, hemiplagic migraines, cluster headaches, migrainous neuralgia, chronic headaches, tension headaches, general headaches, hot flashes, chronic paroxysomal hemicrania, secondary headaches due to an underlying structural problem in the head or neck, cranial neuralgia, sinus headaches (such as for example associated with sinusitis), allergy-induced headaches or migraines, pain, headache-free migraine, abdominal migraine, inflammatory pain, post-operative incision pain, complex regional pain syndrome, cancer pain, primary or metastatic bone cancer pain, fracture pain, chronic pain, osteoporotic fracture pain, pain resulting from burn, osteoporosis, gout joint pain, abdominal pain, pain associated with sickle cell crises, and other nociceptic pain, as well as hepatocellular carcinoma, breast cancer, liver cirrhosis, menstrual pain, neurogenic pain, neuropathic pain, nociceptic pain, trigeminal neuralgia, post-herpetic neuralgia, phantom limb pain, fibromyalgia, menstrual pain, ovarialgia, reflex sympathetic dystrophy, neurogenic pain, osteoarthritis or rheumatoid arthritis pain, lower back pain, diabetic neuropathy, sciatica, or pain or visceral pain associated with: gastro-esophageal reflux, dyspepsia, irritable bowel syndrome, irritable colon, spastic colon, mucous colitis, inflammatory bowel disease, Crohn's disease, ileitis, ulcerative colitis, renal colic, dysmenorrhea, cystitis, menstrual period, labor, menopause, prostatitis, pancreatitis, renal colic, dysmenorrhea, cystitis, including interstitial cystitis (IC), surgery associated with the ileus, diverticulitis, peritonitis, pericarditis, hepatitis, appendicitis, colitis, cholecystitis, endometriosis, chronic and/or acute pancreatitis, myocardial infarction, kidney pain, pleural pain, prostatitis, pelvic pain, trauma to an organ, chronic nociceptive pain, chronic neuropathic pain, chronic inflammatory pain, fibromyalgia, breakthrough pain and persistent pain. Still another preferred embodiment of the invention involves a method of determining a suitable therapeutic dosage or dosage regimen of a candidate anti-CGRP antibody or antibody fragment in humans in order to treat a photophobia-associated condition selected from those identified herein based on the effects of said antibody or antibody fragment in a light aversive behavioral rodent animal model described in detail infra. Common causes of photophobia include migraine headaches, cataracts, or severe ophthalmologic diseases such as uveitis or corneal abrasion. A more extensive list of disorders associated with photophobia includes eye related causes such as Achromatopsia, Aniridia, Anticholinergic drugs may cause photophobia by paralyzing the iris sphincter muscle, Aphakia (absence of the lens of the eye), Buphthalmos (abnormally narrow angle between the cornea and iris), Cataracts, Cone dystrophy, Congenital abnormalities of the eye, Viral conjunctivitis ("pink eye") Corneal abrasion, Corneal dystrophy, Corneal ulcer, disruption of the corneal epithelium, such as that caused by a corneal foreign body or keratitis, Ectopia lentis, Endophthalmitis, Eye trauma caused by disease, injury, or infection such as chalazion, episcleritis, glaucoma, keratoconus, or optic nerve hypoplasia, Hydrophthalmos, or congenital glaucoma Iritis, Optic neuritis, Pigment dispersion syndrom, Pupillary dilation (naturally or chemically induced), Retinal detachment, Scarring of the cornea or sclera and Uveitis. In addition photophobia has nervous-system-related or urological causes including: Autism spectrum disorders, Chiari malformation, Dyslexia, Encephalitis including Myalgic encephalomyelitis aka Chronic fatigue syndrome, Meningitis, Subarachnoid haemorrhage, Tumor of the posterior cranial fossa, as well as other causes such as Ankylosing spondylitis, Albinism, Ariboflavinosis, Benzodiazepines (long term use of or withdrawal from benzodiazepines), Chemotherapy, Chikungunya, Cystinosis, Ehlers-Danlos syndrome, Hangover, Influenza, Infectious Mononucleosis, Magnesium deficiency, Mercury poisoning, Migraine, Rabies, and Tyrosinemia type II, also known as "Richner-Hanhart syndrome". Additionally it is known that photophobia is elevated in depression, bipolar disorder and agoraphobia.

Further another preferred embodiment of the invention relates to methods of assessing based on results in a rodent CGRP animal model a suitable therapeutic dosage or dosage regimen of the candidate anti-CGRP antibody or antibody fragment in humans.

Other preferred embodiments the present invention are directed to screening assays and therapeutic usage of specific antibodies and fragments thereof having binding specificity for CGRP for treatment or prevention of photophobia, in particular antibodies having desired epitopic specificity, high affinity or avidity and/or functional properties. In preferred embodiments this invention relates to assays and usage of the antibodies described herein, comprising the sequences of the V_H, V_L and CDR polypeptides described herein, and the polynucleotides encoding them. A preferred embodiment of the invention is directed to chimeric or humanized antibodies and fragments thereof (including Fab fragments) capable of binding to CGRP and/or inhibiting the biological activities mediated by the binding of CGRP to the CGRP receptor ("CGRP-R").

In a further embodiment of the invention is contemplated a method of reducing, treating or preventing diseases or disorders associated with CGRP by affecting those biological activities mediated via CGRP, especially inhibiting or preventing photophobia thereby avoiding the adverse biological activities mediated via binding of CGRP to CGRP-R. In one embodiment, the disease or disorder associated with photophobia is migraine, headache, pain, or other conditions aforementioned which are associated with photophobia. A further non-limiting listing of diseases and disorders associated with CGRP is provided herein.

Another preferred embodiment of the invention contemplates the use of Fab polypeptide sequences for the treatment of migraines and headaches and especially for treatment or prevention of photophobia in a patient. Non-limiting types of migraines and headaches that may be treated using Fab polypeptide sequences are provided elsewhere in this disclosure.

In another embodiment of the invention, the anti-human CGRP antibody for treatment or prevention of photophobia is an antibody which specifically binds to the same overlapping linear or conformational epitopes on an intact CGRP polypeptide or fragment thereof that is (are) specifically bound by Ab3, Ab6, Ab13, or Ab14 as ascertained by epitopic mapping using overlapping linear peptide fragments which span the full length of the native human CGRP polypeptide.

The invention is also directed to an anti-CGRP antibody for treatment or prevention of photophobia that binds with the same CGRP epitope and/or competes with an anti-CGRP antibody for binding to CGRP as an antibody or antibody fragment disclosed herein, including but not limited to an anti-CGRP antibody selected from Ab1, Ab2, Ab3, Ab4, Ab5, Ab6, Ab7, Ab8, Ab9, Ab10, Ab11, Ab12, Ab13, or Ab14, preferably Ab6, Ab10, Ab12, or Ab3. As mentioned, common causes of photophobia include migraine headaches, cataracts, or severe ophthalmologic diseases such as uveitis or corneal abrasion. A more extensive list of disorders associated with photophobia includes eye related causes such as Achromatopsia, Aniridia, Anticholinergic drugs may cause photophobia by paralyzing the iris sphincter muscle, Aphakia (absence of the lens of the eye), Buphthalmos (abnormally narrow angle between the cornea and iris), Cataracts, Cone dystrophy, Congenital abnormalities of the eye, Viral conjunctivitis ("pink eye") Corneal abrasion, Corneal dystrophy, Corneal ulcer, disruption of the corneal epithelium, such as that caused by a corneal foreign body or keratitis, Ectopia lentis, Endophthalmitis, Eye trauma caused by disease, injury, or infection such as chalazion, episcleritis, glaucoma, keratoconus, or optic nerve hypoplasia, Hydrophthalmos, or congenital glaucoma Iritis, Optic neuritis, Pigment dispersion syndrom, Pupillary dilation (naturally or chemically induced), Retinal detachment, Scarring of the cornea or sclera and Uveitis.

In addition photophobia has nervous-system-related or urological causes including: Autism spectrum disorders, Chiari malformation, Dyslexia, Encephalitis including Myalgic encephalomyelitis aka Chronic fatigue syndrome, Meningitis, Subarachnoid haemorrhage, Tumor of the posterior cranial fossa, as well as other causes such as Ankylosing spondylitis, Albinism, Ariboflavinosis, Benzodiazepines (long term use of or withdrawal from benzodiazepines), Chemotherapy, Chikungunya, Cystinosis, Ehlers-Danlos syndrome, Hangover, Influenza, Infectious Mononucleosis, Magnesium deficiency, Mercury poisoning, Migraine, Rabies, and Tyrosinemia type II, also known as "Richner-Hanhart syndrome". Additionally it is known that photophobia is elevated in depression, bipolar disorder and agoraphobia.

In another embodiment, the invention is also directed to an isolated anti-CGRP antibody or antibody fragment for treatment or prevention of photophobia comprising one or more of the CDRs contained in the V_H polypeptide sequences selected from: 3, 13, 23, 33, 43, 53, 63, 73, 83, 93, 103, 113, 123, or 133, or a variant thereof, and/or one or more of the CDRs contained in the V_L polypeptide sequences selected from: 1, 11, 21, 31, 41, 51, 61, 71, 81, 91, 101, 111, 121, or 131, or a variant thereof.

In one embodiment of the invention, the anti-human CGRP antibody for treatment or prevention of photophobia discussed in the two prior paragraphs comprises at least 2 complementarity determining regions (CDRs) in each the variable light and the variable heavy regions which are identical to those contained in an anti-human CGRP antibody selected from Ab1, Ab2, Ab3, Ab4, Ab5, Ab6, Ab7, Ab8, Ab9, Ab10, Ab11, Ab12, Ab13, or Ab14.

In a preferred embodiment, the anti-human CGRP antibody discussed above for treatment or prevention of photophobia comprises at least 2 complementarity determining regions (CDRs) in each the variable light and the variable heavy regions which are identical to those contained in Ab3 or Ab6. In another embodiment, all of the CDRs of the anti-human CGRP antibody discussed above are identical to the CDRs contained in an anti-human CGRP antibody selected from Ab1, Ab2, Ab3, Ab4, Ab5, Ab6, Ab7, Ab8, Ab9, Ab10, Ab11, Ab12, Ab13, or Ab14. In a preferred embodiment of the invention, all of the CDRs of the anti-human CGRP antibody discussed above are identical to the CDRs contained in an anti-human CGRP antibody selected from Ab3, Ab10, Ab12 or Ab6.

The invention further contemplates that the one or more anti-human CGRP antibodies discussed above for treatment or prevention of photophobia are aglycosylated or minimally glycosylated, e.g., lack N-glycosylation and comprise some O-glycosylation such as some 1 or more mannose residues; e.g., that contain an Fc region that has been modified to alter effector function, half-life, proteolysis, and/or glycosylation; are human, humanized, single chain or chimeric; and are a humanized antibody derived from a rabbit (parent) anti-human CGRP antibody.

The invention further contemplates one or more anti-human CGRP antibodies for treatment or prevention of photophobia wherein the framework regions (FRs) in the variable light region and the variable heavy regions of said antibody respectively are human FRs which are unmodified or which have been modified by the substitution of one or more human FR residues in the variable light or heavy chain region with the corresponding FR residues of the parent rabbit antibody, and wherein said human FRs have been

derived from human variable heavy and light chain antibody sequences which have been selected from a library of human germline antibody sequences based on their high level of homology to the corresponding rabbit variable heavy or light chain regions relative to other human germline antibody sequences contained in the library.

In one embodiment of the invention, the anti-human CGRP antibody or fragment for treatment or prevention of photophobia specifically binds to CGRP expressing human cells and/or to circulating soluble CGRP molecules in vivo, including CGRP expressed on or by human cells in a patient with a disease associated with cells that express CGRP.

In another embodiment, the disease is selected from photophobia or light aversion associated with one or more of: migraines (with or without aura), menstrual headache, menstrual migraine, menopausal migraine or another hormonally related migraine, hemiplegic migraines, cluster headaches, migrainous neuralgia, chronic headaches, tension headaches, general headaches, migraines associated with hot flashes, chronic paroxysomal hemicrania, secondary headaches due to an underlying structural problem in the head or neck, cranial neuralgia, sinus headaches (such as for example associated with sinusitis), allergy-induced headaches or migraines, headache-free migraine, and abdominal migraine.

The invention further contemplates anti-human CGRP antibodies or fragments for treatment or prevention of photophobia directly or indirectly attached to a detectable label or therapeutic agent.

The invention also contemplates one or more nucleic acid sequences which result in the expression of an anti-human CGRP antibody or antibody fragment for treatment or prevention of photophobia as set forth above, including those comprising, or alternatively consisting of, yeast or human preferred codons. The invention also contemplates vectors (including plasmids or recombinant viral vectors) comprising said nucleic acid sequence(s). The invention also contemplates host cells or recombinant host cells expressing at least one of the antibodies set forth above, including a mammalian, yeast, bacterial, and insect cells. In a preferred embodiment, the host cell is a yeast cell. In a further preferred embodiment, the yeast cell is a diploidal yeast cell. In a more preferred embodiment, the yeast cell is a *Pichia* yeast.

The invention also contemplates a method of treatment comprising administering to a patient with a disease or condition associated with CGRP expressing cells a therapeutically effective amount of at least one anti-human CGRP antibody or fragment described herein for treatment or prevention of photophobia. The invention also contemplates that the treatment method may involve the administration of two or more anti-CGRP antibodies or fragments thereof and disclosed herein. If more than one antibody is administered to the patient, the multiple antibodies may be administered simultaneously or concurrently, or may be staggered in their administration. The diseases that may be treated are presented in the non-limiting list set forth above and elsewhere herein. In a preferred embodiment, the disease associated with photophobia is selected from migraine, headache, pain, diarrhea, cancer pain or neuropathic pain. In another embodiment the treatment further includes the administration of another therapeutic agent or regimen selected from chemotherapy, radiotherapy, cytokine administration or gene therapy.

In a non-limiting embodiment of the invention, another therapeutic agent or regimen includes opioids, analgesics such as NSAIDs, Taxol (paclitaxel) or its derivatives, platinum compounds such as carboplatin or cisplatin, anthrocyclines such as doxorubicin, alkylating agents such as cyclophosphamide, anti-metabolites such as 5-fluorouracil, or etoposide.

The invention further contemplates a method of in vivo imaging which detects the presence of cells which express CGRP comprising administering a diagnostically effective amount of at least one anti-human CGRP antibody. In one embodiment, said administration further includes the administration of a radionuclide or fluorophore that facilitates detection of the antibody at CGRP expressing disease sites. In a further embodiment, the results of said in vivo imaging method are used to facilitate the design of an appropriate therapeutic regimen, including therapeutic regimens including radiotherapy, chemotherapy or a combination thereof.

The anti-CGRP activity of the anti-CGRP antibodies of the present invention, and fragments thereof having binding specificity to CGRP for treatment or prevention of photophobia, may also be described by their strength of binding or their affinity for CGRP. In one embodiment of the invention, the anti-CGRP antibodies of the present invention, and fragments thereof having binding specificity to CGRP, bind to CGRP with a dissociation constant (K_D) of less than or equal to 5×10^{-7} M, 10^{-7} M, 5×10^{-8} M, 10^{-8} M, 5×10^{-9} M, 10^{-9} M, 5×10^{-10} M, 10^{-10} M, 5×10^{-11} M, 10^{-11} M, 5×10^{-12} M, 10^{-12} M, 5×10^{-13} M, or 10^{-13} M. Preferably, the anti-CGRP antibodies and fragments thereof bind CGRP with a dissociation constant of less than or equal to 10^{-11} M, 5×10^{-12} M, or 10^{-12} M. In another embodiment of the invention, the anti-CGRP antibodies of the present invention, and fragments thereof having binding specificity to CGRP, bind to a linear or conformational CGRP epitope.

In another embodiment of the invention, the anti-CGRP activity of the anti-CGRP antibodies of the present invention, and fragments thereof having binding specificity to CGRP, bind to CGRP with an off-rate of less than or equal to 10^{-4} S^{-1}, 5×10^{-5} S^{-1}, 10^{-5} S^{-1}, 5×10^{-6} S^{-1}, 10^{-6} S^{-1}, 5×10^{-7} S^{-1}, or 10^{-7} S^{1}.

In a further embodiment of the invention, the anti-CGRP activity of the anti-CGRP antibodies of the present invention, and fragments thereof having binding specificity to CGRP, exhibit anti-CGRP activity by preventing, ameliorating or reducing the symptoms of, or alternatively treating, diseases and disorders associated with CGRP especially for treatment or prevention of photophobia. Non-limiting examples of diseases and disorders associated with CGRP and conditions associated with photophobia are set forth herein.

Polynucleotides Encoding Anti-CGRP Antibody Polypeptides

Antibody Ab1

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 1:

(SEQ ID NO: 141)

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAG

CACAGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATGATAACAACT

ACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATC

TATTCTACATCCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAG

-continued

TGGATCTGGGACACAGTTCACTCTCACCATCAGCGACCTGGAGTGTGCCG

ATGCTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGAT

TGTTTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 2:

(SEQ ID NO: 142)

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAG

CACAGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATGATAACAACT

ACCTAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATC

TATTCTACATCCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAG

TGGATCTGGGACACAGTTCACTCTCACCATCAGCGACCTGGAGTGTGCCG

ATGCTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGAT

TGTTTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGTACGGTGGC

TGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTG

GAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCC

AAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGA

GAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCA

CCCTGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGC

GAAGTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAG

GGGAGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 3:

(SEQ ID NO: 143)

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACC

CCTGACACTCACCTGCACAGTCTCTGGACTCGACCTCAGTAGCTACTACA

TGCAATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTC

ATTGGTATTAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGCCTCGTCGACCACGGTGGATCTGAAAATGACCAGTC

TGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGG

GGCCCAGGCACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 4:

(SEQ ID NO: 144)

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCT

GACACTCACCTGCACAGTCTCTGGACTCGACCTCAGTAGCTACTACATGC

AATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATT

GGTATTAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATTCAC

-continued

CATCTCCAGAGCCTCGTCGACCACGGTGGATCTGAAAATGACCAGTCTGA

CAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGGGGC

CCAGGCACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGT

CTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAGCGGCCC

TGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGG

AACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACA

GTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCA

GCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAAC

ACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAACTCACAC

ATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC

TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAG

GTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAAGTT

CAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGC

GGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTC

CTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAA

CAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGC

AGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATG

ACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCAG

CGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACA

AGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGC

AAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATG

CTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCT

CCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 145; SEQ ID NO: 146; and SEQ ID NO: 147 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 1 or the light chain sequence of SEQ ID NO: 2.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 148; SEQ ID NO: 149; and SEQ ID NO: 150 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 3 or the heavy chain sequence of SEQ ID NO: 4.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 141 encoding the light chain variable sequence of SEQ ID NO: 1; the polynucleotide SEQ ID NO: 142 encoding the light

chain sequence of SEQ ID NO: 2; the polynucleotide SEQ ID NO: 143 encoding the heavy chain variable sequence of SEQ ID NO: 3; the polynucleotide SEQ ID NO: 144 encoding the heavy chain sequence of SEQ ID NO: 4; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 145; SEQ ID NO: 146; and SEQ ID NO: 147) of the light chain variable sequence of SEQ ID NO: 1 or the light chain sequence of SEQ ID NO: 2; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 148; SEQ ID NO: 149; and SEQ ID NO: 150) of the heavy chain variable sequence of SEQ ID NO: 3 or the heavy chain sequence of SEQ ID NO: 4.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab1, the polynucleotides encoding the full length Ab1 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 142 encoding the light chain sequence of SEQ ID NO: 2 and the polynucleotide SEQ ID NO: 144 encoding the heavy chain sequence of SEQ ID NO: 4.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab1 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab1 or Fab fragments thereof may be produced via expression of Ab1 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab2

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 11:

(SEQ ID NO: 151)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGAC

AGAGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATGATAACAACTA

CCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCT

ATTCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGT

GGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGA

TGTTGCAACTTATTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATT

GTTTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 12:

(SEQ ID NO: 152)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGAC

AGAGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATGATAACAACTA

CCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCT

ATTCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGT

GGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGA

TGTTGCAACTTATTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATT

GTTTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCT

GCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGG

AACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCA

AAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAG

AGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCAC

CCTGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCG

AAGTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGG

GGAGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO:

(SEQ ID NO: 153)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGG

GTCCCTGAGACTCTCCTGTGCAGTCTCTGGACTCGACCTCAGTAGCTACT

ACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGA

GTCATTGGTATCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCG

ATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGA

ACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGAC

ATCTGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 14:

(SEQ ID NO: 154)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGACTCGACCTCAGTAGCTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTATCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

-continued

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 155; SEQ ID NO: 156; and SEQ ID NO: 157 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 11 or the light chain sequence of SEQ ID NO: 12.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 158; SEQ ID NO: 159; and SEQ ID NO: 160 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 13 or the heavy chain sequence of SEQ ID NO: 14.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 151 encoding the light chain variable sequence of SEQ ID NO: 11; the polynucleotide SEQ ID NO: 152 encoding the light chain sequence of SEQ ID NO: 12; the polynucleotide SEQ ID NO: 153 encoding the heavy chain variable sequence of SEQ ID NO: 13; the polynucleotide SEQ ID NO: 154 encoding the heavy chain sequence of SEQ ID NO: 14; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 155; SEQ ID NO: 156; and SEQ ID NO: 157) of the light chain variable sequence of SEQ ID NO: 11 or the light chain sequence of SEQ ID NO: 12; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 158; SEQ ID NO: 159; and SEQ ID NO: 160) of the heavy chain variable sequence of SEQ ID NO: 13 or the heavy chain sequence of SEQ ID NO: 14.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab2, the polynucleotides encoding the full length Ab2 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 152 encoding the light chain sequence of SEQ ID NO: 12 and the polynucleotide SEQ ID NO: 154 encoding the heavy chain sequence of SEQ ID NO: 14.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab2 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab2 or Fab fragments thereof may be produced via expression of Ab2 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab3

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 21:

(SEQ ID NO: 161)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGAC

AGAGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATGATAACAACTA

CCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCT

ATTCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGT

GGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGA

TGTTGCAACTTATTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATT

GTTTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 22:

(SEQ ID NO: 162)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGAC

AGAGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATGATAACAACTA

CCTAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCT

ATTCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGT

GGATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGA

TGTTGCAACTTATTACTGTCTAGGCAGTTATGATTGTAGTAGTGGTGATT

GTTTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCT

-continued

GCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGG

AACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCA

AAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAG

AGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCAC

CCTGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCG

AAGTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGG

GGAGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 23:

(SEQ ID NO: 163)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGG

GTCCCTGAGACTCTCCTGTGCAGTCTCTGGACTCGACCTCAGTAGCTACT

ACATGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGA

GTCATTGGTATCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCG

ATTCACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGA

ACAGCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGAC

ATCTGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 24:

(SEQ ID NO: 164)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGACTCGACCTCAGTAGCTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTATCAATGATAACACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACGCGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

-continued

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 165; SEQ ID NO: 166; and SEQ ID NO: 167 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 21 or the light chain sequence of SEQ ID NO: 22.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 168; SEQ ID NO: 169; and SEQ ID NO: 170 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 23 or the heavy chain sequence of SEQ ID NO: 24.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 161 encoding the light chain variable sequence of SEQ ID NO: 21; the polynucleotide SEQ ID NO: 162 encoding the light chain sequence of SEQ ID NO: 22; the polynucleotide SEQ ID NO: 163 encoding the heavy chain variable sequence of SEQ ID NO: 23; the polynucleotide SEQ ID NO: 164 encoding the heavy chain sequence of SEQ ID NO: 24; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 165; SEQ ID NO: 166; and SEQ ID NO: 167) of the light chain variable sequence of SEQ ID NO: 21 or the light chain sequence of SEQ ID NO: 22; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 168; SEQ ID NO: 169; and SEQ ID NO: 170) of the heavy chain variable sequence of SEQ ID NO: 23 or the heavy chain sequence of SEQ ID NO: 24.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab3, the polynucleotides encoding the full length Ab3 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 162 encoding the light chain sequence of SEQ ID NO: 22 and the polynucleotide SEQ ID NO: 164 encoding the heavy chain sequence of SEQ ID NO: 24.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for

expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments for treatment or prevention of photophobia may be produced by enzymatic digestion (e.g., papain) of Ab3 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies for treatment or prevention of photophobia such as Ab3 or Fab fragments thereof may be produced via expression of Ab3 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab4

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 31:

(SEQ ID NO: 171)

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCAC

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATCATAACACCTACC

TGGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAACAACTGATCTAT

GATGCATCCACTCTGGCGTCTGGGGTCCCATCGCGGTTCAGCGGCAGTGG

ATCTGGGACACAGTTCACTCTCACCATCAGCGGCGTGCAGTGTAACGATG

CTGCCGCTTACTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGT

TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 32:

(SEQ ID NO: 172)

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCAC

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATCATAACACCTACC

TGGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAACAACTGATCTAT

GATGCATCCACTCTGGCGTCTGGGGTCCCATCGCGGTTCAGCGGCAGTGG

ATCTGGGACACAGTTCACTCTCACCATCAGCGGCGTGCAGTGTAACGATG

CTGCCGCTTACTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGT

TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

-continued

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 33:

(SEQ ID NO: 173)

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCT

GACACTCACCTGTTCCGTCTCTGGCATCGACCTCAGTGGCTACTACATGA

ACTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATT

GGTATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCAC

CATCTCCAAAACCTCGTCGACCACGGTGGATCTGAAAATGACCAGTCTGA

CAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGGGGC

CCGGGCACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 34:

(SEQ ID NO: 174)

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCT

GACACTCACCTGTTCCGTCTCTGGCATCGACCTCAGTGGCTACTACATGA

ACTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATT

GGTATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATTCAC

CATCTCCAAAACCTCGTCGACCACGGTGGATCTGAAAATGACCAGTCTGA

CAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGGGGC

CCGGGCACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGT

CTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAGCGGCCC

TGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGG

AACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACA

GTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCA

GCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAAC

ACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAACTCACAC

ATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC

TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAG

GTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAAGTT

CAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGC

GGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTC

CTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAA

CAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGC

AGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATG

ACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCAG

-continued

CGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACA

AGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGC

AAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATG

CTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCT

CCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 175; SEQ ID NO: 176; and SEQ ID NO: 177 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 31 or the light chain sequence of SEQ ID NO: 32.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 178; SEQ ID NO: 179; and SEQ ID NO: 180 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 33 or the heavy chain sequence of SEQ ID NO: 34.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 171 encoding the light chain variable sequence of SEQ ID NO: 31; the polynucleotide SEQ ID NO: 172 encoding the light chain sequence of SEQ ID NO: 32; the polynucleotide SEQ ID NO: 173 encoding the heavy chain variable sequence of SEQ ID NO: 33; the polynucleotide SEQ ID NO: 174 encoding the heavy chain sequence of SEQ ID NO: 34; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 175; SEQ ID NO: 176; and SEQ ID NO: 177) of the light chain variable sequence of SEQ ID NO: 31 or the light chain sequence of SEQ ID NO: 32; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 178; SEQ ID NO: 179; and SEQ ID NO: 180) of the heavy chain variable sequence of SEQ ID NO: 33 or the heavy chain sequence of SEQ ID NO: 34.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab4, the polynucleotides encoding the full length Ab4 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 172 encoding the light chain sequence of SEQ ID NO: 32 and the polynucleotide SEQ ID NO: 174 encoding the heavy chain sequence of SEQ ID NO: 34.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab4 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab4 or Fab fragments thereof may be produced via expression of Ab4 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab5

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 41:

(SEQ ID NO: 181)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATCATAACACCTACC

TGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

GATGCATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 42:

(SEQ ID NO: 182)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATCATAACACCTACC

TGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

GATGCATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 43:

(SEQ ID NO: 183)
GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACCTCAGTGGCTACTACA

TGAACTGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 44:

(SEQ ID NO: 184)
GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACCTCAGTGGCTACTACA

TGAACTGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 185; SEQ ID NO: 186; and SEQ ID NO: 187 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 41 or the light chain sequence of SEQ ID NO: 42.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 188; SEQ ID NO: 189; and SEQ ID NO: 190 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 43 or the heavy chain sequence of SEQ ID NO: 44.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 181 encoding the light chain variable sequence of SEQ ID NO: 41; the polynucleotide SEQ ID NO: 182 encoding the light chain sequence of SEQ ID NO: 42; the polynucleotide SEQ ID NO: 183 encoding the heavy chain variable sequence of SEQ ID NO: 43; the polynucleotide SEQ ID NO: 184 encoding the heavy chain sequence of SEQ ID NO: 44; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 185; SEQ ID NO: 186; and SEQ ID NO: 187) of the light chain variable sequence of SEQ ID NO: 41 or the light chain sequence of SEQ ID NO: 42; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 188; SEQ ID NO: 189; and SEQ ID NO: 190) of the heavy chain variable sequence of SEQ ID NO: 43 or the heavy chain sequence of SEQ ID NO: 44.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab5, the polynucleotides encoding the full length Ab5 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 182 encoding the light chain sequence of SEQ ID NO: 42 and the polynucleotide SEQ ID NO: 184 encoding the heavy chain sequence of SEQ ID NO: 44.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab5 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab5 or Fab fragments thereof may be produced via expression of Ab5 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab6

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucle-

otides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 51:

(SEQ ID NO: 191)
CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG
AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATCATAACACCTACC
TGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT
GATGCATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG
ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG
TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGT
TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 52:

(SEQ ID NO: 192)
CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG
AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATCATAACACCTACC
TGGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT
GATGCATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG
ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG
TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTACTAATGGTGATTGT
TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCTGC
ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA
CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA
GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG
TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC
TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA
GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG
AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 53:

(SEQ ID NO: 193)
GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC
CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACCTCAGTGGCTACTACA
TGAACTGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC
ATTGGTATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATT
CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA
GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC
TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 54:

(SEQ ID NO: 194)
GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC
CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACCTCAGTGGCTACTACA
TGAACTGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC
ATTGGTATTAATGGTGCCACATACTACGCGAGCTGGGCGAAAGGCCGATT
CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA
GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC
TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC
ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG
CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG
TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT
CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT
CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC
AGCAACACCAAGGTGGACGCGAGAGTTGAGCCCAAATCTTGTGACAAAAC
TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG
TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC
CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT
CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA
AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC
ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT
CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA
AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG
GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA
TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA
ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC
TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT
CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA
GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 195; SEQ ID NO: 196; and SEQ ID NO: 197 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 51 or the light chain sequence of SEQ ID NO: 52.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 198; SEQ ID NO: 199; and SEQ ID NO: 200 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 53 or the heavy chain sequence of SEQ ID NO: 54.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding

antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 191 encoding the light chain variable sequence of SEQ ID NO: 51; the polynucleotide SEQ ID NO: 192 encoding the light chain sequence of SEQ ID NO: 52; the polynucleotide SEQ ID NO: 193 encoding the heavy chain variable sequence of SEQ ID NO: 53; the polynucleotide SEQ ID NO: 194 encoding the heavy chain sequence of SEQ ID NO: 54; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 195; SEQ ID NO: 196; and SEQ ID NO: 197) of the light chain variable sequence of SEQ ID NO: 51 or the light chain sequence of SEQ ID NO: 52; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 198; SEQ ID NO: 199; and SEQ ID NO: 200) of the heavy chain variable sequence of SEQ ID NO: 53 or the heavy chain sequence of SEQ ID NO: 54.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab6, the polynucleotides encoding the full length Ab6 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 192 encoding the light chain sequence of SEQ ID NO: 52 and the polynucleotide SEQ ID NO: 194 encoding the heavy chain sequence of SEQ ID NO: 54.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab6 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab6 or Fab fragments thereof may be produced via expression of Ab6 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab7

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 61:

(SEQ ID NO: 201)

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAGCAC

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATAATTACAACTACC

TTGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCTCATCGCGATTCAAAGGCAGTGG

ATCTGGGACACAGTTCACTCTCACCATCAGCGACGTGCAGTGTGACGATG

CTGCCACTTACTACTGTCTAGGCAGTTATGACTGTAGTACTGGTGATTGT

TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 62:

(SEQ ID NO: 202)

CAAGTGCTGACCCAGACTGCATCCCCCGTGTCTGCAGCTGTGGGAAGCAC

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTATAATTACAACTACC

TTGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCTCATCGCGATTCAAAGGCAGTGG

ATCTGGGACACAGTTCACTCTCACCATCAGCGACGTGCAGTGTGACGATG

CTGCCACTTACTACTGTCTAGGCAGTTATGACTGTAGTACTGGTGATTGT

TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 63:

(SEQ ID NO: 203)

CAGGAGCAGCTGAAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACATC

CCTGACACTCACCTGCACCGTCTCTGGAATCGACCTCAGTAACCACTACA

TGCAATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGAGTC

GTTGGTATTAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAACCTCGTCGACCACGGTGGATCTGAAAATGACCAGGC

TGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGG

GGCCCAGGCACCCTGGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 64:

(SEQ ID NO: 204)

CAGGAGCAGCTGAAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACATC

CCTGACACTCACCTGCACCGTCTCTGGAATCGACCTCAGTAACCACTACA

TGCAATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGAGTC

GTTGGTATTAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAACCTCGTCGACCACGGTGGATCTGAAAATGACCAGGC

TGACAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGGGACATCTGG

GGCCCAGGCACCCTGGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATC

GGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAGCGG

-continued

CCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCG

TGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCCT

ACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCA

GCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGC

AACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAACTCA

CACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCT

TCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCT

GAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAA

GTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGC

CGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACC

GTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTC

CAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAG

GGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAGGAG

ATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCC

CAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACT

ACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTAC

AGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTC

ATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCC

TCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 205; SEQ ID NO: 206; and SEQ ID NO: 207 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 61 or the light chain sequence of SEQ ID NO: 62.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 208; SEQ ID NO: 209; and SEQ ID NO: 210 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 63 or the heavy chain sequence of SEQ ID NO: 64.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 201 encoding the light chain variable sequence of SEQ ID NO: 61; the polynucleotide SEQ ID NO: 202 encoding the light chain sequence of SEQ ID NO: 62; the polynucleotide SEQ ID NO: 203 encoding the heavy chain variable sequence of SEQ ID NO: 63; the polynucleotide SEQ ID NO: 204 encoding the heavy chain sequence of SEQ ID NO: 64; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 205; SEQ ID NO: 206; and SEQ ID NO: 207) of the light chain variable sequence of SEQ ID NO: 61 or the light chain sequence of SEQ ID NO: 62; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 208; SEQ ID NO: 209; and SEQ ID NO: 210) of the heavy chain variable sequence of SEQ ID NO: 63 or the heavy chain sequence of SEQ ID NO: 64.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab7, the polynucleotides encoding the full length Ab7 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 202 encoding the light chain sequence of SEQ ID NO: 62 and the polynucleotide SEQ ID NO: 204 encoding the heavy chain sequence of SEQ ID NO: 64.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab7 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab7 or Fab fragments thereof may be produced via expression of Ab7 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab8

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 71:

(SEQ ID NO: 211)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTACAATTACAACTACC

TTGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTACTGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 72:

(SEQ ID NO: 212)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAGTGTTTACAATTACAACTACC

TTGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

-continued

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTACTGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 73:

(SEQ ID NO: 213)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACCTCAGTAACCACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

GTTGGTATCAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 74:

(SEQ ID NO: 214)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACCTCAGTAACCACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

GTTGGTATCAATGGTCGCACATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCTAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

-continued

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 215; SEQ ID NO: 216; and SEQ ID NO: 217 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 71 or the light chain sequence of SEQ ID NO: 72.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 218; SEQ ID NO: 219; and SEQ ID NO: 220 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 73 or the heavy chain sequence of SEQ ID NO: 74.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 211 encoding the light chain variable sequence of SEQ ID NO: 71; the polynucleotide SEQ ID NO: 212 encoding the light chain sequence of SEQ ID NO: 72; the polynucleotide SEQ ID NO: 213 encoding the heavy chain variable sequence of SEQ ID NO: 73; the polynucleotide SEQ ID NO: 214 encoding the heavy chain sequence of SEQ ID NO: 74; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 215; SEQ ID NO: 216; and SEQ ID NO: 217) of the light chain variable sequence of SEQ ID NO: 71 or the light chain sequence of SEQ ID NO: 72; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 218; SEQ ID NO: 219; and SEQ ID NO: 220) of the heavy chain variable sequence of SEQ ID NO: 73 or the heavy chain sequence of SEQ ID NO: 74.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab8, the polynucleotides encoding the full length Ab8 antibody comprise, or alternatively consist of,

the polynucleotide SEQ ID NO: 212 encoding the light chain sequence of SEQ ID NO: 72 and the polynucleotide SEQ ID NO: 214 encoding the heavy chain sequence of SEQ ID NO: 74.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab8 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab8 or Fab fragments thereof may be produced via expression of Ab8 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab9

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 81:

(SEQ ID NO: 221)

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCAC

AGTCACCATCAATTGCCAGGCCAGTCAGAATGTTTATAATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTAT

TCTACGTCCACTCTGGCATCTGGGGTCTCATCGCGATTCAGAGGCAGTGG

ATCTGGGACACAGTTCACTCTCACCATCAGCGACGTGCAGTGTGACGATG

CTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTCGTGGTGATTGT

TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 82:

(SEQ ID NO: 222)

CAAGTGCTGACCCAGACTCCATCCCCCGTGTCTGCAGCTGTGGGAAGCAC

AGTCACCATCAATTGCCAGGCCAGTCAGAATGTTTATAATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTAT

TCTACGTCCACTCTGGCATCTGGGGTCTCATCGCGATTCAGAGGCAGTGG

ATCTGGGACACAGTTCACTCTCACCATCAGCGACGTGCAGTGTGACGATG

CTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTCGTGGTGATTGT

TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

-continued

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 83:

(SEQ ID NO: 223)

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCT

GACACTCACCTGCACAGTCTCTGGAATCGGCCTCAGTAGCTACTACATGC

AGTGGGTCCGCCAGTCTCCAGGGAGGGGGCTGGAATGGATCGGAGTCATT

GGTAGTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCAC

CATCTCCAAGACCTCGTCGACCACGGTGGATCTGAGAATGGCCAGTCTGA

CAACCGAGGACACGGCCACCTATTTCTGTACCAGAGGGGACATCTGGGGC

CCGGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 84:

(SEQ ID NO: 224)

CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGGACACCCCT

GACACTCACCTGCACAGTCTCTGGAATCGGCCTCAGTAGCTACTACATGC

AGTGGGTCCGCCAGTCTCCAGGGAGGGGGCTGGAATGGATCGGAGTCATT

GGTAGTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATTCAC

CATCTCCAAGACCTCGTCGACCACGGTGGATCTGAGAATGGCCAGTCTGA

CAACCGAGGACACGGCCACCTATTTCTGTACCAGAGGGGACATCTGGGGC

CCGGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGT

CTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAGCGGCCC

TGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGG

AACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACA

GTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCA

GCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAAC

ACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAACTCACAC

ATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC

TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAG

GTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAAGTT

CAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGC

GGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTC

CTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAA

CAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGC

AGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATG

ACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCAG

-continued

CGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACA

AGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGC

AAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATG

CTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCT

CCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 225; SEQ ID NO: 226; and SEQ ID NO: 227 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 81 or the light chain sequence of SEQ ID NO: 82.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 228; SEQ ID NO: 229; and SEQ ID NO: 230 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 83 or the heavy chain sequence of SEQ ID NO: 84.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 221 encoding the light chain variable sequence of SEQ ID NO: 81; the polynucleotide SEQ ID NO: 222 encoding the light chain sequence of SEQ ID NO: 82; the polynucleotide SEQ ID NO: 223 encoding the heavy chain variable sequence of SEQ ID NO: 83; the polynucleotide SEQ ID NO: 224 encoding the heavy chain sequence of SEQ ID NO: 84; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 225; SEQ ID NO: 226; and SEQ ID NO: 227) of the light chain variable sequence of SEQ ID NO: 81 or the light chain sequence of SEQ ID NO: 82; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 228; SEQ ID NO: 229; and SEQ ID NO: 230) of the heavy chain variable sequence of SEQ ID NO: 83 or the heavy chain sequence of SEQ ID NO: 84.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab9, the polynucleotides encoding the full length Ab9 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 222 encoding the light chain sequence of SEQ ID NO: 82 and the polynucleotide SEQ ID NO: 224 encoding the heavy chain sequence of SEQ ID NO: 84.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab9 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab9 or Fab fragments thereof may be produced via expression of Ab9 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab10

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 91:

(SEQ ID NO: 231)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAATGTTTACAATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTCGTGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 92:

(SEQ ID NO: 232)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAATGTTTACAATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTCGTGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 93:

(SEQ ID NO: 233)
GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGGCCTCAGTAGCTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTAGTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 94:

(SEQ ID NO: 234)
GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGGCCTCAGTAGCTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTAGTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 235; SEQ ID NO: 236; and SEQ ID NO: 237 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 91 or the light chain sequence of SEQ ID NO: 92.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 238; SEQ ID NO: 239; and SEQ ID NO:240 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 93 or the heavy chain sequence of SEQ ID NO: 94.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 231 encoding the light chain variable sequence of SEQ ID NO: 91; the polynucleotide SEQ ID NO: 232 encoding the light chain sequence of SEQ ID NO: 92; the polynucleotide SEQ ID NO: 233 encoding the heavy chain variable sequence of SEQ ID NO: 93; the polynucleotide SEQ ID NO: 234 encoding the heavy chain sequence of SEQ ID NO: 94; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 235; SEQ ID NO: 236; and SEQ ID NO: 237) of the light chain variable sequence of SEQ ID NO: 91 or the light chain sequence of SEQ ID NO: 92; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 238; SEQ ID NO: 239; and SEQ ID NO: 240) of the heavy chain variable sequence of SEQ ID NO: 93 or the heavy chain sequence of SEQ ID NO: 94.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab10, the polynucleotides encoding the full length Ab10 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 232 encoding the light chain sequence of SEQ ID NO: 92 and the polynucleotide SEQ ID NO: 234 encoding the heavy chain sequence of SEQ ID NO: 94.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab10 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab10 or Fab fragments thereof may be produced via expression of Ab10 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab11

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucle-

otides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 101:

(SEQ ID NO: 241)
CAGGTGCTGACCCAGACTGCATCCCCCGTGTCTCCAGCTGTGGGAAGCAC
AGTCACCATCAATTGCCGGGCCAGTCAGAGTGTTTATTATAACAACTACC
TAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTAT
TCTACATCCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAGTGG
ATCTGGGACACAGTTCACTCTCACCATCAGCGACGTGCAGTGTGACGATG
CTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTAATGGTGATTGT
TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 102:

(SEQ ID NO: 242)
CAGGTGCTGACCCAGACTGCATCCCCCGTGTCTCCAGCTGTGGGAAGCAC
AGTCACCATCAATTGCCGGGCCAGTCAGAGTGTTTATTATAACAACTACC
TAGCCTGGTATCAGCAGAAACCAGGGCAGCCTCCCAAGCAACTGATCTAT
TCTACATCCACTCTGGCATCTGGGGTCTCATCGCGGTTCAAAGGCAGTGG
ATCTGGGACACAGTTCACTCTCACCATCAGCGACGTGCAGTGTGACGATG
CTGCCACTTACTACTGTCTAGGCAGTTATGATTGTAGTAATGGTGATTGT
TTTGTTTTCGGCGGAGGGACCGAGGTGGTGGTCAAACGTACGGTGGCTGC
ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA
CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA
GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG
TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC
TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA
GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG
AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 103:

(SEQ ID NO: 243)
CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGAGGATCCCT
GACACTCACCTGCACAGTCTCTGGAATCGACGTCACTAACTACTATATGC
AATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATT
GGTGTGAATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATTCAC
CATCTCCAAAACCTCGTCGACCACGGTGGATCTGAAAATGACCAGTCTGA
CAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGCGACATCTGGGGC
CCGGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 104:

(SEQ ID NO: 244)
CAGTCGCTGGAGGAGTCCGGGGGTCGCCTGGTCACGCCTGGAGGATCCCT
GACACTCACCTGCACAGTCTCTGGAATCGACGTCACTAACTACTATATGC
AATGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAATGGATCGGAGTCATT
GGTGTGAATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATTCAC
CATCTCCAAAACCTCGTCGACCACGGTGGATCTGAAAATGACCAGTCTGA
CAACCGAGGACACGGCCACCTATTTCTGTGCCAGAGGCGACATCTGGGGC
CCGGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCCATCGGT
CTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAGCGGCCC
TGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTCGTGG
AACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCCTACA
GTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCA
GCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAAC
ACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAACTCACAC
ATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAGTCTTCC
TCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAG
GTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAAGTT
CAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGC
GGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTCACCGTC
CTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAA
CAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCAAAGGGC
AGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAGGAGATG
ACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCAG
CGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAACTACA
AGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTCTACAGC
AAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTTCTCATG
CTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGAGCCTCT
CCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 245; SEQ ID NO: 246; and SEQ ID NO: 247 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 101 or the light chain sequence of SEQ ID NO: 102.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 248; SEQ ID NO: 249; and SEQ ID NO: 250 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 103 or the heavy chain sequence of SEQ ID NO: 104.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding

antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 241 encoding the light chain variable sequence of SEQ ID NO: 101; the polynucleotide SEQ ID NO: 242 encoding the light chain sequence of SEQ ID NO: 102; the polynucleotide SEQ ID NO: 243 encoding the heavy chain variable sequence of SEQ ID NO: 103; the polynucleotide SEQ ID NO: 244 encoding the heavy chain sequence of SEQ ID NO: 104; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 245; SEQ ID NO: 246; and SEQ ID NO: 247) of the light chain variable sequence of SEQ ID NO: 101 or the light chain sequence of SEQ ID NO: 102; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 248; SEQ ID NO: 249; and SEQ ID NO: 250) of the heavy chain variable sequence of SEQ ID NO: 103 or the heavy chain sequence of SEQ ID NO: 104.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab11, the polynucleotides encoding the full length Ab11 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 242 encoding the light chain sequence of SEQ ID NO: 102 and the polynucleotide SEQ ID NO: 244 encoding the heavy chain sequence of SEQ ID NO: 104.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab11 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab11 or Fab fragments thereof may be produced via expression of Ab11 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab12

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 111:

(SEQ ID NO: 251)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCGGGCCAGTCAGAGTGTTTACTATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTAATGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 112:

(SEQ ID NO: 252)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCGGGCCAGTCAGAGTGTTTACTATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTAATGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 113:

(SEQ ID NO: 253)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACGTCACTAACTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTGTGAATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCCAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 114:

(SEQ ID NO: 254)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGACGTCACTAACTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTGTGAATGGTAAGAGATACTACGCGAGCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTGCCAGAGGGGACATC

-continued

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 255; SEQ ID NO: 256; and SEQ ID NO: 257 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 111 or the light chain sequence of SEQ ID NO: 112.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 258; SEQ ID NO: 259; and SEQ ID NO: 260 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 113 or the heavy chain sequence of SEQ ID NO: 114.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 251 encoding the light chain variable sequence of SEQ ID NO: 111; the polynucleotide SEQ ID NO: 252 encoding the light chain sequence of SEQ ID NO: 112; the polynucleotide SEQ ID NO: 253 encoding the heavy chain variable sequence of SEQ ID NO: 113; the polynucleotide SEQ ID NO: 254 encoding the heavy chain sequence of SEQ ID NO: 114; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 255; SEQ ID NO: 256; and SEQ ID NO: 257) of the light chain variable sequence of SEQ ID NO: 111 or the light chain sequence of SEQ ID NO: 112; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 258; SEQ ID NO: 259; and SEQ ID NO: 260) of the heavy chain variable sequence of SEQ ID NO: 113 or the heavy chain sequence of SEQ ID NO: 114.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab12, the polynucleotides encoding the full length Ab12 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 252 encoding the light chain sequence of SEQ ID NO: 112 and the polynucleotide SEQ ID NO: 254 encoding the heavy chain sequence of SEQ ID NO: 114.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab12 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab12 or Fab fragments thereof may be produced via expression of Ab12 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab13

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 121:

(SEQ ID NO: 261)

GCCATCGTGATGACCCAGACTCCATCTTCCAAGTCTGTCCCTGTGGGAGA

CACAGTCACCATCAATTGCCAGGCCAGTGAGAGTCTTTATAATAACAACG

CCTTGGCCTGGTTTCAGCAGAAACCAGGGCAGCCTCCCAAGCGCCTGATC

TATGATGCATCCAAACTGGCATCTGGGGTCCCATCGCGGTTCAGTGGCGG

TGGGTCTGGGACACAGTTCACTCTCACCATCAGTGGCGTGCAGTGTGACG

ATGCTGCCACTTACTACTGTGGAGGCTACAGAAGTGATAGTGTTGATGGT

GTTGCTTTCGCCGGAGGGACCGAGGTGGTGGTCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 122:

(SEQ ID NO: 262)

GCCATCGTGATGACCCAGACTCCATCTTCCAAGTCTGTCCCTGTGGGAGA

CACAGTCACCATCAATTGCCAGGCCAGTGAGAGTCTTTATAATAACAACG

-continued

CCTTGGCCTGGTTTCAGCAGAAACCAGGGCAGCCTCCCAAGCGCCTGATC

TATGATGCATCCAAACTGGCATCTGGGGTCCCATCGCGGTTCAGTGGCGG

TGGGTCTGGGACACAGTTCACTCTCACCATCAGTGGCGTGCAGTGTGACG

ATGCTGCCACTTACTACTGTGGAGGCTACAGAAGTGATAGTGTTGATGGT

GTTGCTTTCGCCGGAGGGACCGAGGTGGTGGTCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 123:

(SEQ ID NO: 263)

CAGTCGGTGGAGGAGTCCGGGGGAGGCCTGGTCCAGCCTGAGGGATCCCT

GACACTCACCTGCACAGCCTCTGGATTCGACTTCAGTAGCAATGCAATGT

GGTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGATGCATT

TACAATGGTGATGGCAGCACATACTACGCGAGCTGGGTGAATGGCCGATT

CTCCATCTCCAAAACCTCGTCGACCACGGTGACTCTGCAACTGAATAGTC

TGACAGTCGCGGACACGGCCACGTATTATTGTGCGAGAGATCTTGACTTG

TGGGGCCCGGGCACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 124:

(SEQ ID NO: 264)

CAGTCGGTGGAGGAGTCCGGGGGAGGCCTGGTCCAGCCTGAGGGATCCCT

GACACTCACCTGCACAGCCTCTGGATTCGACTTCAGTAGCAATGCAATGT

GGTGGGTCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGATCGGATGCATT

TACAATGGTGATGGCAGCACATACTACGCGAGCTGGGTGAATGGCCGATT

CTCCATCTCCAAAACCTCGTCGACCACGGTGACTCTGCAACTGAATAGTC

TGACAGTCGCGGACACGGCCACGTATTATTGTGCGAGAGATCTTGACTTG

TGGGGCCCGGGCACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACAAGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

-continued

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 265; SEQ ID NO: 266; and SEQ ID NO: 267 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 121 or the light chain sequence of SEQ ID NO: 122.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 268; SEQ ID NO: 269; and SEQ ID NO: 270 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 123 or the heavy chain sequence of SEQ ID NO: 124.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 261 encoding the light chain variable sequence of SEQ ID NO: 121; the polynucleotide SEQ ID NO: 262 encoding the light chain sequence of SEQ ID NO: 122; the polynucleotide SEQ ID NO: 263 encoding the heavy chain variable sequence of SEQ ID NO: 123; the polynucleotide SEQ ID NO: 264 encoding the heavy chain sequence of SEQ ID NO: 124; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 265; SEQ ID NO: 266; and SEQ ID NO: 267) of the light chain variable sequence of SEQ ID NO: 121 or the light chain sequence of SEQ ID NO: 122; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 268; SEQ ID NO: 269; and SEQ ID NO: 270) of the heavy chain variable sequence of SEQ ID NO: 123 or the heavy chain sequence of SEQ ID NO: 124.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding)

fragments having binding specificity for CGRP. With respect to antibody Ab13, the polynucleotides encoding the full length Ab13 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 262 encoding the light chain sequence of SEQ ID NO: 122 and the polynucleotide SEQ ID NO: 264 encoding the heavy chain sequence of SEQ ID NO: 124.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab13 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab13 or Fab fragments thereof may be produced via expression of Ab13 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

Antibody Ab14

The invention is further directed to polynucleotides encoding antibody polypeptides having binding specificity to CGRP. In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable light chain polypeptide sequence of SEQ ID NO: 131:

(SEQ ID NO: 271)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAATGTTTACAATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTCGTGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGT.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the light chain polypeptide sequence of SEQ ID NO: 132:

(SEQ ID NO: 272)

CAAGTGCTGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAG

AGTCACCATCAATTGCCAGGCCAGTCAGAATGTTTACAATAACAACTACC

TAGCCTGGTATCAGCAGAAACCAGGGAAAGTTCCTAAGCAACTGATCTAT

TCTACATCCACTCTGGCATCTGGGGTCCCATCTCGTTTCAGTGGCAGTGG

ATCTGGGACAGATTTCACTCTCACCATCAGCAGCCTGCAGCCTGAAGATG

TTGCAACTTATTACTGTCTGGGCAGTTATGATTGTAGTCGTGGTGATTGT

TTTGTTTTCGGCGGAGGAACCAAGGTGGAAATCAAACGTACGGTGGCTGC

ACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA

CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAA

-continued

GTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAG

TGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCC

TGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA

GTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGG

AGAGTGTTAG.

In another embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the variable heavy chain polypeptide sequence of SEQ ID NO: 133:

(SEQ ID NO: 273)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGGCCTCAGTAGCTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTAGTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGC.

In one embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, the following polynucleotide sequence encoding the heavy chain polypeptide sequence of SEQ ID NO: 134:

(SEQ ID NO: 274)

GAGGTGCAGCTTGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTC

CCTGAGACTCTCCTGTGCAGTCTCTGGAATCGGCCTCAGTAGCTACTACA

TGCAATGGGTCCGTCAGGCTCCAGGGAAGGGGCTGGAGTGGGTCGGAGTC

ATTGGTAGTGATGGTAAGACATACTACGCGACCTGGGCGAAAGGCCGATT

CACCATCTCCAGAGACAATTCCAAGACCACGGTGTATCTTCAAATGAACA

GCCTGAGAGCTGAGGACACTGCTGTGTATTTCTGTACCAGAGGGGACATC

TGGGGCCAAGGGACCCTCGTCACCGTCTCGAGCGCCTCCACCAAGGGCCC

ATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAGCACCTCTGGGGGCACAG

CGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTG

TCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGT

CCTACAGTCCTCAGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCT

CCAGCAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCC

AGCAACACCAAGGTGGACGCGAGAGTTGAGCCCAAATCTTGTGACAAAAC

TCACACATGCCCACCGTGCCCAGCACCTGAACTCCTGGGGGGACCGTCAG

TCTTCCTCTTCCCCCCAAAACCCAAGGACACCCTCATGATCTCCCGGACC

CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGT

CAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAA

AGCCGCGGGAGGAGCAGTACGCCAGCACGTACCGTGTGGTCAGCGTCCTC

ACCGTCCTGCACCAGGACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGT

CTCCAACAAAGCCCTCCCAGCCCCCATCGAGAAAACCATCTCCAAAGCCA

AAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCCATCCCGGGAG

-continued

GAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTA

TCCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACA

ACTACAAGACCACGCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCCTC

TACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAGCAGGGGAACGTCTT

CTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAGAAGA

GCCTCTCCCTGTCTCCGGGTAAATGA.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 275; SEQ ID NO: 276; and SEQ ID NO: 277 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the light chain variable sequence of SEQ ID NO: 131 or the light chain sequence of SEQ ID NO: 132.

In a further embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one or more of the polynucleotide sequences of SEQ ID NO: 278; SEQ ID NO: 279; and SEQ ID NO: 280 which correspond to polynucleotides encoding the complementarity-determining regions (CDRs, or hypervariable regions) of the heavy chain variable sequence of SEQ ID NO: 133 or the heavy chain sequence of SEQ ID NO: 134.

The invention also contemplates polynucleotide sequences including one or more of the polynucleotide sequences encoding antibody fragments described herein. In one embodiment of the invention, polynucleotides encoding antibody fragments having binding specificity to CGRP comprise, or alternatively consist of, one, two, three or more, including all of the following polynucleotides encoding antibody fragments: the polynucleotide SEQ ID NO: 271 encoding the light chain variable sequence of SEQ ID NO: 131; the polynucleotide SEQ ID NO: 272 encoding the light chain sequence of SEQ ID NO: 132; the polynucleotide SEQ ID NO: 273 encoding the heavy chain variable sequence of SEQ ID NO: 133; the polynucleotide SEQ ID NO: 274 encoding the heavy chain sequence of SEQ ID NO: 134; polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 275; SEQ ID NO: 276; and SEQ ID NO: 277) of the light chain variable sequence of SEQ ID NO: 131 or the light chain sequence of SEQ ID NO: 132; and polynucleotides encoding the complementarity-determining regions (SEQ ID NO: 278; SEQ ID NO: 279; and SEQ ID NO: 280) of the heavy chain variable sequence of SEQ ID NO: 133 or the heavy chain sequence of SEQ ID NO: 134.

In a preferred embodiment of the invention, polynucleotides of the invention comprise, or alternatively consist of, polynucleotides encoding Fab (fragment antigen binding) fragments having binding specificity for CGRP. With respect to antibody Ab14, the polynucleotides encoding the full length Ab14 antibody comprise, or alternatively consist of, the polynucleotide SEQ ID NO: 272 encoding the light chain sequence of SEQ ID NO: 132 and the polynucleotide SEQ ID NO: 274 encoding the heavy chain sequence of SEQ ID NO: 134.

Another embodiment of the invention contemplates these polynucleotides incorporated into an expression vector for expression in mammalian cells such as CHO, NSO, HEK-293, or in fungal, insect, or microbial systems such as yeast cells such as the yeast *Pichia*. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*. In one embodiment of the invention described herein (infra), Fab fragments may be produced by enzymatic digestion (e.g., papain) of Ab14 following expression of the full-length polynucleotides in a suitable host. In another embodiment of the invention, anti-CGRP antibodies such as Ab14 or Fab fragments thereof may be produced via expression of Ab14 polynucleotides in mammalian cells such as CHO, NSO or HEK 293 cells, fungal, insect, or microbial systems such as yeast cells (for example diploid yeast such as diploid *Pichia*) and other yeast strains. Suitable *Pichia* species include, but are not limited to, *Pichia pastoris*.

In one embodiment, the invention is directed to an isolated polynucleotide comprising a polynucleotide encoding an anti-CGRP V_H antibody amino acid sequence selected from SEQ ID NO: 3, 13, 23, 33, 43, 53, 63, 73, 83, 93, 103, 113, 123, or 133, or encoding a variant thereof wherein at least one framework residue (FR residue) has been substituted with an amino acid present at the corresponding position in a rabbit anti-CGRP antibody V_H polypeptide or a conservative amino acid substitution.

In another embodiment, the invention is directed to an isolated polynucleotide comprising the polynucleotide sequence encoding an anti-CGRP V_L antibody amino acid sequence of 1, 11, 21, 31, 41, 51, 61, 71, 81, 91, 101, 111, 121, or 131, or encoding a variant thereof wherein at least one framework residue (FR residue) has been substituted with an amino acid present at the corresponding position in a rabbit anti-CGRP antibody V_L polypeptide or a conservative amino acid substitution.

In yet another embodiment, the invention is directed to one or more heterologous polynucleotides comprising a sequence encoding the polypeptides contained in SEQ ID NO:1 and SEQ ID NO:3; SEQ ID NO:11 and SEQ ID NO:13; SEQ ID NO:21 and SEQ ID NO:23; SEQ ID NO:31 and SEQ ID NO:33; SEQ ID NO:41 and SEQ ID NO:43; SEQ ID NO:51 and SEQ ID NO:53, SEQ ID NO:61 and SEQ ID NO:63; SEQ ID NO:71 and SEQ ID NO:73; SEQ ID NO:81 and SEQ ID NO:83; SEQ ID NO:91 and SEQ ID NO:93; SEQ ID NO:101 and SEQ ID NO:103; SEQ ID NO:111 and SEQ ID NO:113; SEQ ID NO:121 and SEQ ID NO:123; or SEQ ID NO:131 and SEQ ID NO:133.

In another embodiment, the invention is directed to an isolated polynucleotide that expresses a polypeptide containing at least one CDR polypeptide derived from an anti-CGRP antibody wherein said expressed polypeptide alone specifically binds CGRP or specifically binds CGRP when expressed in association with another polynucleotide sequence that expresses a polypeptide containing at least one CDR polypeptide derived from an anti-CGRP antibody wherein said at least one CDR is selected from those contained in the V_L or V_H polypeptides of SEQ ID NO: 1, 3, 11, 13, 21, 23, 31, 33, 41, 43, 51, 53, 61, 63, 71, 73, 81, 83, 91, 93, 101, 103, 111, 113, 121, 123, 131, or SEQ ID NO: 133.

Host cells and vectors comprising said polynucleotides are also contemplated.

The invention further contemplates vectors comprising the polynucleotide sequences encoding the variable heavy and light chain polypeptide sequences, as well as the individual complementarity-determining regions (CDRs, or hypervariable regions), as set forth herein, as well as host cells comprising said vector sequences. In one embodiment of the invention, the host cell is a yeast cell. In another embodiment of the invention, the yeast host cell belongs to the genus *Pichia*.

B-Cell Screening and Isolation

In one embodiment, the present invention contemplates the preparation and isolation of a clonal population of antigen-specific B cells that may be used for isolating at least one CGRP antigen-specific cell, which can be used to produce a monoclonal antibody against CGRP, which is specific to a desired CGRP antigen, or a nucleic acid sequence corresponding to such an antibody. Methods of preparing and isolating said clonal population of antigen-specific B cells are taught, for example, in U.S. patent publication no. US 2007/0269868 to Carvalho-Jensen et al., the disclosure of which is herein incorporated by reference in its entirety. Methods of preparing and isolating said clonal population of antigen-specific B cells are also taught herein in the examples. Methods of "enriching" a cell population by size or density are known in the art. See, e.g., U.S. Pat. No. 5,627,052. These steps can be used in addition to enriching the cell population by antigen-specificity.

Methods of Humanizing Antibodies

In another embodiment, the present invention contemplates methods for humanizing antibody heavy and light chains. Methods for humanizing antibody heavy and light chains which may be applied to anti-CGRP antibodies are taught, for example, in U.S. patent application publication no. US 2009/0022659 to Olson et al., and in U.S. Pat. No. 7,935,340 to Garcia-Martinez et al., the disclosures of each of which are herein incorporated by reference in their entireties.

Methods of Producing Antibodies and Fragments Thereof

In another embodiment, the present invention contemplates methods for producing anti-CGRP antibodies and fragments thereof. Methods for producing anti-CGRP antibodies and fragments thereof secreted from polyploidal, preferably diploid or tetraploid strains of mating competent yeast are taught, for example, in U.S. patent application publication no. US 2009/0022659 to Olson et al., and in U.S. Pat. No. 7,935,340 to Garcia-Martinez et al., the disclosures of each of which are herein incorporated by reference in their entireties.

Other methods of producing antibodies are well known to those of ordinary skill in the art. For example, methods of producing chimeric antibodies are now well known in the art (See, for example, U.S. Pat. No. 4,816,567 to Cabilly et al.; Morrison et al., P.N.A.S. USA, 81:8651-55 (1984); Neuberger, M. S. et al., Nature, 314:268-270 (1985); Boulianne, G. L. et al., Nature, 312:643-46 (1984), the disclosures of each of which are herein incorporated by reference in their entireties).

Likewise, other methods of producing humanized antibodies are now well known in the art (See, for example, U.S. Pat. Nos. 5,530,101, 5,585,089, 5,693,762, and 6,180,370 to Queen et al; U.S. Pat. Nos. 5,225,539 and 6,548,640 to Winter; U.S. Pat. Nos. 6,054,297, 6,407,213 and 6,639,055 to Carter et al; U.S. Pat. No. 6,632,927 to Adair; Jones, P. T. et al, Nature, 321:522-525 (1986); Reichmann, L., et al, Nature, 332:323-327 (1988); Verhoeyen, M, et al, Science, 239:1534-36 (1988), the disclosures of each of which are herein incorporated by reference in their entireties).

Antibody polypeptides of the invention having CGRP binding specificity may also be produced by constructing, using conventional techniques well known to those of ordinary skill in the art, an expression vector containing an operon and a DNA sequence encoding an antibody heavy chain in which the DNA sequence encoding the CDRs required for antibody specificity is derived from a non-human cell source, preferably a rabbit B-cell source, while the DNA sequence encoding the remaining parts of the antibody chain is derived from a human cell source.

A second expression vector is produced using the same conventional means well known to those of ordinary skill in the art, said expression vector containing an operon and a DNA sequence encoding an antibody light chain in which the DNA sequence encoding the CDRs required for antibody specificity is derived from a non-human cell source, preferably a rabbit B-cell source, while the DNA sequence encoding the remaining parts of the antibody chain is derived from a human cell source.

The expression vectors are transfected into a host cell by convention techniques well known to those of ordinary skill in the art to produce a transfected host cell, said transfected host cell cultured by conventional techniques well known to those of ordinary skill in the art to produce said antibody polypeptides.

The host cell may be co-transfected with the two expression vectors described above, the first expression vector containing DNA encoding an operon and a light chain-derived polypeptide and the second vector containing DNA encoding an operon and a heavy chain-derived polypeptide. The two vectors contain different selectable markers, but preferably achieve substantially equal expression of the heavy and light chain polypeptides. Alternatively, a single vector may be used, the vector including DNA encoding both the heavy and light chain polypeptides. The coding sequences for the heavy and light chains may comprise cDNA, genomic DNA, or both.

The host cells used to express the antibody polypeptides may be either a bacterial cell such as *E. coli*, or a eukaryotic cell such as *P. pastoris*. In one embodiment of the invention, a mammalian cell of a well-defined type for this purpose, such as a myeloma cell, a Chinese hamster ovary (CHO) cell line, a NSO cell line, or a HEK293 cell line may be used.

The general methods by which the vectors may be constructed, transfection methods required to produce the host cell and culturing methods required to produce the antibody polypeptides from said host cells all include conventional techniques. Although preferably the cell line used to produce the antibody is a mammalian cell line, any other suitable cell line, such as a bacterial cell line such as an *E. coli*-derived bacterial strain, or a yeast cell line, may alternatively be used.

Similarly, once produced the antibody polypeptides may be purified according to standard procedures in the art, such as for example cross-flow filtration, ammonium sulphate precipitation, affinity column chromatography and the like.

The antibody polypeptides described herein may also be used for the design and synthesis of either peptide or non-peptide mimetics that would be useful for the same therapeutic applications as the antibody polypeptides of the invention. See, for example, Saragobi et al, Science, 253: 792-795 (1991), the contents of which is herein incorporated by reference in its entirety.

Screening Assays

The invention also includes screening assays designed to assist in the identification of diseases and disorders associated with CGRP especially conditions associated with photophobia such as migraine, other headache and pain conditions, depression, bipolar disorder, agoraphobia and others in patients exhibiting symptoms of photophobia or a CGRP associated disease or disorder.

In one embodiment of the invention, the anti-CGRP antibodies of the invention, or CGRP binding fragments thereof, are used to detect the presence of CGRP in a biological sample obtained from a patient exhibiting symp-

toms of a disease or disorder associated with CGRP especially one associated with photophobia. The presence of CGRP, or elevated levels thereof when compared to predisease levels of CGRP in a comparable biological sample, may be beneficial in diagnosing a disease or disorder associated with CGRP.

Another embodiment of the invention provides a diagnostic or screening assay to assist in diagnosis of diseases or disorders associated with CGRP and photophobia in patients exhibiting symptoms of a CGRP associated disease or disorder identified herein, comprising assaying the level of CGRP expression in a biological sample from said patient using a post-translationally modified anti-CGRP antibody or binding fragment thereof. The anti-CGRP antibody or binding fragment thereof may be post-translationally modified to include a detectable moiety such as set forth previously in the disclosure.

The CGRP level in the biological sample is determined using a modified anti-CGRP antibody or binding fragment thereof as set forth herein, and comparing the level of CGRP in the biological sample against a standard level of CGRP (e.g., the level in normal biological samples). The skilled clinician would understand that some variability may exist between normal biological samples, and would take that into consideration when evaluating results. In one embodiment of the invention, the anti-CGRP antibodies of the invention may be used to correlate CGRP expression levels with a particular stage of cancerous development. One skilled in the art would be able to measure CGRP in numerous subjects in order to establish ranges of CGRP expression that correspond to clinically defined stages of cancerous development. These ranges will allow the skilled practitioner to measure CGRP in a subject diagnosed with a cancer and correlate the levels in each subject with a range that corresponds to a stage of said cancer. One skilled in the art would understand that by measuring CGRP in the patient at different intervals, the progression of the cancer can be determined.

The above-recited assay may also be useful in monitoring a disease or disorder, where the level of CGRP obtained in a biological sample from a patient believed to have a CGRP associated disease or disorder especially one associated with photophobia is compared with the level of CGRP in prior biological samples from the same patient, in order to ascertain whether the CGRP level in said patient has changed with, for example, a treatment regimen.

The invention is also directed to a method of in vivo imaging which detects the presence of cells which express CGRP comprising administering a diagnostically effective amount of a diagnostic composition. Said in vivo imaging is useful for the detection or imaging of CGRP expressing tumors or metastases, for example, and can be useful as part of a planning regimen for the design of an effective cancer treatment protocol. The treatment protocol may include, for example, one or more of radiation, chemotherapy, cytokine therapy, gene therapy, and antibody therapy, as well as an anti-CGRP antibody or fragment thereof.

The present invention further provides for a kit for detecting binding of an anti-CGRP antibody of the invention to CGRP. In particular, the kit may be used to detect the presence of a CGRP specifically reactive with an anti-CGRP antibody of the invention or an immunoreactive fragment thereof. The kit may also include an antibody bound to a substrate, a secondary antibody reactive with the antigen and a reagent for detecting a reaction of the secondary antibody with the antigen. Such a kit may be an ELISA kit and can comprise the substrate, primary and secondary antibodies when appropriate, and any other necessary reagents such as detectable moieties, enzyme substrates, and color reagents, for example as described herein. The diagnostic kit may also be in the form of an immunoblot kit. The diagnostic kit may also be in the form of a chemiluminescent kit (Meso Scale Discovery, Gaithersburg, Md.). The diagnostic kit may also be a lanthanide-based detection kit (PerkinElmer, San Jose, Calif.).

A skilled clinician would understand that a biological sample includes, but is not limited to, sera, plasma, urine, saliva, mucous, pleural fluid, synovial fluid and spinal fluid.

Methods of Ameliorating or Reducing Symptoms of, or Treating, or Preventing, Diseases and Disorders Associated with, CGRP

In another embodiment of the invention, anti-CGRP antibodies described herein, or fragments thereof, are useful for ameliorating or reducing the symptoms of, or treating, or preventing, diseases and disorders associated with CGRP especially for treatment or prevention of photophobia. In a preferred embodiment the anti-CGRP antibodies or antibody fragments will be shown to be efficacious (block adverse side effects associated with excess circulating CGRP including light aversive behavior) in the rodent animal model disclosed in Example 8.

Anti-CGRP antibodies described herein, or fragments thereof, as well as combinations, can also be administered in a therapeutically effective amount to patients in need of treatment of diseases and disorders associated with CGRP for treatment or prevention of photophobia in the form of a pharmaceutical composition as described in greater detail below.

In another embodiment of the invention, anti-CGRP antibodies described herein, or fragments thereof, are useful for ameliorating or reducing the symptoms of, or treating, or preventing, migraines (with or without aura), weight loss, cancer or tumors, angiogenesis associated with cancer or tumor growth, angiogenesis associated with cancer or tumor survival, pain, hemiplagic migraines, cluster headaches, migrainous neuralgia, chronic headaches, tension headaches, general headaches, headache-free migraine, abdominal migraine, hot flashes, chronic paroxysomal hemicrania, secondary headaches due to an underlying structural problem in the head or neck, cranial neuralgia, sinus headaches (such as for example associated with sinusitis), and allergy-induced headaches or migraines.

In another embodiment of the invention, anti-CGRP antibodies described herein, or fragments thereof and/or with a second agent, are useful for ameliorating or reducing the symptoms of, or treating, or preventing, photophobia associated with the following non-limiting listing of diseases and disorders: neurogenic, neuropathic or nociceptic pain. Neuropathic pain may include, but is not limited to, trigeminal neuralgia, post-herpetic neuralgia, phantom limb pain, fibromyalgia, menstrual pain, ovarialgia, reflex sympathetic dystrophy and neurogenic pain. In other preferred embodiments, anti-CGRP antibodies described herein, or fragments thereof and/or with a second agent, are useful for ameliorating or reducing the symptoms of, or treating, or preventing, photophobia associated with osteoarthritis or rheumatoid arthritis pain, lower back pain, diabetic neuropathy, sciatica, and other neuropathic pain.

In another embodiment of the invention, anti-CGRP antibodies described herein, or fragments thereof and/or with a second agent, are useful for ameliorating or reducing the symptoms of, or treating, or preventing, photophobia associated with the following non-limiting listing of diseases and disorders: visceral pain or more specifically associated with

gastro-esophageal reflux, dyspepsia, irritable bowel syndrome, inflammatory bowel disease, Crohn's disease, ileitis, ulcerative colitis, renal colic, dysmenorrhea, cystitis, menstrual period, labor, menopause, prostatitis, or pancreatitis.

Administration

In one embodiment of the invention, the anti-CGRP antibodies or fragments described herein, or anti-CGRP-R antibodies or fragments thereof, as well as combinations of said antibodies or antibody fragments, for treatment or prevention of photophobia, are administered to a subject at a concentration of between about 0.1 and 100.0 mg/kg of body weight of recipient subject. In a preferred embodiment of the invention, the anti-CGRP antibodies described herein, or CGRP binding fragments thereof, as well as combinations of said antibodies or antibody fragments, are administered to a subject at a concentration of about 0.4 mg/kg of body weight of recipient subject. In a preferred embodiment of the invention, the anti-CGRP antibodies described herein, or CGRP binding fragments thereof, as well as combinations of said antibodies or antibody fragments, are administered to a recipient subject with a frequency of once every twenty-six weeks or less, such as once every sixteen weeks or less, once every eight weeks or less, once every four weeks or less, once every two weeks or less, once every week or less, or once daily or less.

Fab fragments for treatment or prevention of photophobia may be administered every two weeks or less, every week or less, once daily or less, multiple times per day, and/or every few hours. In one embodiment of the invention, a patient receives Fab fragments of 0.1 mg/kg to 40 mg/kg per day given in divided doses of 1 to 6 times a day, or in a sustained release form, effective to obtain desired results.

It is to be understood that the concentration of the antibody or Fab administered to a given patient for treatment or prevention of photophobia may be greater or lower than the exemplary administration concentrations set forth above in paragraphs [0566] and [0567].

A person of skill in the art would be able to determine an effective dosage and frequency of administration for treatment or prevention of photophobia through routine experimentation, for example guided by the disclosure herein and the teachings in Goodman, L. S., Gilman, A., Brunton, L. L., Lazo, J. S., & Parker, K. L. (2006). Goodman & Gilman's the pharmacological basis of therapeutics. New York: McGraw-Hill; Howland, R. D., Mycek, M. J., Harvey, R. A., Champe, P. C., & Mycek, M. J. (2006). Pharmacology. Lippincott's illustrated reviews. Philadelphia: Lippincott Williams & Wilkins; and Golan, D. E. (2008). Principles of pharmacology: the pathophysiologic basis of drug therapy. Philadelphia, Pa., [etc.]: Lippincott Williams & Wilkins.

In another embodiment of the invention, the anti-CGRP antibodies described herein, or CGRP binding fragments thereof, as well as combinations of said antibodies or antibody fragments, for treatment or prevention of photophobia are administered to a subject in a pharmaceutical formulation.

A "pharmaceutical composition" refers to a chemical or biological composition suitable for administration to a mammal. Such compositions may be specifically formulated for administration via one or more of a number of routes, including but not limited to buccal, epicutaneous, epidural, inhalation, intraarterial, intracardial, intracerebroventricular, intradermal, intramuscular, intranasal, intraocular, intraperitoneal, intraspinal, intrathecal, intravenous, oral, parenteral, rectally via an enema or suppository, subcutaneous, subdermal, sublingual, transdermal, and transmucosal. In addition, administration can occur by means of injection, powder, liquid, gel, drops, or other means of administration.

In one embodiment of the invention, the anti-CGRP antibodies described herein, or CGRP binding fragments thereof, as well as combinations of said antibodies or antibody fragments, for treatment or prevention of photophobia may be optionally administered in combination with one or more active agents. Such active agents include analgesic, anti-histamine, antipyretic, anti-inflammatory, antibiotic, antiviral, and anti-cytokine agents. Active agents include agonists, antagonists, and modulators of TNF-α, IL-2, IL-4, IL-6, IL-10, IL-12, IL-13, IL-18, IFN-α, IFN-γ, BAFF, CXCL13, IP-10, VEGF, EPO, EGF, HRG, Hepatocyte Growth Factor (HGF), Hepcidin, including antibodies reactive against any of the foregoing, and antibodies reactive against any of their receptors. Active agents also include but are not limited to 2-Arylpropionic acids, Aceclofenac, Acemetacin, Acetylsalicylic acid (Aspirin), Alclofenac, Alminoprofen, Amoxiprin, Ampyrone, Arylalkanoic acids, Azapropazone, Benorylate/Benorilate, Benoxaprofen, Bromfenac, Carprofen, Celecoxib, Choline magnesium salicylate, Clofezone, COX-2 inhibitors, Dexibuprofen, Dexketoprofen, Diclofenac, Diflunisal, Droxicam, Ethenzamide, Etodolac, Etoricoxib, Faislamine, fenamic acids, Fenbufen, Fenoprofen, Flufenamic acid, Flunoxaprofen, Flurbiprofen, Ibuprofen, Ibuproxam, Indometacin, Indoprofen, Kebuzone, Ketoprofen, Ketorolac, Lornoxicam, Loxoprofen, Lumiracoxib, Magnesium salicylate, Meclofenamic acid, Mefenamic acid, Meloxicam, Metamizole, Methyl salicylate, Mofebutazone, Nabumetone, Naproxen, N-Arylanthranilic acids, Nerve Growth Factor (NGF), Oxametacin, Oxaprozin, Oxicams, Oxyphenbutazone, Parecoxib, Phenazone, Phenylbutazone, Phenylbutazone, Piroxicam, Pirprofen, profens, Proglumetacin, Pyrazolidine derivatives, Rofecoxib, Salicyl salicylate, Salicylamide, Salicylates, Substance P, Sulfinpyrazone, Sulindac, Suprofen, Tenoxicam, Tiaprofenic acid, Tolfenamic acid, Tolmetin, and Valdecoxib.

An anti-histamine can be any compound that opposes the action of histamine or its release from cells (e.g., mast cells). Anti-histamines include but are not limited to acrivastine, astemizole, azatadine, azelastine, betatastine, brompheniramine, buclizine, cetirizine, cetirizine analogues, chlorpheniramine, clemastine, CS 560, cyproheptadine, desloratadine, dexchlorpheniramine, ebastine, epinastine, fexofenadine, HSR 609, hydroxyzine, levocabastine, loratidine, methscopolamine, mizolastine, norastemizole, phenindamine, promethazine, pyrilamine, terfenadine, and tranilast.

Antibiotics include but are not limited to Amikacin, Aminoglycosides, Amoxicillin, Ampicillin, Ansamycins, Arsphenamine, Azithromycin, Azlocillin, Aztreonam, Bacitracin, Carbacephem, Carbapenems, Carbenicillin, Cefaclor, Cefadroxil, Cefalexin, Cefalothin, Cefalotin, Cefamandole, Cefazolin, Cefdinir, Cefditoren, Cefepime, Cefixime, Cefoperazone, Cefotaxime, Cefoxitin, Cefpodoxime, Cefprozil, Ceftazidime, Ceftibuten, Ceftizoxime, Ceftobiprole, Ceftriaxone, Cefuroxime, Cephalosporins, Chloramphenicol, Cilastatin, Ciprofloxacin, Clarithromycin, Clindamycin, Cloxacillin, Colistin, Co-trimoxazole, Dalfopristin, Demeclocycline, Dicloxacillin, Dirithromycin, Doripenem, Doxycycline, Enoxacin, Ertapenem, Erythromycin, Ethambutol, Flucloxacillin, Fosfomycin, Furazolidone, Fusidic acid, Gatifloxacin, Geldanamycin, Gentamicin, Glycopeptides, Herbimycin, Imipenem, Isoniazid, Kanamycin, Levofloxacin, Lincomycin, Linezolid, Lomefloxacin, Loracarbef, Macrolides, Mafenide, Meropenem, Meticillin, Metronida-

zole, Mezlocillin, Minocycline, Monobactams, Moxifloxacin, Mupirocin, Nafcillin, Neomycin, Netilmicin, Nitrofurantoin, Norfloxacin, Ofloxacin, Oxacillin, Oxytetracycline, Paromomycin, Penicillin, Penicillins, Piperacillin, Platensimycin, Polymyxin B, Polypeptides, Prontosil, Pyrazinamide, Quinolones, Quinupristin, Rifampicin, Rifampin, Roxithromycin, Spectinomycin, Streptomycin, Sulfacetamide, Sulfamethizole, Sulfanilimide, Sulfasalazine, Sulfisoxazole, Sulfonamides, Teicoplanin, Telithromycin, Tetracycline, Tetracyclines, Ticarcillin, Tinidazole, Tobramycin, Trimethoprim, Trimethoprim-Sulfamethoxazole, Troleandomycin, Trovafloxacin, and Vancomycin.

Active agents also include Aldosterone, Beclometasone, Betamethasone, Corticosteroids, Cortisol, Cortisone acetate, Deoxycorticosterone acetate, Dexamethasone, Fludrocortisone acetate, Glucocorticoids, Hydrocortisone, Methylprednisolone, Prednisolone, Prednisone, Steroids, and Triamcinolone. Any suitable combination of these active agents is also contemplated.

In preferred embodiments, the subject antibodies and antibody fragments may be administered in a therapeutic regimen that includes compounds typically used to treat migraines, including migraines associated with photophobia. Examples hereof include analgesics such as NSAIDs. Examples include those afore-mentioned such as Ibuprofen, naproxen, sumatriptan, Paracetamol/acetaminophen, either alone or in combination with metoclopramide, and caffeine.

Triptans such as sumatriptan are commonly used as are Ergotamines such as Ergotamine. In addition, corticosteroids may be used.

Also, antimimetics may help relieve symptoms of nausea and help prevent vomiting, which can diminish the effectiveness of orally taken analgesics. In addition, some antiemetics, such as metoclopramide, are prokinetics and help gastric emptying, which is often impaired during episodes of migraine. Three combination antiemetic and analgesic preparations used for migraines include (aspirin with metoclopramide), (paracetamol/codeine for analgesia, with buclizine as the antiemetic) and paracetamol/metoclopramide.

A "pharmaceutical excipient" or a "pharmaceutically acceptable excipient" is a carrier, usually a liquid, in which an active therapeutic agent is formulated. In one embodiment of the invention, the active therapeutic agent is a humanized antibody described herein, or one or more fragments thereof. The excipient generally does not provide any pharmacological activity to the formulation, though it may provide chemical and/or biological stability, and release characteristics. Exemplary formulations can be found, for example, in Remington's Pharmaceutical Sciences, 19th Ed., Grennaro, A., Ed., 1995 which is incorporated by reference.

As used herein "pharmaceutically acceptable carrier" or "excipient" includes any and all solvents, dispersion media, coatings, antibacterial and antifungal agents, isotonic and absorption delaying agents that are physiologically compatible. In one embodiment, the carrier is suitable for parenteral administration. Alternatively, the carrier can be suitable for intravenous, intraperitoneal, intramuscular, or sublingual administration. Pharmaceutically acceptable carriers include sterile aqueous solutions or dispersions and sterile powders for the extemporaneous preparation of sterile injectable solutions or dispersions. The use of such media and agents for pharmaceutically active substances is well known in the art. Except insofar as any conventional media or agent is incompatible with the active compound, use thereof in the pharmaceutical compositions of the invention is contemplated. Supplementary active compounds can also be incorporated into the compositions.

Pharmaceutical compositions typically must be sterile and stable under the conditions of manufacture and storage. The invention contemplates that the pharmaceutical composition is present in lyophilized form. The composition can be formulated as a solution, microemulsion, liposome, or other ordered structure suitable to high drug concentration. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol, and liquid polyethylene glycol), and suitable mixtures thereof. The invention further contemplates the inclusion of a stabilizer in the pharmaceutical composition. The proper fluidity can be maintained, for example, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants.

In many cases, it will be preferable to include isotonic agents, for example, sugars, polyalcohols such as mannitol, sorbitol, or sodium chloride in the composition. Prolonged absorption of the injectable compositions can be brought about by including in the composition an agent which delays absorption, for example, monostearate salts and gelatin. Moreover, the alkaline polypeptide can be formulated in a time release formulation, for example in a composition which includes a slow release polymer. The active compounds can be prepared with carriers that will protect the compound against rapid release, such as a controlled release formulation, including implants and microencapsulated delivery systems. Biodegradable, biocompatible polymers can be used, such as ethylene vinyl acetate, polyanhydrides, polyglycolic acid, collagen, polyorthoesters, polylactic acid and polylactic, polyglycolic copolymers (PLG). Many methods for the preparation of such formulations are known to those skilled in the art.

For each of the recited embodiments, the compounds can be administered by a variety of dosage forms. Any biologically-acceptable dosage form known to persons of ordinary skill in the art, and combinations thereof, are contemplated. Examples of such dosage forms include, without limitation, reconstitutable powders, elixirs, liquids, solutions, suspensions, emulsions, powders, granules, particles, microparticles, dispersible granules, cachets, inhalants, aerosol inhalants, patches, particle inhalants, implants, depot implants, injectables (including subcutaneous, intramuscular, intravenous, and intradermal), infusions, and combinations thereof.

The above description of various illustrated embodiments of the invention is not intended to be exhaustive or to limit the invention to the precise form disclosed. While specific embodiments of, and examples for, the invention are described herein for illustrative purposes, various equivalent modifications are possible within the scope of the invention, as those skilled in the relevant art will recognize. The teachings provided herein of the invention can be applied to other purposes, other than the examples described above.

These and other changes can be made to the invention in light of the above detailed description. In general, in the following claims, the terms used should not be construed to limit the invention to the specific embodiments disclosed in the specification and the claims. Accordingly, the invention is not limited by the disclosure, but instead the scope of the invention is to be determined entirely by the following claims.

The invention may be practiced in ways other than those particularly described in the foregoing description and examples. Numerous modifications and variations of the invention are possible in light of the above teachings and, therefore, are within the scope of the appended claims.

Certain teachings related to methods for obtaining a clonal population of antigen-specific B cells were disclosed

in U.S. Provisional patent application No. 60/801,412, filed May 19, 2006, the disclosure of which is herein incorporated by reference in its entirety.

Certain teachings related to humanization of rabbit-derived monoclonal antibodies and preferred sequence modifications to maintain antigen binding affinity were disclosed in International Application No. PCT/US2008/064421, corresponding to International Publication No. WO/2008/144757, entitled "Novel Rabbit Antibody Humanization Methods and Humanized Rabbit Antibodies", filed May 21, 2008, the disclosure of which is herein incorporated by reference in its entirety.

Certain teachings related to producing antibodies or fragments thereof using mating competent yeast and corresponding methods were disclosed in U.S. patent application Ser. No. 11/429,053, filed May 8, 2006, (U.S. Patent Application Publication No. US2006/0270045), the disclosure of which is herein incorporated by reference in its entirety.

Certain anti-CGRP antibody polynucleotides and polypeptides are disclosed in the sequence listing accompanying this patent application filing, and the disclosure of said sequence listing is herein incorporated by reference in its entirety.

The entire disclosure of each document cited (including patents, patent applications, journal articles, abstracts, manuals, books, or other disclosures) in the Background of the Invention, Detailed Description, and Examples is herein incorporated by reference in their entireties.

The following examples are put forth so as to provide those of ordinary skill in the art with a complete disclosure and description of how to make and use the subject invention, and are not intended to limit the scope of what is regarded as the invention. Efforts have been made to ensure accuracy with respect to the numbers used (e.g. amounts, temperature, concentrations, etc.) but some experimental errors and deviations should be allowed for. Unless otherwise indicated, parts are parts by weight, molecular weight is average molecular weight, temperature is in degrees centigrade; and pressure is at or near atmospheric.

EXAMPLES

Example 1 Preparation of Antibodies that Bind CGRP

By using the antibody selection protocol described herein, one can generate an extensive panel of antibodies.

Immunization Strategy

Rabbits were immunized with human CGRPα (American Peptides, Sunnyvale Calif. and Bachem, Torrance Calif.). Immunization consisted of a first subcutaneous (sc) injection of 100 μg of antigen mixed with 100 μg of KLH in complete Freund's adjuvant (CFA) (Sigma) followed by two boosts, two weeks apart each containing 50 μg antigen mixed with 50 μg in incomplete Freund's adjuvant (IFA) (Sigma). Animals were bled on day 55, and serum titers were determined by ELISA (antigen recognition) and by inhibition of CGRP driven cAMP increase in SK-N-MC.

Antibody Selection Titer Assessment

To identify and characterize antibodies that bind to human CGRPα, antibody-containing solutions were tested by ELISA. Briefly, neutravidin coated plates (Thermo Scientific), were coated with N-term biotinylated human CGRPα (50 μL per well, 1 μg/mL) diluted in ELISA buffer (0.5% fish skin gelatin in PBS pH 7.4) either for approximately 1 hr at room temperature or alternatively overnight at 4° C. The plates were then further blocked with ELISA buffer for one hour at room temperature and washed using wash buffer (PBS, 0.05% tween 20). Serum samples tested were serially diluted using ELISA buffer. Fifty microliters of diluted serum samples were transferred onto the wells and incubated for one hour at room temperature for one hour. After this incubation, the plate was washed with wash buffer. For development, an anti-rabbit specific Fc-HRP (1:5000 dilution in ELISA buffer) was added onto the wells and incubated for 45 min at RT. After a 3× wash step with wash solution, the plate was developed using TMB substrate for two minutes at room temperature and the reaction was quenched using 0.5M HCl. The well absorbance was read at 450 nm.

Titer Determination of Serum Samples by Functional Activity (Inhibition of CGRP Driven cAMP Levels)

To identify and characterize antibodies with functional activity, an inhibition of CGRP driven increase of cAMP levels assay was done using electrochemiluminescence (Meso Scale Discovery, MSD). Briefly, antibody preparations to be tested were serially diluted in MSD assay buffer (Hepes, MgCl2, pH 7.3, 1 mg/mL blocker A, Meso Scale Discovery) in a 96 well round bottom polystyrene plate (Costar). To this plate, human CGRPα was added (10 ng/mL final concentration) diluted in MSD assay buffer and incubated for one hour at 37 C. Appropriate controls were used as suggested by the assay-kit manufacturer. Human neuroepithelioma cells (SK-N-MC, ATCC) were detached using an EDTA solution (5 mM in PBS) and washed using growth media (MEM, 100/FBS, antibiotics) by centrifugation. The cell number was adjusted to 2 million cells per mL in assay buffer, and IBMX (3-Isobutyl-1Methylxanthine, Sigma) was added to a final concentration of 0.2 mM right before loading cells onto cAMP assay plate. After the antibody human CGRPα solution was incubated for one hour 20 microliters of solution containing cells were transferred to the cAMP assay plate. All tested samples were run in duplicates with appropriate controls. Ten microliters of cells were added to the wells and the plate was incubated for 30 minutes with shaking at room temperature. While cells were being incubated with the CGRP solution, the stop solution was prepared by making a 1:200 solution of TAG labeled cAMP (MSD) in lysis buffer (MSD). To stop the cells-CGRP incubation, 20 microliters of stop solution was added to the cells and the plate was incubated for one hour with shaking at room temperature. The read buffer (MSD) was diluted four times with water and 100 microliters were added to all wells on the plate. The plate was then read using a Sector Imager 2400 (MSD) and the Prism software was used for data fit and IC50 determination.

Tissue Harvesting

Once acceptable titers were established, the rabbit(s) were sacrificed. Spleen, lymph nodes, and whole blood were harvested and processed as follows:

Spleen and lymph nodes were processed into a single cell suspension by disassociating the tissue and pushing through sterile wire mesh at 70 μm (Fisher) with a plunger of a 20 cc syringe. Cells were collected in PBS. Cells were washed twice by centrifugation. After the last wash, cell density was determined by trypan blue. Cells were centrifuged at 1500 rpm for 10 minutes; the supernatant was discarded. Cells were resuspended in the appropriate volume of 10% dimethyl sulfoxide (DMSO, Sigma) in FBS (Hyclone) and dispensed at 1 ml/vial. Vials were stored at −70° C. in a slow freezing chamber for 24 hours and stored in liquid nitrogen.

Peripheral blood mononuclear cells (PBMCs) were isolated by mixing whole blood with equal parts of the low glucose medium described above without FBS. 35 ml of the

whole blood mixture was carefully layered onto 8 ml of Lympholyte Rabbit (Cedarlane) into a 45 ml conical tube (Corning) and centrifuged 30 minutes at 2500 rpm at room temperature without brakes. After centrifugation, the PBMC layers were carefully removed using a glass Pasteur pipette (VWR), combined, and placed into a clean 50 ml vial. Cells were washed twice with the modified medium described above by centrifugation at 1500 rpm for 10 minutes at room temperature, and cell density was determined by trypan blue staining. After the last wash, cells were resuspended in an appropriate volume of 10% DMSO/FBS medium and frozen as described above.

B Cell Selection, Enrichment and Culture Conditions

On the day of setting up B cell culture, PBMC, splenocyte, or lymph node vials were thawed for use. Vials were removed from LN2 tank and placed in a 37° C. water bath until thawed. Contents of vials were transferred into 15 ml conical centrifuge tube (Corning) and 10 ml of modified RPMI described above was slowly added to the tube. Cells were centrifuged for 5 minutes at 2K RPM, and the supernatant was discarded. Cells were resuspended in 10 ml of fresh media. Cell density and viability was determined by trypan blue.

a) The Following Protocol was Used for Ab1 and Ab13

Cells were pre-mixed with the biotinylated human CGRPα as follows. Cells were washed again and resuspended at 1E07 cells/80 μL medium. Biotinylated human CGRPα was added to the cell suspension at the final concentration of 5 ug/mL and incubated for 30 minutes at 4° C. Unbound biotinylated human CGRPα was removed performing two 10 ml washes using PBF [Ca/Mg free PBS (Hyclone), 2 mM ethylenediamine tetraacetic acid (EDTA), 0.5% bovine serum albumin (BSA) (Sigma-biotin free)]. After the second wash, cells were resuspended at 1E07 cells/80 μl PBF and 20 μl of MACS® streptavidin beads (Miltenyi Biotech, Auburn Calif.) per 10E7 cells were added to the cell suspension. Cells and beads were incubated at 4° C. for 15 minutes and washed once with 2 ml of PBF per 10E7 cells.

b) The Following Protocol was Used for Ab4, Ab7, Ab9 and Ab11:

Biotinylated human CGRPα was pre-loaded onto the streptavidin beads as follows. Seventy five microliters of streptavidin beads (Milteny Biotec, Auburn Calif.) were mixed with N-terminally biotinylated huCGRPα (10 ug/ml final concentration) and 300 μl PBF. This mixture was incubated at 4° C. for 30 min and unbound biotinylated human CGRPα was removed using a MACS® separation column (Miltenyi Biotec, with a 1 ml rinse to remove unbound material. Then material was plunged out, then used to resuspend cells from above in 100 ul per 1E7 cells, the mixture was then incubated at 4° C. for 30 min and washed once with 10 ml of PBF.

For both a) and b) protocols the following applied: After washing, the cells were resuspended in 500 μl of PBF and set aside. A MACS® MS column (Miltenyi Biotec, Auburn Calif.) was pre-rinsed with 500 ml of PBF on a magnetic stand (Milteni). Cell suspension was applied to the column through a pre-filter, and unbound fraction was collected. The column was washed with 2.5 ml of PBF buffer. The column was removed from the magnet stand and placed onto a clean, sterile 1.5 ml eppendorf tube. 1 ml of PBF buffer was added to the top of the column, and positive selected cells were collected. The yield and viability of positive cell fraction was determined by trypan blue staining. Positive selection yielded an average of 1% of the starting cell concentration.

A pilot cell screen was established to provide information on seeding levels for the culture. Plates were seeded at 10, 25, 50, 100, or 200 enriched B cells/well. In addition, each well contained 50K cells/well of irradiated EL-4.B5 cells (5,000 Rads) and an appropriate level of activated rabbit T cell supernatant (See U.S. Patent Application Publication No. 20070269868) (ranging from 1-5% depending on preparation) in high glucose modified RPMI medium at a final volume of 250 μl/well. Cultures were incubated for 5 to 7 days at 37° C. in 4% CO_2.

B-Cell Culture Screening by Antigen-Recognition (ELISA)

To identify wells producing anti-human CGRPα antibodies, the same protocol as described for titer determination of serum samples by antigen-recognition (ELISA) was used with the following changes. Briefly, neutravidin coated plates were coated with a mixture of both N- and C-terminally biotinylated human CGRPα (50 μL per well, 1 μg/mL each). B-cell supernatant samples (50 μL) were tested without prior dilution.

Identification of Functional Activity in B-Cell Supernatants Using CGRP Driven cAMP Production

To determine functional activity contained in B-cell supernatants, a similar procedure to that described for the determination of functional titer of serum samples was used with the following modifications. Briefly, B-cell supernatant (20 μL) were used in place of the diluted polyclonal serum samples.

Isolation of Antigen-Specific B-Cells

Plates containing wells of interest were removed from −70° C., and the cells from each well were recovered using five washes of 200 microliters of medium (10% RPMI complete, 55 μM BME) per well. The recovered cells were pelleted by centrifugation and the supernatant was carefully removed. Pelleted cells were resuspended in 100 μl of medium. To identify antibody expressing cells, streptavidin coated magnetic beads (M280 dynabeads, Invitrogen) were coated with a combination of both N- and C-terminal biotinylated human CGRPα. Individual biotinylated human CGRPα lots were optimized by serial dilution. One hundred microliters containing approximately 4×10E7 coated beads were then mixed with the resuspended cells. To this mixture 15 microliters of goat anti-rabbit H&L IgG-FITC (Jackson Immunoresearch) diluted 1:100 in medium were added.

Twenty microliters of cell/beads/anti-rabbit H&L suspension were removed and 5 microliter droplets were dispensed on a one-well glass slide previously treated with Sigmacote (Sigma) totaling 35 to 40 droplets per slide. An impermeable barrier of paraffin oil (JT Baker) was used to submerge the droplets, and the slide was incubated for 90 minutes at 37° C. in a 4% CO2 incubator in the dark.

Specific B cells that produce antibody can be identified by the fluorescent ring around produced by the antibody secretion, recognition of the bead-associated biotinylated antigen, and subsequent detection by the fluorescent-IgG detection reagent. Once a cell of interest was identified it was recovered via a micromanipulator (Eppendorf). The single cell synthesizing and exporting the antibody was transferred into a microcentrifuge tube, frozen using dry ice and stored at −70° C.

Amplification and Sequence Determination of Antibody Sequences from Antigen-Specific B Cells

Antibody sequences were recovered using a combined RT-PCR based method from a single isolated B-cell. Primers containing restriction enzymes were designed to anneal in conserved and constant regions of the target immunoglobulin genes (heavy and light), such as rabbit immunoglobulin sequences, and a two-step nested PCR recovery was used to

amplify the antibody sequence. Amplicons from each well were analyzed for recovery and size integrity. The resulting fragments are then digested with AluI to fingerprint the sequence clonality. Identical sequences displayed a common fragmentation pattern in their electrophoretic analysis. The original heavy and light chain amplicon fragments were then digested using the restriction enzyme sites contained within the PCR primers and cloned into an expression vector. Vector containing subcloned DNA fragments were amplified and purified. Sequence of the subcloned heavy and light chains were verified prior to expression.

Recombinant Production of Monoclonal Antibody of Desired Antigen Specificity and/or Functional Properties

To determine antigen specificity and functional properties of recovered antibodies from specific B-cells, vectors driving the expression of the desired paired heavy and light chain sequences were transfected into HEK-293 cells.

Antigen-Recognition of Recombinant Antibodies by ELISA

To characterize recombinant expressed antibodies for their ability to bind to human-CGRPα, antibody-containing solutions were tested by ELISA. All incubations were done at room temperature. Briefly, Immulon IV plagtes (Thermo Scientific), were coated with a CGRPα containing solution (1 ut/mL in PBS) for 2 hours. CGRPα-coated plates were then washed three times in wash buffer (PBS, 0.05% Tween-20). The plates were then blocked using a blocking solution (PBS, 0.5% fish skin gelatin, 0.05% Tween-20) for approximately one hour. The blocking solution was then removed and the plates were then incubated with a dilution series of the antibody being tested for approximately one hour. At the end of this incubation, the plate was washed three times with wash buffer and further incubated with a secondary antibody containing solution (Peroxidase conjugated affinipure $F(ab')_2$ fragment goat anti-human IgG, Fc fragment specific (Jackson Immunoresearch) for approximately 45 minutes and washed three times. At that point a substrate solution (TMB peroxidase substrate, BioFx) and incubated for 3 to 5 minutes in the dark. The reaction was stopped by addition of a HCl containing solution (0.5M) and the plate was read at 450 nm in a plate-reader.

Results: FIGS. **15-18** demonstrate that anti-CGRP antibodies Ab1-Ab14 bind to and recognize CGRPα.

Functional Characterization of Recombinant Antibodies by Modulation of CGRP Driven Intracellular cAMP Levels and Cross Reactivity to Rats

To characterize recombinant expressed antibody for their ability to inhibit CGRPα mediated increased cellular levels of cAMP assay, an electrochemiluminescence assay-kit (Meso Scale Discovery, MSD) was used. Briefly, antibody preparations to be tested were serially diluted in MSD assay buffer (Hepes, MgCl2, pH 7.3, 1 mg/mL blocker A, Meso Scale Discovery) in a 96 well round bottom polystyrene plate (Costar). To this plate, human CGRPα was added (25 ng/mL final concentration) diluted in MSD assay buffer and incubated for one hour at 37° C. Appropriate controls were used as suggested by the assay-kit manufacturer. Human neuroepithelioma cells (SK-N-MC, ATCC) were detached using an EDTA solution (5 mM) and washed using growth media (MEM, 10% FBS, antibiotics) by centrifugation. The cell number was adjusted to 2 million cells per mL in assay buffer, and IBMX (3-Isobutyl-1Methylxanthine, 50 mM Sigma) was added to a final concentration of 0.2 mM right before loading cells onto cAMP assay plate. The antibody human CGRPα solution was incubated for one hour after which 20 microliters of solution containing cells were transferred to the cAMP assay plate. All tested samples were run in duplicates with appropriate controls. Ten microliters of cells were added to the wells and the plate was incubated for 30 minutes with shaking. While cells were being incubated with the CGRP solution, the stop solution was prepared by making a 1:200 solution of TAG labeled cAMP (MSD) in lysis buffer (MSD). To stop the cells-CGRP incubation, 20 microliters of stop solution was added to the cells and the plate was incubated for one hour with shaking. The read buffer (MSD) was diluted four times with water and 100 microliters were added to all wells on the plate. The plate was then read using a Sector Imager 2400 (MSD) and the Prism software was used for data fit and IC50 determination.

To test for the ability of recombinant antibodies to antagonize human CGRPβ a similar assay was performed with the substitution of the CGRP agonist (CGRPβ 10 ng/mL final concentration). Evaluation of the recombinant antibodies to recognize and inhibit rat CGRP-mediated cAMP generation was conducted using rat CGRP (5 ng/mL final concentration) and the rat L6 cell line (ATCC).

Results: FIGS. **19-37** demonstrate that anti-CGRP antibodies Ab1-Ab14 inhibit CGRPα, CGRPβ, and rat CGRP mediated increased cellular levels of cAMP.

Example 2: Enzymatic Production of Fab Fragments

Papain digestions were conducted using immobilized papain (Thermo/Pierce) as per manufacturer's instructions. Briefly, purified antibodies were incubated in a cystein/HCl-containing buffer with immobilized papain at 37° C. with gentle rocking. The digestion was monitored by taking an aliquot and analyzing using SDS-PAGE for cleavage of the heavy chain. To stop the reaction, the immobilized papain was spun out and washed using 50 mM Tris pH 7.5 and filtered. Undigested full length antibody and Fc fragments were removed by using a MabSelectSure (GE) column.

Example 3 Yeast Cell Expression

Construction of *Pichia pastoris* Expression Vectors for Heavy and Light Chain.

The humanized light and heavy chain fragments were commercially synthesized and subcloned into a pGAP expression vector. The pGAP expression vector uses the GAP promoter to drive expression of the immunoglobulin chain and the human serum albumin (HSA) leader sequence for export. In addition, this vector contains common elements such as a bacterial origin of replication, and a copy of the kanamycin resistance gene which confers resistance to the antibiotic G418 in *P. pastoris*. G418 provides a means of selection for strains that contain the desired expression vector integrated into their genome.

Transformation of Expression Vectors into Haploid Met1 and Lys3 Host Strains of *Pichia pastoris*

All methods used for transformation of haploid *P. pastoris* strains and manipulation of the *P. pastoris* sexual cycle were done as described in *Pichia* Protocols (Methods in Molecular Biology Higgings, D R, and Cregg, J M, Eds. 1998. Humana Press, Totowa, N.J.). Prior to transformation each vector was linearized within the GAP promoter sequences to direct the integration of the vector into the GAP promoter locus of the *P. pastoris* genome. Haploid strains were transfected using electroporation and successful transformants were selected on YPDS (yeast extract, peptone dextrose with sorbitol) G418 agar plates. Copy numbers of heavy and light chain genes were determined for haploid strains by Southern blot analysis. Haploid strains were then

mated and selected for their ability to grow in the absence of the amino acid markers (i.e., Lys and Met). Resulting diploid clones were then subjected to a final Southern blot to confirm copy numbers of heavy and light chain genes. A clone expressing the antibody of interest was selected using biolayer interferometry Protein-A biosensors to monitor expression (Octet, ForteBio).

Example 4 Expression of Ab3, Ab6 and Ab14 in *Pichia pastoris*

Three *Pichia* strains for expression of full-length antibody were made. For all the full length antibody expressing strains, haploids strains were created and subsequently mated. One haploid strain expressed full-length light chain sequence and another haploid strain expressed the full-length heavy chain sequence. Each diploid strain was used to generate a research cell bank and used for expression in a bioreactor.

First an inoculum was expanded using the research cell bank using medium comprised of the following nutrients (% w/v): yeast extract 3%, anhydrous dextrose 4%, YNB 1.34%, Biotin 0.004% and 100 mM potassium phosphate. To generate the inoculum for the fermenters, the cell bank was expanded for approximately 24 hours in a shaking incubator at 30° C. and 300 rpm. A 10% inoculum was then added to Labfors 2.5 L working volume vessels containing 1 L sterile growth medium. The growth medium was comprised of the following nutrients: potassium sulfate 18.2 g/L, ammonium phosphate monobasic 36.4 g/L, potassium phosphate dibasic 12.8 g/L, magnesium sulfate heptahydrate 3.72 g/L, sodium citrate dihydrate 10 g/L, glycerol 40 g/L, yeast extract 30 g/L, PTM1 trace metals 4.35 mL/L, and antifoam 204 1.67 mL/L. The PTM1 trace metal solution was comprised of the following components: cupric sulfate pentahydrate 6 g/L, sodium iodide 0.08 g/L, manganese sulfate hydrate 3 g/L, sodium molybdate dihydrate 0.2 g/L, boric acid 0.02 g/L, cobalt chloride 0.5 g/L, zinc chloride 20 g/L, ferrous sulfate heptahydrate 65 g/L, biotin 0.2 g/L, and sulfuric acid 5 mL/L.

The bioreactor process control parameters were set as follows: Agitation 1000 rpm, airflow 1.35 standard liter per minute, temperature 28° C. and pH was controlled at six using ammonium hydroxide. No oxygen supplementation was provided.

Fermentation cultures were grown for approximately 12 to 16 hours until the initial glycerol was consumed as denoted by a dissolved oxygen spike. The cultures were starved for approximately three hours after the dissolved oxygen spike. After this starvation period, a bolus addition of ethanol was added to the reactor to reach 1% ethanol (w/v). The fermentation cultures were allowed to equilibrate for 15 to 30 minutes. Feed addition was initiated 30 minutes post-ethanol bolus and set at a constant rate of 1 mL/min for 40 minutes, then the feed pump was controlled by an ethanol sensor keeping the concentration of ethanol at 1% for the remainder of the run using an ethanol sensing probe (Raven Biotech). The feed was comprised of the following components: yeast extract 50 g/L, dextrose 500 g/L, magnesium sulfate heptahydrate 3 g/L, and PTM1 trace metals 12 mL/L. For fermentation of the full length Ab6 and Ab14, sodium citrate dihydrate (0.5 g/L) was also added to the feed. The total fermentation time was approximately 90 hours.

Example 5 Methods of Humanizing Antibodies

Methods of humanizing antibodies have been described previously in issued U.S. Pat. No. 7,935,340, the disclosure of which is incorporated herein by reference in its entirety. In some instances, a determination of whether additional rabbit framework residues are required to maintain activity is necessary. In some instances the humanized antibodies still requires some critical rabbit framework residues to be retained to minimize loss of affinity or activity. In these cases, it is necessary to change single or multiple framework amino acids from human germline sequences back to the original rabbit amino acids in order to have desired activity. These changes are determined experimentally to identify which rabbit residues are necessary to preserve affinity and activity. This is now the end of the variable heavy and light chain humanized amino acid sequence.

Example 6 Inhibition of CGRP Binding to its Cellular Receptor

To characterize recombinantly expressed antibodies for their ability to inhibit CGRP binding to its cellular receptor, a radioligand-binding assay was performed as previously described [Elshourbagy et al, Endocrinology 139:1678 (1998); Zimmerman et al, Peptides, 16:421 (1995)]. Membrane preparations of recombinant human CGRP receptors, calcitonin receptor-like receptor and RAMP1 (Chemiscreen, Millipore) were used. Antibody dilutions were preincubated with ^{125}I radiolabeled human CGRPα (0.03 nM) for 30 minutes at room temperature. Non-specific binding was estimated in the presence of 0.1 μM human CGRPα. Membranes were filtered and washed. The filters were then counted to determine ^{125}I radiolabeled human CGRPα specifically bound.

Results: FIG. **38** demonstrates that anti-CGRP antibodies Ab1-Ab13 inhibit CGRP binding to its cellular receptor.

Example 7 Inhibition of Neurogenic Vasodilation by Anti-CGRP Antibodies in Rats

CGRP is a potent vasodilator (Nature 313: 54-56 (1985) and Br J. Clin. Pharmacol. 26(6):691-5. (1988)). A pharmacodynamic assay to measure CGRP receptor antagonist activity non-invasively was used to characterize anti-CGRP antibodies. The model relied on changes in dermal blood flow measured using a laser Doppler imaging following the topical application of a capsaicin solution. Capsaicin activates the transient receptor potential vanilloid type 1 receptor (TRPV-1), producing neurogenic inflammation and vasodilatation via the local release of vasoactive mediators including CGRP and substance P (Br. J. Pharmacol. 110: 772-776 (1993)).

On the day prior to the vasodilatation assay, animals were dosed with the test agent or control via IP (intraperitoneal). Following dosing, the animals were shaved and depilated in the lower back region of their dorsal side, in an area approximately 2×6 cm. The animals were then returned to their cages overnight. On the day of test, approximately 24 hours post dosing, animals were anesthetized with isoflurane gas and placed on a temperature controlled heating pad and fitted with a nose cone for continuous delivery of isoflurane. A laser doppler imager was used for the observation of vasodilatation. A beam of coherent red light generated by a 633 nm helium-neon laser was directed to the shaved area, a rectangle (2×6 cm), and scanned at a medium resolution mode. A baseline Doppler scan was obtained first and the location of O-ring placement predetermined by identifying two similar low flux areas. Two rubber Orings (~1 cm in diameter) were placed in the selected regions and a baseline scan was performed. Immediately after completion of the

scan, 1 mg of capsaicin in 5 μL of an ethanol:acetone solution (1:1) was applied within each of the two O-rings Doppler scans were repeated at 2.5, 5, 7.5, 10, 12.5, 15, 17.5, 20, 22.5, 25, 27.5 and 30 minutes after the application of capsaicin. Percent change from baseline mean Flux within each of the two O-rings, was plotted as the results of vasodilatation due to capsaicin.

In order to test recombinantly expressed antibodies for their ability to inhibit CGRP binding to its cellular receptor, a radioligand-binding assay was performed as previously described.

Results: FIGS. **39** and **40** demonstrates that anti-CGRP antibodies Ab3 and Ab6 reduced vasodilation in this model following capsaicin administration.

Example 8 Inhibition of Light Aversion or Photophobia by Systemic (IP) Injection of Anti-CGRP Antibody in Transgenic Nestin/Ramp1 Mice

As discussed supra, one of the hallmarks of migraines is photophobia, or increased sensitivity to light [Mulleners et al, Headache 41: 31-39 (2001); Recober et al, J. Neuroscience 29:8798:8804 (2009)]. It is also known that migraineurs, but not non-migraineurs, are sensitive to CGRP-induced headache [reviewed in Neurology 22:241-246 (2009)]. CGRP binds to a G protein coupled receptor called CLR (calcitonin like receptor) that works concomitantly with the receptor activity-modifying protein 1 (RAMP1) in mediating CGRP binding and signaling. In-vitro, the activity of CGRP is strongly enhanced by over-expression of the RAMP1 subunit of the CGRP receptor [(J. Neurosci. 27:2693-2703 (2007)]. To study light aversion behavior in mice, a nestin/human-RAMP1 transgenic mouse model was developed [Recober et al, J. Neuroscience 29: 8798-8804 (2009); Russo et al, Mol. Cell. Pharmacol., 1:264-270 (2009)]. These mice when exposed to CGRP present symptoms associated with migraines in particular light aversion (ibid). This protocol is detailed below.

To test the ability of anti-CGRP antibodies to block CGRP-induced light aversion or photophobia, mice are housed under standard conditions in groups of 2-5 per cage with a 12 hour light cycle (lights on at 0500 CST)/0600 CDT and off at 1700 CST/1800 CDT) and access to water and food ad libitum. The mice used in the studies are comprised in mice colonies of genotype nestin/hRAMP1 that contain two transgene alleles Tg(Nes-cre)1Kln/J and Tg(RAMP1) alleles (B6;SJL-Tg(Nes-cre)1Kln Tg(RAMP1). Nes-cre was introduced in these mice by an intercross involving mice obtained from The Jackson Laboratory (stock 003771) on a B6 genetic background yielding mice.

The control mice used in the protocol are littermates that are either non-transgenic, or single transgenic (not expressing hRAMP1) containing either transgene: nestin-cre or Cx1-GFP-hRAMP1. The stock colony is maintained by backcrossing CX1-GFP-hRAMP1 mice with non-transgenic littermates in the barrier facility. For behavior studies, the colony is maintained by crossing CX1-GFP-hRAMP1 single transgenic with nestin-cre mice in non-barrier facilities. All of these mice are cared for by animal care and procedures approved by the University of Iowa Animal Care and Use Committee and further are performed in accordance with the standard set by the National Institutes of Health.

The materials and equipment used in this protocol include a light-dark box and testing chambers comprising a plexiglass open field (27×27×20.3 cm) containing 16 beam infrared arrays (Med Associates Inc., St. Albans, Vt.). The light/dark box is divided in two equally sized zones by a dark insert that is opaque to visible light. There is an opening (5.2×6.8 cm) in the dark insert that allows the mouse to freely move between the two zones. This testing chamber is placed inside a sound-attenuating cubicle (56×38×36 cm) with a fan for ventilation (Med Associates Inc.). There are six chambers for the overall system that integrates with a computer containing software for recording and data collection (Med Associates Inc.).

The software used to monitor results is Activity Monitor v 6.02 (Med Associates Inc.). The software settings used for recording comprise: Resolution (ms): 50, Box Size: 3, Resting Delay (ms): 500, Ambulatory Trigger: 3, Session Type: C, Session Time (min): 20, Block Interval (sec): 300, and Compressed File: DEFAULT.ZIP.

In the protocol, the light source for each chamber is an LED panel, which was installed to the ceiling of the sound-attenuating cubicle. The LED panel contains 36 collimated—1 watt LED bulbs (5500 k Daylight White) (LED-wholesalers, Burlingame, Calif.). To control light intensity, each LED panel is connected to a dimmable LED driver (LINEARdrive; eldoLED America Inc., San Jose, Calif.) leading to a potential range of light intensity from ~300 to 27,000 lux. The standard light intensity is ~1000-1200 lux unless otherwise stated. Alternatively, lower light intensities have been achieved by using layers of wax paper to filter the light leading to an intensity of ~55 lux.

The injectors used are hand-made by inserting a stripped 30 gauge×½" needle into non-radiopaque polyethylene tubing (inner diameter 0.38 mm; outer diameter 1.09 mm). Using the tubing described above, a stopper (~1 cm in length) is placed over the needle leaving approximately 2.5 mm of the bevel uncovered. These injectors are connected to a 10 μL Hamilton syringe.

The mice are injected with rat α-CGRP (Sigma) diluted in Dulbecco phosphate-buffered saline (D-PBS) (Hyclone). The total dose delivery is 0.5 nmol. For example, 250 or 500 μg CGRP is diluted in 250 or 500 μL sterile PBS for a final concentration of 1 μg/μL. The CGRP is stored at −20° C. and aliquots are freeze-thawed at most one time. The PBS is stored at 4° C.

The mice are administered one of the anti-CGRP antibodies disclosed herein (Ab3), vehicle or a control antibody, which are stored at 4° C. prior to administration. In this protocol prior to the administration of the CGRP i.e., approximately 24 hours prior to testing, the mice are weighed and then receive an intraperitoneal (ip) injection of either: vehicle, control antibody, or CGRP-binding antibody at a dosage of 30 mg/kg. The mice are also screened to detect any abnormal physical conditions that could affect the assay such as a missing eye, cataracts, or other abnormalities such as grooming, etc. The day after antibody administration, mice are transported in cages from animal housing on a cart and then the mice are placed in the behavior room for acclimation at least 1 hour prior to any injection or testing. Any coverings required for transport are removed from the cages and normal light conditions (standard overhead fluorescent lighting) are turned on during acclimation and remain on for the remainder of the procedure. In addition, all equipment that produces sound including anesthetic devices, light/dark chambers, and LED panels are turned on during acclimation and remain until testing is complete. Typically there is minimal human presence in the room during acclimation.

After acclimation each mouse is placed in an induction chamber and administered 3.5% isoflurane. After the mouse is anesthetized, it is transferred to a nose cone maintaining

3.5% isoflurane administration, so that it remains anesthetized during injection. Thereafter drug administration is effected using the injector by direct injection into the right lateral ventricle through the intact scalp aiming at 1 mm posterior to bregma and 1 mm right from the midline.

Typically for consistency all the injections are performed by the same person after a period of training yielding a success rate of >90% as demonstrated by injections of dye into the ventricles. The drugs injected are either 2.0 μL vehicle (D-PBS) μL or 2.0 μg CGRP in 2.0 μL vehicle (1 μg/μL) administered as a direct intracerebroventricular injection into the right lateral ventricle of the brain through the intact scalp aiming at 1 mm posterior to bregma and 1 mm right from the midline as described before [Recober et al, J. Neuroscience 29: 8798-8804 (2009)] After all 2.0 μL is delivered, the needle remains in place for 10 sec and then removed. The time of injection is then recorded.

After injection the mice are allowed to recover for 30 minutes prior to testing in an empty, uncovered cage containing a paper towel for bedding. During recovery, the following is recorded: diarrhea, excessive urination, bleeding post-injection, abnormal behavior such as lack of movement, seizures, etc. After a 30 minute recovery testing is effected. Each mouse is placed along the back wall (furthest from the opening between the two zones) in the light zone approximately in the center. This triggers the recording to begin. Up to six mice are tested at one time (one mouse per chamber). During testing the shelf with the chamber is pushed back into the cabinet and the doors closed. The software records mouse movement for 20 minutes. After the recording is completed, each mouse is removed and placed back in home cage for transport back to animal housing.

Results

Using this protocol an anti-CGRP antibody developed by Alder Biopharmaceuticals (Ab3) was tested to determine its potential suitability for treating migraine, particularly chronic migraine in human subjects and more particularly for treatment or prevention of CGRP-associated photophobia. The results of these studies are shown in FIG. **41** and FIG. **42**. FIG. **41** contains data that compares the effect of ICV injection of CGRP in hRAMP1 tg mice and control littermate mice. The data reveals that the CGRP administration results in decreased time in light behavior in the hRAMP1 tg mice relative to their control littermates.

FIG. **42** contains data which compares the effect of systemic (IP) injection of anti-CGRP antibody (Ab3) in vehicle, vehicle alone, and control antibody in vehicle in nestin/RAMP1 mice which are administered these moieties intraperitoneally at 30 mg/kg about 24 hours prior to administration of CGRP. The data in the left side of the graph is the total time in light (seconds) for the first 10 minutes, and the data on the right side of the graph is the total time in light (seconds) for the first 20 minutes measured after CGRP injection (administered via ICV injection) and the recovery period. Light intensity in light zone was approximately 1×10^3 lx. The data reveal that the mice who received the anti-CGRP antibody Ab3 according to the invention had a statistically significant increase in the amount of time spent in the light relative to the mice who received the controls.

These results indicate that Ab3 inhibits CGRP-associated photophobia or light aversion and should be well suited for treating migraine or other disorders that involve photophobia, especially CGRP related photophobia. Based it is anticipated that other anti-CGRP antibodies including others disclosed herein may behave similarly. These results further indicate that the subject light aversion behavior assay may be used to assess the potential therapeutic efficacy (ability to antagonize effects of CGRP in vivo) of candidate of anti-CGRP antibodies and antibody fragments. This was unanticipated as it was unforeseeable that a large polypeptide such as an anti-CGRP antibody would go through the blood-brain barrier and inhibit photophobia or light aversion.

The results reveal that the excess CGRP that induces light aversive behavior in mice is reduced by the systemic administration of anti-CGRP antibody suggesting that the antibody is able to bind a sufficient amount of the circulating CGRP to counteract the light aversive behavior. These results suggest that the anti-CGRP antibody may be crossing the blood-brain barrier and thereby inhibiting the neurological effects of CGRP, in particular migraine associated photophobia and pain.

This is the first demonstration that the subject animal light aversive behavior assay may be used to assess therapeutic efficacy of a polypeptide such as an anti-CGRP antibody or anti-CGRP antibody fragment. In addition these results suggest that this animal model potentially may be useful in determining effective dosages of a candidate anti-CGRP antibody or antibody fragment, effective modes of administration, as well as a suitable dosage regimen.

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Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Arg Ala Ser Ser Thr Thr Val Asp Leu Lys Met
65                  70                  75                  80

Thr Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser
            100                 105


<210> SEQ ID NO 4
<211> LENGTH: 439
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 4

Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr Pro
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Val Ser Gly Leu Asp Leu Ser Ser Tyr Tyr
            20                  25                  30

Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Arg Ala Ser Ser Thr Thr Val Asp Leu Lys Met
65                  70                  75                  80

Thr Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr
            100                 105                 110

Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser
        115                 120                 125

Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu
    130                 135                 140

Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His
145                 150                 155                 160

Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser
                165                 170                 175

Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys
            180                 185                 190

Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu
        195                 200                 205

Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro
    210                 215                 220

Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys
225                 230                 235                 240

Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val
                245                 250                 255
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Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp
            260                 265                 270

Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr
        275                 280                 285

Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp
    290                 295                 300

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu
305                 310                 315                 320

Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg
                325                 330                 335

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys
            340                 345                 350

Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp
        355                 360                 365

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys
    370                 375                 380

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser
385                 390                 395                 400

Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser
                405                 410                 415

Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser
            420                 425                 430

Leu Ser Leu Ser Pro Gly Lys
        435


<210> SEQ ID NO 5
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 5

Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 6
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 6

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 7
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 7

Leu Gly Ser Tyr Asp Cys Ser Ser Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 8
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 8

Ser Tyr Tyr Met Gln
1               5
```

<210> SEQ ID NO 9
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 9

Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 10
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 10

Gly Asp Ile
1

<210> SEQ ID NO 11
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 11

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Ser Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg

<210> SEQ ID NO 12
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 12

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

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Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Ser Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215


<210> SEQ ID NO 13
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 13

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Leu Asp Leu Ser Ser Tyr
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Gly Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
            100                 105                 110


<210> SEQ ID NO 14
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 14

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Leu Asp Leu Ser Ser Tyr
            20                  25                  30
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Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Gly Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
            100                 105                 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
        115                 120                 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
    130                 135                 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145                 150                 155                 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
                165                 170                 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
            180                 185                 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
        195                 200                 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440
```

<210> SEQ ID NO 15
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 15

Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn Tyr Leu Ala
1 5 10

<210> SEQ ID NO 16
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 16

Ser Thr Ser Thr Leu Ala Ser
1 5

<210> SEQ ID NO 17
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 17

Leu Gly Ser Tyr Asp Cys Ser Ser Gly Asp Cys Phe Val
1 5 10

<210> SEQ ID NO 18
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 18

Ser Tyr Tyr Met Gln
1 5

<210> SEQ ID NO 19
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 19

Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 20
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 20

Gly Asp Ile
1

<210> SEQ ID NO 21
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
polypeptide

<400> SEQUENCE: 21

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

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Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
    50                  55                  60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Ser Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg
```

<210> SEQ ID NO 22
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 22

```
Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1               5                   10                  15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
    50                  55                  60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Ser Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215
```

<210> SEQ ID NO 23
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence

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<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 23

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Leu Asp Leu Ser Ser Tyr
20 25 30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
100 105 110

<210> SEQ ID NO 24
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 24

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Leu Asp Leu Ser Ser Tyr
20 25 30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
100 105 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
115 120 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
130 135 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145 150 155 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
165 170 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
180 185 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Ala Arg
195 200 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
210 215 220

```
Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 25
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 25

Gln Ala Ser Gln Ser Val Tyr Asp Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 26
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 26

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 27
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 27

Leu Gly Ser Tyr Asp Cys Ser Ser Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 28
<211> LENGTH: 5
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-continued

<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 28

Ser Tyr Tyr Met Gln
1 5

<210> SEQ ID NO 29
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 29

Val Ile Gly Ile Asn Asp Asn Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 30
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 30

Gly Asp Ile
1

<210> SEQ ID NO 31
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 31

Gln Val Leu Thr Gln Thr Pro Ser Pro Val Ser Ala Ala Val Gly Ser
1 5 10 15

Thr Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr His Asn Thr
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
35 40 45

Ile Tyr Asp Ala Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Gly Val Gln
65 70 75 80

Cys Asn Asp Ala Ala Ala Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Thr
85 90 95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
100 105 110

Arg

<210> SEQ ID NO 32
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 32

Gln Val Leu Thr Gln Thr Pro Ser Pro Val Ser Ala Ala Val Gly Ser
1 5 10 15

Thr Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr His Asn Thr

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            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Asp Ala Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
    50                  55                  60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Gly Val Gln
65                  70                  75                  80

Cys Asn Asp Ala Ala Ala Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Thr
                85                  90                  95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215


<210> SEQ ID NO 33
<211> LENGTH: 109
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 33

Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr Pro
1               5                   10                  15

Leu Thr Leu Thr Cys Ser Val Ser Gly Ile Asp Leu Ser Gly Tyr Tyr
            20                  25                  30

Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Lys Thr Ser Ser Thr Thr Val Asp Leu Lys Met
65                  70                  75                  80

Thr Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser
            100                 105


<210> SEQ ID NO 34
<211> LENGTH: 439
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide
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<400> SEQUENCE: 34

```
Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr Pro
1               5                  10                  15

Leu Thr Leu Thr Cys Ser Val Ser Gly Ile Asp Leu Ser Gly Tyr Tyr
            20                  25                  30

Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Lys Thr Ser Ser Thr Thr Val Asp Leu Lys Met
65                  70                  75                  80

Thr Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr
            100                 105                 110

Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser
        115                 120                 125

Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu
    130                 135                 140

Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His
145                 150                 155                 160

Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser
                165                 170                 175

Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys
            180                 185                 190

Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu
        195                 200                 205

Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro
    210                 215                 220

Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys
225                 230                 235                 240

Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val
                245                 250                 255

Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp
            260                 265                 270

Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr
        275                 280                 285

Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp
    290                 295                 300

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu
305                 310                 315                 320

Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg
                325                 330                 335

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys
            340                 345                 350

Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp
        355                 360                 365

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys
    370                 375                 380

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser
385                 390                 395                 400

Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser
                405                 410                 415
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Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser
420 425 430

Leu Ser Leu Ser Pro Gly Lys
435

<210> SEQ ID NO 35
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 35

Gln Ala Ser Gln Ser Val Tyr His Asn Thr Tyr Leu Ala
1 5 10

<210> SEQ ID NO 36
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 36

Asp Ala Ser Thr Leu Ala Ser
1 5

<210> SEQ ID NO 37
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 37

Leu Gly Ser Tyr Asp Cys Thr Asn Gly Asp Cys Phe Val
1 5 10

<210> SEQ ID NO 38
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 38

Gly Tyr Tyr Met Asn
1 5

<210> SEQ ID NO 39
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 39

Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 40
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 40

Gly Asp Ile
1

<210> SEQ ID NO 41
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 41

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr His Asn Thr
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Asp Ala Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Thr
85 90 95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg

<210> SEQ ID NO 42
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 42

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr His Asn Thr
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Asp Ala Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Thr
85 90 95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
115 120 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
130 135 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145 150 155 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
165 170 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
180 185 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
195 200 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys

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210 215

<210> SEQ ID NO 43
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 43

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Leu Ser Gly Tyr
20 25 30

Tyr Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
100 105 110

<210> SEQ ID NO 44
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 44

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Leu Ser Gly Tyr
20 25 30

Tyr Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
100 105 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
115 120 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
130 135 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145 150 155 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
165 170 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr

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           180                 185                 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
        195                 200                 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 45
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 45

Gln Ala Ser Gln Ser Val Tyr His Asn Thr Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 46
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 46

Asp Ala Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 47
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 47
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Leu Gly Ser Tyr Asp Cys Thr Asn Gly Asp Cys Phe Val
1 5 10

<210> SEQ ID NO 48
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 48

Gly Tyr Tyr Met Asn
1 5

<210> SEQ ID NO 49
<211> LENGTH: 15
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 49

Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 50
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 50

Gly Asp Ile
1

<210> SEQ ID NO 51
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 51

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr His Asn Thr
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Asp Ala Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Thr
85 90 95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg

<210> SEQ ID NO 52
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 52

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr His Asn Thr
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Asp Ala Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Thr
85 90 95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
115 120 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
130 135 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145 150 155 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
165 170 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
180 185 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
195 200 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215

<210> SEQ ID NO 53
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 53

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Leu Ser Gly Tyr
20 25 30

Tyr Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
100 105 110

<210> SEQ ID NO 54

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<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 54

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Leu Ser Gly Tyr
20 25 30

Tyr Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
100 105 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
115 120 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
130 135 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145 150 155 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
165 170 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
180 185 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Ala Arg
195 200 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
210 215 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225 230 235 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
245 250 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
260 265 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
275 280 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
290 295 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305 310 315 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
325 330 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
340 345 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
355 360 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn

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    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 55
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 55

Gln Ala Ser Gln Ser Val Tyr His Asn Thr Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 56
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 56

Asp Ala Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 57
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 57

Leu Gly Ser Tyr Asp Cys Thr Asn Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 58
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 58

Gly Tyr Tyr Met Asn
1               5


<210> SEQ ID NO 59
<211> LENGTH: 15
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 59

Ile Gly Ile Asn Gly Ala Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1               5                   10                  15


<210> SEQ ID NO 60
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 60

Gly Asp Ile
1
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<210> SEQ ID NO 61
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 61

Gln Val Leu Thr Gln Thr Ala Ser Pro Val Ser Ala Ala Val Gly Ser
1 5 10 15

Thr Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asn Tyr Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Ser Ser Arg Phe Lys
50 55 60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Asp Val Gln
65 70 75 80

Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Thr Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
100 105 110

Arg

<210> SEQ ID NO 62
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 62

Gln Val Leu Thr Gln Thr Ala Ser Pro Val Ser Ala Ala Val Gly Ser
1 5 10 15

Thr Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asn Tyr Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Ser Ser Arg Phe Lys
50 55 60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Asp Val Gln
65 70 75 80

Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Thr Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
100 105 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
115 120 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
130 135 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145 150 155 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
165 170 175

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Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215


<210> SEQ ID NO 63
<211> LENGTH: 110
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 63

Gln Glu Gln Leu Lys Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr
1               5                   10                  15

Ser Leu Thr Leu Thr Cys Thr Val Ser Gly Ile Asp Leu Ser Asn His
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile
        35                  40                  45

Gly Val Val Gly Ile Asn Gly Arg Thr Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Thr Ser Ser Thr Thr Val Asp Leu Lys
65                  70                  75                  80

Met Thr Arg Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg
                85                  90                  95

Gly Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser
            100                 105                 110


<210> SEQ ID NO 64
<211> LENGTH: 440
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 64

Gln Glu Gln Leu Lys Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr
1               5                   10                  15

Ser Leu Thr Leu Thr Cys Thr Val Ser Gly Ile Asp Leu Ser Asn His
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile
        35                  40                  45

Gly Val Val Gly Ile Asn Gly Arg Thr Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Thr Ser Ser Thr Thr Val Asp Leu Lys
65                  70                  75                  80

Met Thr Arg Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg
                85                  90                  95

Gly Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser Ala Ser
            100                 105                 110

Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr
        115                 120                 125

Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro
    130                 135                 140
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Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val
145                 150                 155                 160

His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser
                165                 170                 175

Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile
            180                 185                 190

Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val
        195                 200                 205

Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala
    210                 215                 220

Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro
225                 230                 235                 240

Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val
                245                 250                 255

Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val
            260                 265                 270

Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln
        275                 280                 285

Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln
    290                 295                 300

Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala
305                 310                 315                 320

Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro
                325                 330                 335

Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr
            340                 345                 350

Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser
        355                 360                 365

Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr
    370                 375                 380

Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr
385                 390                 395                 400

Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe
                405                 410                 415

Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys
            420                 425                 430

Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 65
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 65

Gln Ala Ser Gln Ser Val Tyr Asn Tyr Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 66
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 66

Ser Thr Ser Thr Leu Ala Ser
1               5
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<210> SEQ ID NO 67
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 67

Leu Gly Ser Tyr Asp Cys Ser Thr Gly Asp Cys Phe Val
1 5 10

<210> SEQ ID NO 68
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 68

Asn His Tyr Met Gln
1 5

<210> SEQ ID NO 69
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 69

Val Val Gly Ile Asn Gly Arg Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 70
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 70

Gly Asp Ile
1

<210> SEQ ID NO 71
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 71

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asn Tyr Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Thr Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg

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<210> SEQ ID NO 72
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 72

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Ser Val Tyr Asn Tyr Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Thr Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
115 120 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
130 135 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145 150 155 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
165 170 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
180 185 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
195 200 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215

<210> SEQ ID NO 73
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 73

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Leu Ser Asn His
20 25 30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Val Gly Ile Asn Gly Arg Thr Tyr Tyr Ala Ser Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala

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                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
            100                 105                 110


<210> SEQ ID NO 74
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 74

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Leu Ser Asn His
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Gly Val Val Gly Ile Asn Gly Arg Thr Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
            100                 105                 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
        115                 120                 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
    130                 135                 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145                 150                 155                 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
                165                 170                 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
            180                 185                 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
        195                 200                 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335
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Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 75
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 75

Gln Ala Ser Gln Ser Val Tyr Asn Tyr Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 76
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 76

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 77
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 77

Leu Gly Ser Tyr Asp Cys Ser Thr Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 78
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 78

Asn His Tyr Met Gln
1               5


<210> SEQ ID NO 79
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 79

Val Val Gly Ile Asn Gly Arg Thr Tyr Tyr Ala Ser Trp Ala Lys Gly
1               5                   10                  15


<210> SEQ ID NO 80
<211> LENGTH: 3
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-continued

<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 80

Gly Asp Ile
1

<210> SEQ ID NO 81
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 81

Gln Val Leu Thr Gln Thr Pro Ser Pro Val Ser Ala Ala Val Gly Ser
1 5 10 15

Thr Val Thr Ile Asn Cys Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Ser Ser Arg Phe Arg
50 55 60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Asp Val Gln
65 70 75 80

Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Arg Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
100 105 110

Arg

<210> SEQ ID NO 82
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 82

Gln Val Leu Thr Gln Thr Pro Ser Pro Val Ser Ala Ala Val Gly Ser
1 5 10 15

Thr Val Thr Ile Asn Cys Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Ser Ser Arg Phe Arg
50 55 60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Asp Val Gln
65 70 75 80

Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Arg Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
100 105 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
115 120 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
130 135 140

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Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215


<210> SEQ ID NO 83
<211> LENGTH: 109
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 83

Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr Pro
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Val Ser Gly Ile Gly Leu Ser Ser Tyr Tyr
            20                  25                  30

Met Gln Trp Val Arg Gln Ser Pro Gly Arg Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Lys Thr Ser Ser Thr Thr Val Asp Leu Arg Met
65                  70                  75                  80

Ala Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Thr Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser
            100                 105


<210> SEQ ID NO 84
<211> LENGTH: 439
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 84

Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Thr Pro
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Val Ser Gly Ile Gly Leu Ser Ser Tyr Tyr
            20                  25                  30

Met Gln Trp Val Arg Gln Ser Pro Gly Arg Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Lys Thr Ser Ser Thr Thr Val Asp Leu Arg Met
65                  70                  75                  80

Ala Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Thr Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr
            100                 105                 110
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Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser
        115                 120                 125

Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu
    130                 135                 140

Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His
145                 150                 155                 160

Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser
                165                 170                 175

Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys
            180                 185                 190

Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu
        195                 200                 205

Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro
    210                 215                 220

Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys
225                 230                 235                 240

Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val
                245                 250                 255

Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp
            260                 265                 270

Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr
        275                 280                 285

Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp
    290                 295                 300

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu
305                 310                 315                 320

Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg
                325                 330                 335

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys
            340                 345                 350

Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp
        355                 360                 365

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys
    370                 375                 380

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser
385                 390                 395                 400

Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser
                405                 410                 415

Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser
            420                 425                 430

Leu Ser Leu Ser Pro Gly Lys
        435


<210> SEQ ID NO 85
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 85

Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 86
<211> LENGTH: 7
<212> TYPE: PRT
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<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 86

Ser Thr Ser Thr Leu Ala Ser
1 5

<210> SEQ ID NO 87
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 87

Leu Gly Ser Tyr Asp Cys Ser Arg Gly Asp Cys Phe Val
1 5 10

<210> SEQ ID NO 88
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 88

Ser Tyr Tyr Met Gln
1 5

<210> SEQ ID NO 89
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 89

Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 90
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 90

Gly Asp Ile
1

<210> SEQ ID NO 91
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 91

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Arg Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg

<210> SEQ ID NO 92
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 92

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1 5 10 15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn
20 25 30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
35 40 45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65 70 75 80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
85 90 95

Arg Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
100 105 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
115 120 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
130 135 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145 150 155 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
165 170 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
180 185 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
195 200 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215

<210> SEQ ID NO 93
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 93

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Gly Leu Ser Ser Tyr
20 25 30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

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Gly Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Thr
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
            100                 105                 110


<210> SEQ ID NO 94
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 94

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Gly Leu Ser Ser Tyr
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Gly Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Thr
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
            100                 105                 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
        115                 120                 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
    130                 135                 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145                 150                 155                 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
                165                 170                 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
            180                 185                 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
        195                 200                 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300
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Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 95
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 95

Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 96
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 96

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 97
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 97

Leu Gly Ser Tyr Asp Cys Ser Arg Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 98
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 98

Ser Tyr Tyr Met Gln
1               5


<210> SEQ ID NO 99
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 99
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Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys Gly
1               5                   10                  15


<210> SEQ ID NO 100
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 100

Gly Asp Ile
1


<210> SEQ ID NO 101
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 101

Gln Val Leu Thr Gln Thr Ala Ser Pro Val Ser Pro Ala Val Gly Ser
1               5                   10                  15

Thr Val Thr Ile Asn Cys Arg Ala Ser Gln Ser Val Tyr Tyr Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Ser Ser Arg Phe Lys
    50                  55                  60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Asp Val Gln
65                  70                  75                  80

Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
            100                 105                 110

Arg


<210> SEQ ID NO 102
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 102

Gln Val Leu Thr Gln Thr Ala Ser Pro Val Ser Pro Ala Val Gly Ser
1               5                   10                  15

Thr Val Thr Ile Asn Cys Arg Ala Ser Gln Ser Val Tyr Tyr Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Ser Ser Arg Phe Lys
    50                  55                  60

Gly Ser Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Asp Val Gln
65                  70                  75                  80

Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Glu Val Val Val Lys
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            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215


<210> SEQ ID NO 103
<211> LENGTH: 109
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 103

Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Gly Ser
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Val Ser Gly Ile Asp Val Thr Asn Tyr Tyr
            20                  25                  30

Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Val Asn Gly Lys Arg Tyr Tyr Ala Ser Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Lys Thr Ser Ser Thr Thr Val Asp Leu Lys Met
65                  70                  75                  80

Thr Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser
            100                 105


<210> SEQ ID NO 104
<211> LENGTH: 439
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 104

Gln Ser Leu Glu Glu Ser Gly Gly Arg Leu Val Thr Pro Gly Gly Ser
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Val Ser Gly Ile Asp Val Thr Asn Tyr Tyr
            20                  25                  30

Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Val Ile Gly Val Asn Gly Lys Arg Tyr Tyr Ala Ser Trp Ala Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Lys Thr Ser Ser Thr Thr Val Asp Leu Lys Met
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65                  70                  75                  80

Thr Ser Leu Thr Thr Glu Asp Thr Ala Thr Tyr Phe Cys Ala Arg Gly
                85                  90                  95

Asp Ile Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr
            100                 105                 110

Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser
        115                 120                 125

Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu
    130                 135                 140

Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His
145                 150                 155                 160

Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser
                165                 170                 175

Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys
            180                 185                 190

Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu
        195                 200                 205

Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro
    210                 215                 220

Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys
225                 230                 235                 240

Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val
                245                 250                 255

Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp
            260                 265                 270

Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr
        275                 280                 285

Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp
    290                 295                 300

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu
305                 310                 315                 320

Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg
                325                 330                 335

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys
            340                 345                 350

Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp
        355                 360                 365

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys
    370                 375                 380

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser
385                 390                 395                 400

Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser
                405                 410                 415

Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser
            420                 425                 430

Leu Ser Leu Ser Pro Gly Lys
        435


<210> SEQ ID NO 105
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 105
```

-continued

```
Arg Ala Ser Gln Ser Val Tyr Tyr Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 106
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 106

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 107
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 107

Leu Gly Ser Tyr Asp Cys Ser Asn Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 108
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 108

Asn Tyr Tyr Met Gln
1               5


<210> SEQ ID NO 109
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 109

Val Ile Gly Val Asn Gly Lys Arg Tyr Tyr Ala Ser Trp Ala Lys Gly
1               5                   10                  15


<210> SEQ ID NO 110
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 110

Gly Asp Ile
1


<210> SEQ ID NO 111
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 111

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1               5                   10                  15

Arg Val Thr Ile Asn Cys Arg Ala Ser Gln Ser Val Tyr Tyr Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
```

-continued

```
    50                  55                  60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg
```

<210> SEQ ID NO 112
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 112

```
Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1               5                   10                  15

Arg Val Thr Ile Asn Cys Arg Ala Ser Gln Ser Val Tyr Tyr Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
    50                  55                  60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Asn Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215
```

<210> SEQ ID NO 113
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 113

```
Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15
```

-continued

```
Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Val Thr Asn Tyr
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Gly Val Ile Gly Val Asn Gly Lys Arg Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
            100                 105                 110


<210> SEQ ID NO 114
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 114

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Asp Val Thr Asn Tyr
            20                  25                  30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Gly Val Ile Gly Val Asn Gly Lys Arg Tyr Tyr Ala Ser Trp Ala Lys
    50                  55                  60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65                  70                  75                  80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Ala
                85                  90                  95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
            100                 105                 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
        115                 120                 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
    130                 135                 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145                 150                 155                 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
                165                 170                 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
            180                 185                 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
        195                 200                 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
```

-continued

```
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 115
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 115

Arg Ala Ser Gln Ser Val Tyr Tyr Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 116
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 116

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 117
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 117

Leu Gly Ser Tyr Asp Cys Ser Asn Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 118
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 118

Asn Tyr Tyr Met Gln
1               5
```

<210> SEQ ID NO 119
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 119

Val Ile Gly Val Asn Gly Lys Arg Tyr Tyr Ala Ser Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 120
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 120

Gly Asp Ile
1

<210> SEQ ID NO 121
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 121

Ala Ile Val Met Thr Gln Thr Pro Ser Ser Lys Ser Val Pro Val Gly
1 5 10 15

Asp Thr Val Thr Ile Asn Cys Gln Ala Ser Glu Ser Leu Tyr Asn Asn
20 25 30

Asn Ala Leu Ala Trp Phe Gln Gln Lys Pro Gly Gln Pro Pro Lys Arg
35 40 45

Leu Ile Tyr Asp Ala Ser Lys Leu Ala Ser Gly Val Pro Ser Arg Phe
50 55 60

Ser Gly Gly Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Gly Val
65 70 75 80

Gln Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Gly Gly Tyr Arg Ser Asp
85 90 95

Ser Val Asp Gly Val Ala Phe Ala Gly Gly Thr Glu Val Val Val Lys
100 105 110

Arg

<210> SEQ ID NO 122
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 122

Ala Ile Val Met Thr Gln Thr Pro Ser Ser Lys Ser Val Pro Val Gly
1 5 10 15

Asp Thr Val Thr Ile Asn Cys Gln Ala Ser Glu Ser Leu Tyr Asn Asn
20 25 30

Asn Ala Leu Ala Trp Phe Gln Gln Lys Pro Gly Gln Pro Pro Lys Arg
35 40 45

Leu Ile Tyr Asp Ala Ser Lys Leu Ala Ser Gly Val Pro Ser Arg Phe
50 55 60

-continued

```
Ser Gly Gly Gly Ser Gly Thr Gln Phe Thr Leu Thr Ile Ser Gly Val
65                  70                  75                  80

Gln Cys Asp Asp Ala Ala Thr Tyr Tyr Cys Gly Gly Tyr Arg Ser Asp
                85                  90                  95

Ser Val Asp Gly Val Ala Phe Ala Gly Gly Thr Glu Val Val Val Lys
            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215


<210> SEQ ID NO 123
<211> LENGTH: 111
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 123

Gln Ser Val Glu Glu Ser Gly Gly Gly Leu Val Gln Pro Glu Gly Ser
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Ala Ser Gly Phe Asp Phe Ser Ser Asn Ala
            20                  25                  30

Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Cys Ile Tyr Asn Gly Asp Gly Ser Thr Tyr Tyr Ala Ser Trp Val Asn
    50                  55                  60

Gly Arg Phe Ser Ile Ser Lys Thr Ser Ser Thr Thr Val Thr Leu Gln
65                  70                  75                  80

Leu Asn Ser Leu Thr Val Ala Asp Thr Ala Thr Tyr Tyr Cys Ala Arg
                85                  90                  95

Asp Leu Asp Leu Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser
            100                 105                 110


<210> SEQ ID NO 124
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      polypeptide

<400> SEQUENCE: 124

Gln Ser Val Glu Glu Ser Gly Gly Gly Leu Val Gln Pro Glu Gly Ser
1               5                   10                  15

Leu Thr Leu Thr Cys Thr Ala Ser Gly Phe Asp Phe Ser Ser Asn Ala
            20                  25                  30
```

-continued

```
Met Trp Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Ile Gly
        35                  40                  45

Cys Ile Tyr Asn Gly Asp Gly Ser Thr Tyr Tyr Ala Ser Trp Val Asn
    50                  55                  60

Gly Arg Phe Ser Ile Ser Lys Thr Ser Ser Thr Thr Val Thr Leu Gln
65                  70                  75                  80

Leu Asn Ser Leu Thr Val Ala Asp Thr Ala Thr Tyr Tyr Cys Ala Arg
                85                  90                  95

Asp Leu Asp Leu Trp Gly Pro Gly Thr Leu Val Thr Val Ser Ser Ala
            100                 105                 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
        115                 120                 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
    130                 135                 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145                 150                 155                 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
                165                 170                 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
            180                 185                 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg
        195                 200                 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440
```

-continued

<210> SEQ ID NO 125
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 125

Gln Ala Ser Glu Ser Leu Tyr Asn Asn Asn Ala Leu Ala
1 5 10

<210> SEQ ID NO 126
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 126

Asp Ala Ser Lys Leu Ala Ser
1 5

<210> SEQ ID NO 127
<211> LENGTH: 12
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 127

Gly Gly Tyr Arg Ser Asp Ser Val Asp Gly Val Ala
1 5 10

<210> SEQ ID NO 128
<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 128

Ser Asn Ala Met Trp
1 5

<210> SEQ ID NO 129
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 129

Cys Ile Tyr Asn Gly Asp Gly Ser Thr Tyr Tyr Ala Ser Trp Val Asn
1 5 10 15

Gly

<210> SEQ ID NO 130
<211> LENGTH: 4
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 130

Asp Leu Asp Leu
1

<210> SEQ ID NO 131
<211> LENGTH: 113
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
polypeptide

<400> SEQUENCE: 131

Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp

-continued

```
1               5                   10                  15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
    50                  55                  60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Arg Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg
```

<210> SEQ ID NO 132
<211> LENGTH: 219
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 132

```
Gln Val Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly Asp
1               5                   10                  15

Arg Val Thr Ile Asn Cys Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn
            20                  25                  30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Val Pro Lys Gln Leu
        35                  40                  45

Ile Tyr Ser Thr Ser Thr Leu Ala Ser Gly Val Pro Ser Arg Phe Ser
    50                  55                  60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln
65                  70                  75                  80

Pro Glu Asp Val Ala Thr Tyr Tyr Cys Leu Gly Ser Tyr Asp Cys Ser
                85                  90                  95

Arg Gly Asp Cys Phe Val Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100                 105                 110

Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
        115                 120                 125

Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
    130                 135                 140

Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
145                 150                 155                 160

Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
                165                 170                 175

Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
            180                 185                 190

Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
        195                 200                 205

Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
    210                 215
```

<210> SEQ ID NO 133
<211> LENGTH: 111
<212> TYPE: PRT

-continued

<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 133

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Gly Leu Ser Ser Tyr
20 25 30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Thr
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
100 105 110

<210> SEQ ID NO 134
<211> LENGTH: 441
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polypeptide

<400> SEQUENCE: 134

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Val Ser Gly Ile Gly Leu Ser Ser Tyr
20 25 30

Tyr Met Gln Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Gly Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys
50 55 60

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Thr Thr Val Tyr Leu
65 70 75 80

Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys Thr
85 90 95

Arg Gly Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala
100 105 110

Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser
115 120 125

Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe
130 135 140

Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly
145 150 155 160

Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu
165 170 175

Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr
180 185 190

Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Ala Arg
195 200 205

Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro

```
    210                 215                 220

Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
225                 230                 235                 240

Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
                245                 250                 255

Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
            260                 265                 270

Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
        275                 280                 285

Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
    290                 295                 300

Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
305                 310                 315                 320

Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln
                325                 330                 335

Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met
            340                 345                 350

Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro
        355                 360                 365

Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn
    370                 375                 380

Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu
385                 390                 395                 400

Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val
                405                 410                 415

Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln
            420                 425                 430

Lys Ser Leu Ser Leu Ser Pro Gly Lys
        435                 440


<210> SEQ ID NO 135
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 135

Gln Ala Ser Gln Asn Val Tyr Asn Asn Asn Tyr Leu Ala
1               5                   10


<210> SEQ ID NO 136
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 136

Ser Thr Ser Thr Leu Ala Ser
1               5


<210> SEQ ID NO 137
<211> LENGTH: 13
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 137

Leu Gly Ser Tyr Asp Cys Ser Arg Gly Asp Cys Phe Val
1               5                   10


<210> SEQ ID NO 138
```

-continued

<211> LENGTH: 5
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 138

Ser Tyr Tyr Met Gln
1 5

<210> SEQ ID NO 139
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 139

Val Ile Gly Ser Asp Gly Lys Thr Tyr Tyr Ala Thr Trp Ala Lys Gly
1 5 10 15

<210> SEQ ID NO 140
<211> LENGTH: 3
<212> TYPE: PRT
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 140

Gly Asp Ile
1

<210> SEQ ID NO 141
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 141

caagtgctga cccagactgc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagag tgtttatgat aacaactacc tagcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacatcca ctctggcatc tggggtctca 180

tcgcggttca aaggcagtgg atctgggaca cagttcactc tcaccatcag cgacctggag 240

tgtgccgatg ctgccactta ctactgtcta ggcagttatg attgtagtag tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgt 339

<210> SEQ ID NO 142
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 142

caagtgctga cccagactgc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagag tgtttatgat aacaactacc tagcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacatcca ctctggcatc tggggtctca 180

tcgcggttca aaggcagtgg atctgggaca cagttcactc tcaccatcag cgacctggag 240

tgtgccgatg ctgccactta ctactgtcta ggcagttatg attgtagtag tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

-continued

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 143
<211> LENGTH: 327
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 143

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg ggacacccct gacactcacc 60

tgcacagtct ctggactcga cctcagtagc tactacatgc aatgggtccg ccaggctcca 120

gggaaggggc tggaatggat cggagtcatt ggtattaatg ataacacata ctacgcgagc 180

tgggcgaaag gccgattcac catctccaga gcctcgtcga ccacggtgga tctgaaaatg 240

accagtctga caaccgagga cacggccacc tatttctgtg ccagagggga catctggggc 300

ccaggcaccc tcgtcaccgt ctcgagc 327

<210> SEQ ID NO 144
<211> LENGTH: 1320
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 144

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg ggacacccct gacactcacc 60

tgcacagtct ctggactcga cctcagtagc tactacatgc aatgggtccg ccaggctcca 120

gggaaggggc tggaatggat cggagtcatt ggtattaatg ataacacata ctacgcgagc 180

tgggcgaaag gccgattcac catctccaga gcctcgtcga ccacggtgga tctgaaaatg 240

accagtctga caaccgagga cacggccacc tatttctgtg ccagagggga catctggggc 300

ccaggcaccc tcgtcaccgt ctcgagcgcc tccaccaagg gcccatcggt cttccccctg 360

gcaccctcct ccaagagcac ctctgggggc acagcggccc tgggctgcct ggtcaaggac 420

tacttccccg aaccggtgac ggtgtcgtgg aactcaggcg ccctgaccag cggcgtgcac 480

accttcccgg ctgtcctaca gtcctcagga ctctactccc tcagcagcgt ggtgaccgtg 540

ccctccagca gcttgggcac ccagacctac atctgcaacg tgaatcacaa gcccagcaac 600

accaaggtgg acaagagagt tgagcccaaa tcttgtgaca aaactcacac atgcccaccg 660

tgcccagcac ctgaactcct gggggggaccg tcagtcttcc tcttcccccc aaaacccaag 720

gacaccctca tgatctcccg gacccctgag gtcacatgcg tggtggtgga cgtgagccac 780

gaagaccctg aggtcaagtt caactggtac gtggacggcg tggaggtgca taatgccaag 840

acaaagccgc gggaggagca gtacgccagc acgtaccgtg tggtcagcgt cctcaccgtc 900

ctgcaccagg actggctgaa tggcaaggag tacaagtgca aggtctccaa caaagccctc 960

ccagccccca tcgagaaaac catctccaaa gccaaagggc agccccgaga accacaggtg 1020

tacaccctgc ccccatcccg ggaggagatg accaagaacc aggtcagcct gacctgcctg 1080

gtcaaaggct tctatcccag cgacatcgcc gtggagtggg agagcaatgg gcagccggag 1140

-continued

aacaactaca agaccacgcc tcccgtgctg gactccgacg gctccttctt cctctacagc 1200

aagctcaccg tggacaagag caggtggcag caggggaacg tcttctcatg ctccgtgatg 1260

catgaggctc tgcacaacca ctacacgcag aagagcctct ccctgtctcc gggtaaatga 1320

<210> SEQ ID NO 145
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 145

caggccagtc agagtgttta tgataacaac tacctagcc 39

<210> SEQ ID NO 146
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 146

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 147
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 147

ctaggcagtt atgattgtag tagtggtgat tgttttgtt 39

<210> SEQ ID NO 148
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 148

agctactaca tgcaa 15

<210> SEQ ID NO 149
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 149

gtcattggta ttaatgataa cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 150
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 150

ggggacatc 9

<210> SEQ ID NO 151
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 151

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatgat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtcta ggcagttatg attgtagtag tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 152
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 152

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatgat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtcta ggcagttatg attgtagtag tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 153
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 153

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggggtc cctgagactc 60

tcctgtgcag tctctggact cgacctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatca atgataacac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 154
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 154

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggggtc cctgagactc 60

tcctgtgcag tctctggact cgacctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatca atgataacac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacaa gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt ccccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 155
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 155

caggccagtc agagtgttta tgataacaac tacctagcc 39

<210> SEQ ID NO 156
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 156

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 157
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 157

ctaggcagtt atgattgtag tagtggtgat tgttttgtt 39

<210> SEQ ID NO 158
<211> LENGTH: 15
<212> TYPE: DNA

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<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 158

agctactaca tgcaa 15

<210> SEQ ID NO 159
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 159

gtcattggta tcaatgataa cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 160
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 160

ggggacatc 9

<210> SEQ ID NO 161
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 161

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatgat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtcta ggcagttatg attgtagtag tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 162
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 162

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatgat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtcta ggcagttatg attgtagtag tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 163
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 163

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggact cgacctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatca atgataacac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 164
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 164

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggact cgacctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatca atgataacac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacgc gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt cccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

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gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 165
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 165

caggccagtc agagtgttta tgataacaac tacctagcc 39

<210> SEQ ID NO 166
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 166

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 167
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 167

ctaggcagtt atgattgtag tagtggtgat tgttttgtt 39

<210> SEQ ID NO 168
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 168

agctactaca tgcaa 15

<210> SEQ ID NO 169
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 169

gtcattggta tcaatgataa cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 170
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 170

ggggacatc 9

<210> SEQ ID NO 171
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 171

caagtgctga cccagactcc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagag tgtttatcat aacacctacc tggcctggta tcagcagaaa 120

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ccagggcagc ctcccaaaca actgatctat gatgcatcca ctctggcgtc tggggtccca 180

tcgcggttca gcggcagtgg atctgggaca cagttcactc tcaccatcag cggcgtgcag 240

tgtaacgatg ctgccgctta ctactgtctg ggcagttatg attgtactaa tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgt 339

<210> SEQ ID NO 172
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 172

caagtgctga cccagactcc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagag tgtttatcat aacacctacc tggcctggta tcagcagaaa 120

ccagggcagc ctcccaaaca actgatctat gatgcatcca ctctggcgtc tggggtccca 180

tcgcggttca gcggcagtgg atctgggaca cagttcactc tcaccatcag cggcgtgcag 240

tgtaacgatg ctgccgctta ctactgtctg ggcagttatg attgtactaa tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 173
<211> LENGTH: 327
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 173

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg ggacacccct gacactcacc 60

tgttccgtct ctggcatcga cctcagtggc tactacatga actgggtccg ccaggctcca 120

gggaaggggc tggaatggat cggagtcatt ggtattaatg gtgccacata ctacgcgagc 180

tgggcgaaag gccgattcac catctccaaa acctcgtcga ccacggtgga tctgaaaatg 240

accagtctga caaccgagga cacggccacc tatttctgtg ccagagggga catctggggc 300

ccgggcaccc tcgtcaccgt ctcgagc 327

<210> SEQ ID NO 174
<211> LENGTH: 1320
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 174

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg ggacacccct gacactcacc 60

tgttccgtct ctggcatcga cctcagtggc tactacatga actgggtccg ccaggctcca 120

-continued

gggaaggggc tggaatggat cggagtcatt ggtattaatg gtgccacata ctacgcgagc 180

tgggcgaaag gccgattcac catctccaaa acctcgtcga ccacggtgga tctgaaaatg 240

accagtctga caaccgagga cacggccacc tatttctgtg ccagagggga catctggggc 300

ccgggcaccc tcgtcaccgt ctcgagcgcc tccaccaagg gcccatcggt cttccccctg 360

gcaccctcct ccaagagcac ctctgggggc acagcggccc tgggctgcct ggtcaaggac 420

tacttccccg aaccggtgac ggtgtcgtgg aactcaggcg ccctgaccag cggcgtgcac 480

accttcccgg ctgtcctaca gtcctcagga ctctactccc tcagcagcgt ggtgaccgtg 540

ccctccagca gcttgggcac ccagacctac atctgcaacg tgaatcacaa gcccagcaac 600

accaaggtgg acaagagagt tgagcccaaa tcttgtgaca aaactcacac atgcccaccg 660

tgcccagcac ctgaactcct gggggggaccg tcagtcttcc tcttcccccc aaaacccaag 720

gacaccctca tgatctcccg gacccctgag gtcacatgcg tggtggtgga cgtgagccac 780

gaagaccctg aggtcaagtt caactggtac gtggacggcg tggaggtgca taatgccaag 840

acaaagccgc gggaggagca gtacgccagc acgtaccgtg tggtcagcgt cctcaccgtc 900

ctgcaccagg actggctgaa tggcaaggag tacaagtgca aggtctccaa caaagccctc 960

ccagccccca tcgagaaaac catctccaaa gccaaagggc agccccgaga accacaggtg 1020

tacaccctgc ccccatcccg ggaggagatg accaagaacc aggtcagcct gacctgcctg 1080

gtcaaaggct tctatcccag cgacatcgcc gtggagtggg agagcaatgg gcagccggag 1140

aacaactaca agaccacgcc tcccgtgctg gactccgacg gctccttctt cctctacagc 1200

aagctcaccg tggacaagag caggtggcag caggggaacg tcttctcatg ctccgtgatg 1260

catgaggctc tgcacaacca ctacacgcag aagagcctct ccctgtctcc gggtaaatga 1320

<210> SEQ ID NO 175
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 175

caggccagtc agagtgttta tcataacacc tacctggcc 39

<210> SEQ ID NO 176
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 176

gatgcatcca ctctggcgtc t 21

<210> SEQ ID NO 177
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 177

ctgggcagtt atgattgtac taatggtgat tgttttgtt 39

<210> SEQ ID NO 178
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 178

ggctactaca tgaac 15

<210> SEQ ID NO 179
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 179

gtcattggta ttaatggtgc cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 180
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 180

ggggacatc 9

<210> SEQ ID NO 181
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 181

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatcat aacacctacc tggcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat gatgcatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtactaa tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 182
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 182

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatcat aacacctacc tggcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat gatgcatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtactaa tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 183

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<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 183

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacctcagt ggctactaca tgaactgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatta atggtgccac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 184
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 184

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacctcagt ggctactaca tgaactgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatta atggtgccac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacaa gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt cccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

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<210> SEQ ID NO 185
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 185

caggccagtc agagtgttta tcataacacc tacctggcc 39

<210> SEQ ID NO 186
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 186

gatgcatcca ctctggcatc t 21

<210> SEQ ID NO 187
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 187

ctgggcagtt atgattgtac taatggtgat tgttttgtt 39

<210> SEQ ID NO 188
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 188

ggctactaca tgaac 15

<210> SEQ ID NO 189
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 189

gtcattggta ttaatggtgc cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 190
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 190

ggggacatc 9

<210> SEQ ID NO 191
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 191

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatcat aacacctacc tggcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat gatgcatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtactaa tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 192
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 192

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttatcat aacacctacc tggcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat gatgcatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtactaa tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 193
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 193

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacctcagt ggctactaca tgaactgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatta atggtgccac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 194
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 194

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacctcagt ggctactaca tgaactgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtatta atggtgccac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacgc gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt cccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 195
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 195

caggccagtc agagtgttta tcataacacc tacctggcc 39

<210> SEQ ID NO 196
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 196

gatgcatcca ctctggcatc t 21

<210> SEQ ID NO 197
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 197

ctgggcagtt atgattgtac taatggtgat tgttttgtt 39

<210> SEQ ID NO 198
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 198

ggctactaca tgaac 15

<210> SEQ ID NO 199

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<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 199

gtcattggta ttaatggtgc cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 200
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 200

ggggacatc 9

<210> SEQ ID NO 201
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 201

caagtgctga cccagactgc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagag tgtttataat tacaactacc ttgcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacatcca ctctggcatc tggggtctca 180

tcgcgattca aaggcagtgg atctgggaca cagttcactc tcaccatcag cgacgtgcag 240

tgtgacgatg ctgccactta ctactgtcta ggcagttatg actgtagtac tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgt 339

<210> SEQ ID NO 202
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 202

caagtgctga cccagactgc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagag tgtttataat tacaactacc ttgcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacatcca ctctggcatc tggggtctca 180

tcgcgattca aaggcagtgg atctgggaca cagttcactc tcaccatcag cgacgtgcag 240

tgtgacgatg ctgccactta ctactgtcta ggcagttatg actgtagtac tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 203
<211> LENGTH: 330
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence

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<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 203

caggagcagc tgaaggagtc cgggggtcgc ctggtcacgc ctgggacatc cctgacactc 60

acctgcaccg tctctggaat cgacctcagt aaccactaca tgcaatgggt ccgccaggct 120

ccagggaagg ggctggagtg gatcggagtc gttggtatta atggtcgcac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agaacctcgt cgaccacggt ggatctgaaa 240

atgaccaggc tgacaaccga ggacacggcc acctatttct gtgccagagg ggacatctgg 300

ggcccaggca ccctggtcac cgtctcgagc 330

<210> SEQ ID NO 204
<211> LENGTH: 1323
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 204

caggagcagc tgaaggagtc cgggggtcgc ctggtcacgc ctgggacatc cctgacactc 60

acctgcaccg tctctggaat cgacctcagt aaccactaca tgcaatgggt ccgccaggct 120

ccagggaagg ggctggagtg gatcggagtc gttggtatta atggtcgcac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agaacctcgt cgaccacggt ggatctgaaa 240

atgaccaggc tgacaaccga ggacacggcc acctatttct gtgccagagg ggacatctgg 300

ggcccaggca ccctggtcac cgtctcgagc gcctccacca agggcccatc ggtcttcccc 360

ctggcaccct cctccaagag cacctctggg ggcacagcgg ccctgggctg cctggtcaag 420

gactacttcc ccgaaccggt gacggtgtcg tggaactcag gcgccctgac cagcggcgtg 480

cacaccttcc cggctgtcct acagtcctca ggactctact ccctcagcag cgtggtgacc 540

gtgccctcca gcagcttggg cacccagacc tacatctgca acgtgaatca caagcccagc 600

aacaccaagg tggacaagag agttgagccc aaatcttgtg acaaaactca cacatgccca 660

ccgtgcccag cacctgaact cctgggggga ccgtcagtct tcctcttccc cccaaaaccc 720

aaggacaccc tcatgatctc ccggacccct gaggtcacat gcgtggtggt ggacgtgagc 780

cacgaagacc ctgaggtcaa gttcaactgg tacgtggacg gcgtggaggt gcataatgcc 840

aagacaaagc cgcgggagga gcagtacgcc agcacgtacc gtgtggtcag cgtcctcacc 900

gtcctgcacc aggactggct gaatggcaag gagtacaagt gcaaggtctc caacaaagcc 960

ctcccagccc ccatcgagaa aaccatctcc aaagccaaag ggcagccccg agaaccacag 1020

gtgtacaccc tgcccccatc ccgggaggag atgaccaaga accaggtcag cctgacctgc 1080

ctggtcaaag gcttctatcc cagcgacatc gccgtggagt gggagagcaa tgggcagccg 1140

gagaacaact acaagaccac gcctcccgtg ctggactccg acggctcctt cttcctctac 1200

agcaagctca ccgtggacaa gagcaggtgg cagcagggga acgtcttctc atgctccgtg 1260

atgcatgagg ctctgcacaa ccactacacg cagaagagcc tctccctgtc tccgggtaaa 1320

tga 1323

<210> SEQ ID NO 205
<211> LENGTH: 39
<212> TYPE: DNA

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<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 205

caggccagtc agagtgttta taattacaac taccttgcc 39

<210> SEQ ID NO 206
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 206

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 207
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 207

ctaggcagtt atgactgtag tactggtgat tgttttgtt 39

<210> SEQ ID NO 208
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 208

aaccactaca tgcaa 15

<210> SEQ ID NO 209
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 209

gtcgttggta ttaatggtcg cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 210
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 210

ggggacatc 9

<210> SEQ ID NO 211
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 211

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttacaat tacaactacc ttgcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtac tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 212
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 212

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagag tgtttacaat tacaactacc ttgcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtac tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 213
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 213

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacctcagt aaccactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc gttggtatca atggtcgcac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 214
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 214

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacctcagt aaccactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc gttggtatca atggtcgcac atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgctag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

-continued

ccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacaa gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt cccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 215
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 215

caggccagtc agagtgttta caattacaac taccttgcc 39

<210> SEQ ID NO 216
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 216

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 217
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 217

ctgggcagtt atgattgtag tactggtgat tgttttgtt 39

<210> SEQ ID NO 218
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 218

aaccactaca tgcaa 15

<210> SEQ ID NO 219
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

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<400> SEQUENCE: 219

gtcgttggta tcaatggtcg cacatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 220
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 220

ggggacatc 9

<210> SEQ ID NO 221
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 221

caagtgctga cccagactcc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagaa tgtttataat aacaactacc tagcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacgtcca ctctggcatc tggggtctca 180

tcgcgattca gaggcagtgg atctgggaca cagttcactc tcaccatcag cgacgtgcag 240

tgtgacgatg ctgccactta ctactgtcta ggcagttatg attgtagtcg tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgt 339

<210> SEQ ID NO 222
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 222

caagtgctga cccagactcc atcccccgtg tctgcagctg tgggaagcac agtcaccatc 60

aattgccagg ccagtcagaa tgtttataat aacaactacc tagcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacgtcca ctctggcatc tggggtctca 180

tcgcgattca gaggcagtgg atctgggaca cagttcactc tcaccatcag cgacgtgcag 240

tgtgacgatg ctgccactta ctactgtcta ggcagttatg attgtagtcg tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 223
<211> LENGTH: 327
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

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<400> SEQUENCE: 223

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg ggacacccct gacactcacc 60

tgcacagtct ctggaatcgg cctcagtagc tactacatgc agtgggtccg ccagtctcca 120

gggaggggggc tggaatggat cggagtcatt ggtagtgatg gtaagacata ctacgcgacc 180

tgggcgaaag gccgattcac catctccaag acctcgtcga ccacggtgga tctgagaatg 240

gccagtctga caaccgagga cacggccacc tatttctgta ccagagggga catctggggc 300

ccggggaccc tcgtcaccgt ctcgagc 327

<210> SEQ ID NO 224
<211> LENGTH: 1320
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 224

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg ggacacccct gacactcacc 60

tgcacagtct ctggaatcgg cctcagtagc tactacatgc agtgggtccg ccagtctcca 120

gggaggggggc tggaatggat cggagtcatt ggtagtgatg gtaagacata ctacgcgacc 180

tgggcgaaag gccgattcac catctccaag acctcgtcga ccacggtgga tctgagaatg 240

gccagtctga caaccgagga cacggccacc tatttctgta ccagagggga catctggggc 300

ccggggaccc tcgtcaccgt ctcgagcgcc tccaccaagg gcccatcggt cttccccctg 360

gcaccctcct ccaagagcac ctctgggggc acagcggccc tgggctgcct ggtcaaggac 420

tacttccccg aaccggtgac ggtgtcgtgg aactcaggcg ccctgaccag cggcgtgcac 480

accttcccgg ctgtcctaca gtcctcagga ctctactccc tcagcagcgt ggtgaccgtg 540

ccctccagca gcttgggcac ccagacctac atctgcaacg tgaatcacaa gcccagcaac 600

accaaggtgg acaagagagt tgagcccaaa tcttgtgaca aaactcacac atgcccaccg 660

tgcccagcac ctgaactcct gggggggaccg tcagtcttcc tcttcccccc aaaacccaag 720

gacaccctca tgatctcccg gacccctgag gtcacatgcg tggtggtgga cgtgagccac 780

gaagaccctg aggtcaagtt caactggtac gtggacggcg tggaggtgca taatgccaag 840

acaaagccgc gggaggagca gtacgccagc acgtaccgtg tggtcagcgt cctcaccgtc 900

ctgcaccagg actggctgaa tggcaaggag tacaagtgca aggtctccaa caaagccctc 960

ccagccccca tcgagaaaac catctccaaa gccaaagggc agccccgaga accacaggtg 1020

tacaccctgc ccccatcccg ggaggagatg accaagaacc aggtcagcct gacctgcctg 1080

gtcaaaggct tctatcccag cgacatcgcc gtggagtggg agagcaatgg gcagccggag 1140

aacaactaca agaccacgcc tcccgtgctg gactccgacg gctccttctt cctctacagc 1200

aagctcaccg tggacaagag caggtggcag caggggaacg tcttctcatg ctccgtgatg 1260

catgaggctc tgcacaacca ctacacgcag aagagcctct ccctgtctcc gggtaaatga 1320

<210> SEQ ID NO 225
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 225

caggccagtc agaatgttta taataacaac tacctagcc 39

<210> SEQ ID NO 226
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 226

tctacgtcca ctctggcatc t 21

<210> SEQ ID NO 227
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 227

ctaggcagtt atgattgtag tcgtggtgat tgttttgtt 39

<210> SEQ ID NO 228
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 228

agctactaca tgcag 15

<210> SEQ ID NO 229
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 229

gtcattggta gtgatggtaa gacatactac gcgacctggg cgaaaggc 48

<210> SEQ ID NO 230
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 230

ggggacatc 9

<210> SEQ ID NO 231
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 231

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagaa tgtttacaat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtcg tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 232
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence

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<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 232

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagaa tgtttacaat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtcg tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 233
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 233

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cggcctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtagtg atggtaagac atactacgcg 180

acctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtaccag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 234
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 234

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cggcctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtagtg atggtaagac atactacgcg 180

acctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtaccag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacaa gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt cccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 235
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 235

caggccagtc agaatgttta caataacaac tacctagcc 39

<210> SEQ ID NO 236
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 236

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 237
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 237

ctgggcagtt atgattgtag tcgtggtgat tgttttgtt 39

<210> SEQ ID NO 238
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 238

agctactaca tgcaa 15

<210> SEQ ID NO 239
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 239

gtcattggta gtgatggtaa gacatactac gcgacctggg cgaaaggc 48

<210> SEQ ID NO 240
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 240

ggggacatc 9

<210> SEQ ID NO 241
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 241

caggtgctga cccagactgc atcccccgtg tctccagctg tgggaagcac agtcaccatc 60

aattgccggg ccagtcagag tgtttattat aacaactacc tagcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacatcca ctctggcatc tggggtctca 180

tcgcggttca aaggcagtgg atctgggaca cagttcactc tcaccatcag cgacgtgcag 240

tgtgacgatg ctgccactta ctactgtcta ggcagttatg attgtagtaa tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgt 339

<210> SEQ ID NO 242
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 242

caggtgctga cccagactgc atcccccgtg tctccagctg tgggaagcac agtcaccatc 60

aattgccggg ccagtcagag tgtttattat aacaactacc tagcctggta tcagcagaaa 120

ccagggcagc ctcccaagca actgatctat tctacatcca ctctggcatc tggggtctca 180

tcgcggttca aaggcagtgg atctgggaca cagttcactc tcaccatcag cgacgtgcag 240

tgtgacgatg ctgccactta ctactgtcta ggcagttatg attgtagtaa tggtgattgt 300

tttgttttcg gcggagggac cgaggtggtg gtcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 243
<211> LENGTH: 327
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 243

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg gaggatccct gacactcacc 60

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tgcacagtct ctggaatcga cgtcactaac tactatatgc aatgggtccg ccaggctcca 120

gggaaggggc tggaatggat cggagtcatt ggtgtgaatg gtaagagata ctacgcgagc 180

tgggcgaaag gccgattcac catctccaaa acctcgtcga ccacggtgga tctgaaaatg 240

accagtctga caaccgagga cacggccacc tatttctgtg ccagaggcga catctggggc 300

ccggggaccc tcgtcaccgt ctcgagc 327

<210> SEQ ID NO 244
<211> LENGTH: 1320
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 244

cagtcgctgg aggagtccgg gggtcgcctg gtcacgcctg gaggatccct gacactcacc 60

tgcacagtct ctggaatcga cgtcactaac tactatatgc aatgggtccg ccaggctcca 120

gggaaggggc tggaatggat cggagtcatt ggtgtgaatg gtaagagata ctacgcgagc 180

tgggcgaaag gccgattcac catctccaaa acctcgtcga ccacggtgga tctgaaaatg 240

accagtctga caaccgagga cacggccacc tatttctgtg ccagaggcga catctggggc 300

ccggggaccc tcgtcaccgt ctcgagcgcc tccaccaagg gcccatcggt cttccccctg 360

gcaccctcct ccaagagcac ctctgggggc acagcggccc tgggctgcct ggtcaaggac 420

tacttccccg aaccggtgac ggtgtcgtgg aactcaggcg ccctgaccag cggcgtgcac 480

accttccccgg ctgtcctaca gtcctcagga ctctactccc tcagcagcgt ggtgaccgtg 540

ccctccagca gcttgggcac ccagacctac atctgcaacg tgaatcacaa gcccagcaac 600

accaaggtgg acaagagagt tgagcccaaa tcttgtgaca aaactcacac atgcccaccg 660

tgcccagcac ctgaactcct gggggggaccg tcagtcttcc tcttcccccc aaaacccaag 720

gacaccctca tgatctcccg gacccctgag gtcacatgcg tggtggtgga cgtgagccac 780

gaagaccctg aggtcaagtt caactggtac gtggacggcg tggaggtgca taatgccaag 840

acaaagccgc gggaggagca gtacgccagc acgtaccgtg tggtcagcgt cctcaccgtc 900

ctgcaccagg actggctgaa tggcaaggag tacaagtgca aggtctccaa caaagccctc 960

ccagccccca tcgagaaaac catctccaaa gccaaagggc agccccgaga accacaggtg 1020

tacaccctgc ccccatcccg ggaggagatg accaagaacc aggtcagcct gacctgcctg 1080

gtcaaaggct tctatcccag cgacatcgcc gtggagtggg agagcaatgg gcagccggag 1140

aacaactaca agaccacgcc tcccgtgctg gactccgacg gctccttctt cctctacagc 1200

aagctcaccg tggacaagag caggtggcag caggggaacg tcttctcatg ctccgtgatg 1260

catgaggctc tgcacaacca ctacacgcag aagagcctct ccctgtctcc gggtaaatga 1320

<210> SEQ ID NO 245
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 245

cgggccagtc agagtgttta ttataacaac tacctagcc 39

<210> SEQ ID NO 246
<211> LENGTH: 21
<212> TYPE: DNA

<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 246

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 247
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 247

ctaggcagtt atgattgtag taatggtgat tgttttgtt 39

<210> SEQ ID NO 248
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 248

aactactata tgcaa 15

<210> SEQ ID NO 249
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 249

gtcattggtg tgaatggtaa gagatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 250
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 250

ggcgacatc 9

<210> SEQ ID NO 251
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 251

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccggg ccagtcagag tgtttactat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtaa tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 252
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 252

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caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccggg ccagtcagag tgtttactat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtaa tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 253
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 253

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacgtcact aactactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtgtga atggtaagag atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgccag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 254
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 254

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggggtc cctgagactc 60

tcctgtgcag tctctggaat cgacgtcact aactactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtgtga atggtaagag atactacgcg 180

agctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtgccag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacaa gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

-continued

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt ccccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 255
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 255

cgggccagtc agagtgttta ctataacaac tacctagcc 39

<210> SEQ ID NO 256
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 256

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 257
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 257

ctgggcagtt atgattgtag taatggtgat tgttttgtt 39

<210> SEQ ID NO 258
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 258

aactactaca tgcaa 15

<210> SEQ ID NO 259
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 259

gtcattggtg tgaatggtaa gagatactac gcgagctggg cgaaaggc 48

<210> SEQ ID NO 260
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 260

ggggacatc 9

<210> SEQ ID NO 261
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 261

gccatcgtga tgacccagac tccatcttcc aagtctgtcc ctgtgggaga cacagtcacc 60

atcaattgcc aggccagtga gagtctttat aataacaacg ccttggcctg gtttcagcag 120

aaaccagggc agcctcccaa gcgcctgatc tatgatgcat ccaaactggc atctggggtc 180

ccatcgcggt tcagtggcgg tgggtctggg acacagttca ctctcaccat cagtggcgtg 240

cagtgtgacg atgctgccac ttactactgt ggaggctaca gaagtgatag tgttgatggt 300

gttgctttcg ccggagggac cgaggtggtg gtcaaacgt 339

<210> SEQ ID NO 262
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 262

gccatcgtga tgacccagac tccatcttcc aagtctgtcc ctgtgggaga cacagtcacc 60

atcaattgcc aggccagtga gagtctttat aataacaacg ccttggcctg gtttcagcag 120

aaaccagggc agcctcccaa gcgcctgatc tatgatgcat ccaaactggc atctggggtc 180

ccatcgcggt tcagtggcgg tgggtctggg acacagttca ctctcaccat cagtggcgtg 240

cagtgtgacg atgctgccac ttactactgt ggaggctaca gaagtgatag tgttgatggt 300

gttgctttcg ccggagggac cgaggtggtg gtcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 263
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 263

cagtcggtgg aggagtccgg gggaggcctg gtccagcctg agggatccct gacactcacc 60

tgcacagcct ctggattcga cttcagtagc aatgcaatgt ggtgggtccg ccaggctcca 120

gggaaggggc tggagtggat cggatgcatt tacaatggtg atggcagcac atactacgcg 180

agctgggtga atggccgatt ctccatctcc aaaacctcgt cgaccacggt gactctgcaa 240

ctgaatagtc tgacagtcgc ggacacggcc acgtattatt gtgcgagaga tcttgacttg 300

tggggcccgg gcaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 264
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 264

cagtcggtgg aggagtccgg gggaggcctg gtccagcctg agggatccct gacactcacc 60

tgcacagcct ctggattcga cttcagtagc aatgcaatgt ggtgggtccg ccaggctcca 120

gggaaggggc tggagtggat cggatgcatt tacaatggtg atggcagcac atactacgcg 180

agctgggtga atggccgatt ctccatctcc aaaacctcgt cgaccacggt gactctgcaa 240

ctgaatagtc tgacagtcgc ggacacggcc acgtattatt gtgcgagaga tcttgacttg 300

tggggcccgg gcaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacaa gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt ccccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 265
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 265

caggccagtg agagtcttta taataacaac gccttggcc 39

<210> SEQ ID NO 266
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 266

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gatgcatcca aactggcatc t 21

<210> SEQ ID NO 267
<211> LENGTH: 36
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 267

ggaggctaca gaagtgatag tgttgatggt gttgct 36

<210> SEQ ID NO 268
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 268

agcaatgcaa tgtgg 15

<210> SEQ ID NO 269
<211> LENGTH: 51
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 269

tgcatttaca atggtgatgg cagcacatac tacgcgagct gggtgaatgg c 51

<210> SEQ ID NO 270
<211> LENGTH: 12
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 270

gatcttgact tg 12

<210> SEQ ID NO 271
<211> LENGTH: 339
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 271

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

aattgccagg ccagtcagaa tgtttacaat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtcg tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgt 339

<210> SEQ ID NO 272
<211> LENGTH: 660
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 272

caagtgctga cccagtctcc atcctccctg tctgcatctg taggagacag agtcaccatc 60

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aattgccagg ccagtcagaa tgtttacaat aacaactacc tagcctggta tcagcagaaa 120

ccagggaaag ttcctaagca actgatctat tctacatcca ctctggcatc tggggtccca 180

tctcgtttca gtggcagtgg atctgggaca gatttcactc tcaccatcag cagcctgcag 240

cctgaagatg ttgcaactta ttactgtctg ggcagttatg attgtagtcg tggtgattgt 300

tttgttttcg gcggaggaac caaggtggaa atcaaacgta cggtggctgc accatctgtc 360

ttcatcttcc cgccatctga tgagcagttg aaatctggaa ctgcctctgt tgtgtgcctg 420

ctgaataact tctatcccag agaggccaaa gtacagtgga aggtggataa cgccctccaa 480

tcgggtaact cccaggagag tgtcacagag caggacagca aggacagcac ctacagcctc 540

agcagcaccc tgacgctgag caaagcagac tacgagaaac acaaagtcta cgcctgcgaa 600

gtcacccatc agggcctgag ctcgcccgtc acaaagagct tcaacagggg agagtgttag 660

<210> SEQ ID NO 273
<211> LENGTH: 333
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 273

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cggcctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtagtg atggtaagac atactacgcg 180

acctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtaccag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agc 333

<210> SEQ ID NO 274
<211> LENGTH: 1326
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic polynucleotide

<400> SEQUENCE: 274

gaggtgcagc ttgtggagtc tgggggaggc ttggtccagc ctgggggtc cctgagactc 60

tcctgtgcag tctctggaat cggcctcagt agctactaca tgcaatgggt ccgtcaggct 120

ccagggaagg ggctggagtg ggtcggagtc attggtagtg atggtaagac atactacgcg 180

acctgggcga aaggccgatt caccatctcc agagacaatt ccaagaccac ggtgtatctt 240

caaatgaaca gcctgagagc tgaggacact gctgtgtatt tctgtaccag aggggacatc 300

tggggccaag ggaccctcgt caccgtctcg agcgcctcca ccaagggccc atcggtcttc 360

cccctggcac cctcctccaa gagcacctct gggggcacag cggccctggg ctgcctggtc 420

aaggactact tccccgaacc ggtgacggtg tcgtggaact caggcgccct gaccagcggc 480

gtgcacacct tcccggctgt cctacagtcc tcaggactct actccctcag cagcgtggtg 540

accgtgccct ccagcagctt gggcacccag acctacatct gcaacgtgaa tcacaagccc 600

agcaacacca aggtggacgc gagagttgag cccaaatctt gtgacaaaac tcacacatgc 660

ccaccgtgcc cagcacctga actcctgggg ggaccgtcag tcttcctctt cccccaaaa 720

cccaaggaca ccctcatgat ctcccggacc cctgaggtca catgcgtggt ggtggacgtg 780

agccacgaag accctgaggt caagttcaac tggtacgtgg acggcgtgga ggtgcataat 840

gccaagacaa agccgcggga ggagcagtac gccagcacgt accgtgtggt cagcgtcctc 900

accgtcctgc accaggactg gctgaatggc aaggagtaca agtgcaaggt ctccaacaaa 960

gccctcccag cccccatcga gaaaaccatc tccaaagcca aagggcagcc ccgagaacca 1020

caggtgtaca ccctgccccc atcccgggag gagatgacca agaaccaggt cagcctgacc 1080

tgcctggtca aaggcttcta tcccagcgac atcgccgtgg agtgggagag caatgggcag 1140

ccggagaaca actacaagac cacgcctccc gtgctggact ccgacggctc cttcttcctc 1200

tacagcaagc tcaccgtgga caagagcagg tggcagcagg ggaacgtctt ctcatgctcc 1260

gtgatgcatg aggctctgca caaccactac acgcagaaga gcctctccct gtctccgggt 1320

aaatga 1326

<210> SEQ ID NO 275
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 275

caggccagtc agaatgttta caataacaac tacctagcc 39

<210> SEQ ID NO 276
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 276

tctacatcca ctctggcatc t 21

<210> SEQ ID NO 277
<211> LENGTH: 39
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 277

ctgggcagtt atgattgtag tcgtggtgat tgttttgtt 39

<210> SEQ ID NO 278
<211> LENGTH: 15
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 278

agctactaca tgcaa 15

<210> SEQ ID NO 279
<211> LENGTH: 48
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 279

gtcattggta gtgatggtaa gacatactac gcgacctggg cgaaaggc 48

<210> SEQ ID NO 280
<211> LENGTH: 9
<212> TYPE: DNA
<213> ORGANISM: Oryctolagus cuniculus

<400> SEQUENCE: 280

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ggggacatc 9

<210> SEQ ID NO 281
<211> LENGTH: 37
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<220> FEATURE:
<223> OTHER INFORMATION: C-term amidated

<400> SEQUENCE: 281

Ala Cys Asp Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
1 5 10 15

Ser Arg Ser Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn Val
20 25 30

Gly Ser Lys Ala Phe
35

<210> SEQ ID NO 282
<211> LENGTH: 37
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<220> FEATURE:
<223> OTHER INFORMATION: C-term amidated

<400> SEQUENCE: 282

Ala Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
1 5 10 15

Ser Arg Ser Gly Gly Met Val Lys Ser Asn Phe Val Pro Thr Asn Val
20 25 30

Gly Ser Lys Ala Phe
35

<210> SEQ ID NO 283
<211> LENGTH: 105
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 283

Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu
1 5 10 15

Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro
20 25 30

Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly
35 40 45

Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr
50 55 60

Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His
65 70 75 80

Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val
85 90 95

Thr Lys Ser Phe Asn Arg Gly Glu Cys
100 105

<210> SEQ ID NO 284
<211> LENGTH: 330
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 284

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
1 5 10 15

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Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
            20                  25                  30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
        35                  40                  45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser
    50                  55                  60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
65                  70                  75                  80

Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
                85                  90                  95

Arg Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
            100                 105                 110

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
        115                 120                 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
    130                 135                 140

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
145                 150                 155                 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
                165                 170                 175

Glu Gln Tyr Ala Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
            180                 185                 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
        195                 200                 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
    210                 215                 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu
225                 230                 235                 240

Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
                245                 250                 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
            260                 265                 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
        275                 280                 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
    290                 295                 300

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr
305                 310                 315                 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
                325                 330
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What is claimed is:

1. A method of assessing the potential efficacy of a candidate anti-CGRP antibody or antibody fragment in inhibiting Calcitonin Gene Related Peptide (CGRP)-associated photophobia or light aversion or precluding the onset of CGRP-associated photophobia or photoaversion associated with a migrainous condition or cluster headache condition in a subject comprising determining whether the antibody or antibody fragment inhibits light aversive behavior in a first Nestin/hRAMP1 rodent administered CGRP and the candidate antibody or antibody fragment compared to the light aversive behavior in a second Nestin/hRAMP1 rodent administered CGRP in the absence of the candidate anti-CGRP antibody or antibody fragment, wherein said candidate anti-GRP antibody or antibody fragment is a variant of an anti-CGRP antibody comprising a variable light (VL) polypeptide comprising the VL CDRs of SEQ ID NO: 55, 56 and 57 and a variable heavy (VH) polypeptide comprising the VH CDR polypeptides of SEQ ID NO: 58, 59 and 60 wherein said variant comprises said CDR polypeptides with the exception that one of the CDR residues in one of the CDRs of said VH or in one of the CDRs of said VL polypeptide has been substituted with another amino acid residue.

2. The method of claim **1**, wherein said anti-CGRP antibody or antibody fragment is human, humanized, or chimeric.

3. The method of claim **2**, wherein said anti-CGRP antibody or antibody fragment is humanized.

4. The method of claim 3, wherein the humanized anti-CGRP antibody or antibody fragment comprises human constant regions selected from the group consisting of IgG1, IgG2, IgG3 and IgG4 constant regions.

5. The method of claim 1, wherein said anti-CGRP antibody fragment is an scFv, camelbody, nanobody, IgNAR, Fab, Fab', or $F(ab)_2$ fragment.

6. The method of claim 1, wherein said anti-CGRP antibody or antibody fragment is non-glycosylated.

7. The method of claim 1, wherein said anti-CGRP antibody or antibody fragment comprises a variable light (VL) polypeptide having a sequence possessing at least 90% sequence identity SEQ ID NO: 51 and further comprises a variable heavy (VH) polypeptide having a sequence possessing at least 90% sequence identity to SEQ ID NO: 53.

8. The method of claim 1, wherein said anti-CGRP antibody or antibody fragment binds to CGRP with an off-rate (K_{off}) of less than or equal to 10^{-4} S^{-1}, 5×10^{-5} S^{-1}, 10^{-5} S^{-1}, 5×10^{-6} S^{-1}, 10^{-6} S^{-1}, 5×10^{-7} S^{-1}, or 10^{-7} S^{1}.

9. The method of claim 1, wherein said anti-CGRP antibody has an Fc region that contains a mutation that alters or eliminates glycosylation or another Fc effector function.

10. The method of claim 1, wherein said anti-CGRP antibody or antibody fragment comprises a variable light (VL) polypeptide having a sequence possessing at least 95% sequence identity SEQ ID NO: 51 and further comprises a variable heavy (VH) polypeptide having a sequence possessing at least 95% sequence identity to SEQ ID NO: 53.

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