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(54) **ASSOCIATION OF THE MUSCARINIC CHOLINERGIC 2 RECEPTOR (CHRM2) GENE WITH MAJOR DEPRESSION IN WOMEN**

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(51) **Int. Cl.**<sup>7</sup> ..... **C12Q 1/68**; C12P 19/34;  
C07H 21/02; C07H 21/04

(52) **U.S. Cl.** ..... **435/6**; 735/91.2; 536/23.1;  
536/24.3

(58) **Field of Search** ..... 435/6, 91.2; 536/23.1,  
536/24.3, 23.2

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(57) **ABSTRACT**

The present invention relates to the observation that women having an A→T 1890 polymorphism in the 3' UTR of the cholinergic muscarinic receptor 2 (CHRM2) gene have an increased risk for developing major depression. The present invention provides diagnostic, screening and therapeutic methods based on that observation.

**13 Claims, No Drawings**

**ASSOCIATION OF THE MUSCARINIC CHOLINERGIC 2 RECEPTOR (CHRM2) GENE WITH MAJOR DEPRESSION IN WOMEN**

**CROSS-REFERENCES TO RELATED APPLICATIONS**

The present application is related to U.S. provisional application Ser. No. 60/298,108 filed Jun. 15, 2001, incorporated herein by reference.

**STATEMENT REGARDING FEDERALLY SPONSORED RESEARCH OR DEVELOPMENT**

Not applicable.

**BACKGROUND OF THE INVENTION**

1. Field of the Invention

The present invention relates to screening patients to determine their risk for having major depression.

2. Description of the Related Art

A bibliography follows at the end of the Detailed Description of the Invention. The listed references are all incorporated herein by reference.

The lifetime frequency of major depression (MD) is twice as high in women as in men (1). Twin studies have shown a significant genetic contribution to MD in women with heritabilities ranging from 0.33 to 0.45 and higher (2-5), and an important role of stress (6). Some twin studies have suggested a comparable heritability for men and women (4), while others have suggested a greater heritability for women (5). In 1972 and again in 1994, Janowsky (8, 9) reviewed the evidence for a role of cholinergic hypersensitivity in depression. Both REM and non-REM sleep is regulated by cholinergic, serotonergic and noradrenergic neurons in the brain stem (10). The early onset of REM sleep, increased REM density and exaggerated REM response to cholinergic stimulation in depression, is consistent with CNS cholinergic overactivity or muscarinic supersensitivity in depression (10). While the HPA axis has been emphasized in the response to stress, recent studies of Kaufer, et al. (11) have identified an important alternative cholinergic pathway. They showed that acute stress resulted in an immediate increase in synaptosomal acetylcholine with neuronal excitability, with a delayed phase response of increased expression of acetylcholinesterase, decreased choline acetyl transferase and vesicular acetylcholine transporter (CHAT) activity and a resulting decrease in neuronal excitability (11, 12). Others have also emphasized the important role of stress in activating muscarinic systems (13).

**BRIEF SUMMARY OF THE INVENTION**

In one aspect, the present invention relates to a method for screening a patient to determine whether such patient is at increased risk for developing major depression, said method comprising analyzing a sample of said patient's genetic material to determine the patient's genotype with respect to the cholinergic muscarinic receptor 2 (CHRM2) gene, wherein the presence of a T at nucleotide number 1890 in the 3' UTR of said gene correlates with an increased risk for developing major depression.

In another aspect, the present invention relates to a method for diagnosing major depression in a patient, said method comprising analyzing a sample of a patient's genetic material to determine to determine the patient's genotype

with respect to the cholinergic muscarinic receptor 2 (CHRM2) gene, wherein the presence of a T at nucleotide number 1890 in the 3' UTR of said gene is indicative of the presence of major depression.

In another aspect, the present invention relates to a method for screening a subject to determine whether such subject is a candidate for a therapy using a drug which prevents or treats a disorder associated with the presence of a T at nucleotide number 1890 in the 3' UTR of the cholinergic muscarinic receptor 2 (CHRM2) gene, said method comprising determining the subject's genotype with respect to such gene, wherein a subject having a T at said nucleotide number is a candidate for such therapy.

In another aspect, the present invention relates to a method for treating a patient having or at increased risk for developing major depression associated with the presence of a T at nucleotide number 1890 in the 3' UTR of the cholinergic muscarinic receptor 2 (CHRM2) gene, said method comprising administering to said patient an effective amount of a material which diminishes the effect of such gene.

In another aspect, the present invention relates to a method for identifying materials that can be used in the treatment of a patient having or at increased risk for developing a disorder associated with the presence of a T at nucleotide number 1890 in the 3' UTR of the cholinergic muscarinic receptor 2 (CHRM2) gene, said method comprising determining whether the material is capable of diminishing the effect of such gene.

In another aspect, the present invention relates to a pharmaceutical composition which comprises

- a) an effective amount of a material which is capable of diminishing the effect of the cholinergic muscarinic receptor 2 (CHRM2) gene having a T at nucleotide number 1890 in the 3' UTR; and
- b) a pharmaceutically acceptable carrier.

In another aspect, the present invention relates to a kit suitable for screening a subject to determine whether such subject is at increased risk for having or developing a disorder associated with the presence of a T at nucleotide number 1890 in the 3' UTR of the cholinergic muscarinic receptor 2 (CHRM2) gene, said kit comprising

- a) material for determining the subject's genotype with respect to said gene;
- b) suitable packaging material; and optionally
- c) instructional material for use of said kit.

**DETAILED DESCRIPTION OF THE INVENTION**

The present invention is based on the observation that subjects, particularly females, having a T at nucleotide number 1890 in the 3' UTR of the cholinergic muscarinic receptor 2 (CHRM2) gene have an increased risk of having major depression.

The present invention entails the determination of the subject's genotype with respect to the CHRM2 gene. That gene is located on chromosome 7q31-q35. See reference (23) and the references cited therein, the contents of which are incorporated herein by reference. The relevant portion of the sequence is as follows (the T nucleotide at position 1890 is indicated in bold type):  
 acatgggaat taggcaggta gacacagtaa  
 tcatgcaggg **gaagggagat** 50 ttgggagaaa ataatgttgt taaaggag  
 aacaacatt atgtatttta 100 aaccaatgtt tatattatgt ttgttaattt tatte-  
 tattt ccttcagggt 150 ttaaagtgtt attigtacta tggtactga ttgagaacg  
 caaatgaat 200 aactcaaca actcctctaa caatagcctg gctcttaca

3

gtcctataa 250 gacattfgaa ggggtgitta ttgtcctggt ggctggatcc  
 ctacgtttgg 300 tgaccattat cgggaaacatc ctagtcattgg ttccattaa  
 agtcaaccgc 350 caccctccaga ccgtaacaaa ttacttttta ttacgtttgg  
 cctgtgctga 400 cctatcataa ggtgttttct ccaatgaact gtacaccctc  
 tacactgtga 450 ttggtactg gctttggga cctgtggtgt gtagctttg  
 gctagccctg 500 gactatgtgg tcagcaatgc ctacgttatg aatctgctca  
 tcaacagctt 550 tgacaggtac ttctgtgca caaaacctt gacctacca  
 gtaagcgga 600 ccacaaaaat ggcaggtatg atgattcgag ctgctcgggt  
 cctctcttc 650 atcctctggg ctccagccat tctctctgg cagttcattg  
 taggggtgag 700 aactgtggag gatggggagt gctacattca gtttttcc  
 aatgtgctg 750 tcacittfg tacggctatt gcagccttct attgccagt  
 gatacatg 800 actgtgctat attggacat atcccagcc agcaagagca  
 ggataaagaa 850 ggacaagaag gagcctgtg ccaaccaaga  
 ccccccttct ccaagtctg 900 tacaaggaag gatagtgaag ccaacaata  
 acaacatgcc cagcagtgac 950 gatggcctgg agcacaacaa atcca-  
 gaat ggcaagcccc ccagggatcc 1000 tgtgactgaa aactgtgtc agg-  
 gagagga gaaggagagc tcaatgact 1050 ccacctcagt cagtgtctgt  
 gctcttaata tgagagatga tgaataaacc 1100 caggatgaaa acacagttc  
 cacttccctg ggcatttcca aagatgaaa 1150 ctctaagcaa acatgcatca  
 gaattggcac caagaccca aaaagtact 1200 catgtaccce aacta-  
 atatacc accgtggagg tagtggggc tccaggtcag 1250 aatggagatg  
 aaaagcagaa tattgtagcc cgaagattg tgaagatgac 1300 taag-  
 cagcct gcaaaaaaga agcctctcc tcccgggaa aagaaatgca 1350  
 ccaggacaat ctggctatt ctgttgctt tcaatcac ttgggcccc 1400  
 tacaatgca tgggtctcat taacacctt tgtcacctt gcatcccaa 1450  
 cactgtgtg acaattggt actggcttg ttactcaac agcactatca 1500  
 accctcctg ctatgcaact tgcattgcca cctcaagaa gacctttaa 1550  
 caccttctca tgtgtcatta taagaacata ggcgctcaaa ggtaaaatat 1600  
 cttgaaaaa gatagaaggt gggcaagggg agcttgaaa gaataaagg  
 1650 gataaacgag ctctagtgt taaaatctt gccattgac ttatagct  
 1700 gattacaaa cgtgcaatc agggagccag cagtacaca ctat-  
 cagc 1750 ctaggctcca gttgcaaaa atfgacctt ataaactgc  
 agtattagga 1800 gcaatgagac atgaaagaa acatgtggg atctg-  
 gatt taagaaacta 1850 tacactgtt ctataatct ctgagaag ggcttct-  
 gat tcaaat 1900 tatcagctc tgcacaagag gaataacctt gtctcttt  
 1950 gttgtgtg ttctcatg tcttaagag aaggaatgcc  
 acagttaca 2000 ggtaaacatg gagacttaa cataaagaaa taggcac-  
 tat acaatgggga 2050 cataaaaaa gaaatgaaa gaaggatgca  
 gaaattgtc tccggagtgt 2100 taagcatatt ttactcttt gttacggctc  
 tatttagagg atggaaatgt 2150 aataaatgct tatttttgc cttctttt  
 2200 cccaccatga agagaagca 2200 acaaacaga 2210 [SEQ ID  
 NO:1].

Such can be determined, for example, by analysis of the subject's DNA. Suitable analysis techniques are well known to those in the art, and include amplification genotyping (amplification of the desired region by suitable methods, such as PCR, followed by electrophoresis), in situ hybridization techniques, direct DNA sequencing, etc.

A further aspect of the present invention is the treatment of a disorder associated with the presence of the allele described herein, to prevent the development or progression of such disorder. The patient is administered an effective amount of a material which diminishes or eliminates the adverse effects of the allele. The material may act in a number of ways which would be apparent to one of ordinary skill. For example, it may act to decrease the production of the protein, such as by affecting the DNA or RNA responsible for protein production, or by affecting regulatory elements. One way to accomplish diminished protein production is by introduction via gene therapy or gene repair techniques of a gene or gene segment which converts the allele associated with the disorder into a benign allele. See, for example, the techniques described in U.S. Pat. No. 5,776,744, the contents of which are incorporated herein by reference. The material may also act by directly or indirectly affecting the produced protein to diminish the protein's activity or effect.

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It will be apparent that the information regarding a subject's genotype with respect to the CHRM2 gene may also be used to determine whether the subject is a candidate for a therapy using a drug which prevents or treats a disorder associated with the CHRM2 gene in question.

For therapeutic treatment, the materials of the present invention may be formulated into a pharmaceutical composition, which may include, in addition to an effective amount of the active ingredient, pharmaceutically acceptable carriers, diluents, buffers, preservatives, surface active agents, and the like. Compositions may also include one or more other active ingredients if necessary or desirable.

The pharmaceutical compositions of the present invention may be administered in a number of ways as will be apparent to one of ordinary skill in the art. Administration may be done topically, orally, by inhalation, or parenterally, for example.

Topical formulations may include ointments, lotions, creams, gels, drops, suppositories, sprays, liquids and powders. Oral formulations include powders, granules, suspensions or solution in water or non-aqueous media, capsules or tablets, for example. Thickeners, flavorings, diluents, emulsifiers, dispersing aids or binders may be used as needed.

Parenteral formulations may include sterile aqueous solutions which may also contain buffers, diluents and other suitable additives.

The dose regimen of the compounds or compositions of the present invention will depend on a number of factors which may readily be determined, such as severity and responsiveness of the condition to be treated.

The present invention also provides a screening method for identifying materials that may be used in the treatment of a patient having or at increased risk for developing a disorder associated with the presence of the CHRM2 gene having a T at nucleotide number 1890 in the 3' UTR. In practice of such a method, a candidate material is screened in an assay which determines whether the material is capable of diminishing the effect of the gene in question. Suitable assays would be readily apparent to one of ordinary skill, including animal models and in vitro assays. The assays may be designed to test, for example, the effect of the material on the production of the particular protein, or its effect on the activity of the protein.

The present invention also provides a kit suitable for screening a subject for any of the purposes described above. The kit comprises material for determining the subject's genotype with respect to the CHRM2 gene. Preferably, such material comprises at least two primers capable of hybridizing to a region flanking nucleotide number 1890 in the 3' UTR segment of the CHRM2 gene. The primers preferably are suitable for use in an amplification reaction, such as PCR. The kit additionally contains suitable packaging material, and optionally contains instructional material for use of the kit, result interpretation, etc. The kit may also contain additional reactants suitable for use with the primers, such as appropriate concentrations of deoxynucleotide triphosphates, suitable buffers, polymerization enzymes, etc.

The following non-limiting examples are illustrative of the processes and products of the present invention.

EXAMPLE 1

Subjects. The sample consisted of 760 non-Hispanic Caucasian adults from the Minnesota Twin and Family

Study (MTFS) (14). The MTFS is a large, multi-discipline, multi-year study to examine the interaction between genetic and environmental risk factors in the development of adolescent and adult alcoholism and drug abuse. The advantage of the study is that it uses a population based twin ascertainment in which all same sex twins born in the state of Minnesota are identified by public birth records. The recruitment targets 11 and 17 year old twins. Of the eligible families only 17% declined invitations to participate. The present study was restricted to the parents of the twins. They were administered the parent version of the DICA-R (Diagnostic Interview for Children and Adolescents (15) and the Structured Clinical Interview for DSM-III-R (SCID-R) (16). Interviews were administered by individuals who have a bachelor's or master's degree in psychology or a related field. Interviewers also complete an intensive course of training that includes didactic instruction, practice interviews, mentoring by an experienced clinical interviewer, and a written examination covering the DSM disorders assessed. All interviews are tape-recorded. Complete interviews are reviewed in a consensus conference by at least two advanced clinical psychology graduate students. Individual symptoms are reviewed, including listening to the audio tapes as needed, to determine whether the behaviors reported by the interviewees were frequent and severe enough to count as a symptom under DSM. In a study of the reliability of the diagnostic and consensus procedures that involved review of clinical material by two independent teams for clinicians, the kappa coefficient for the depression diagnosis was estimated at 0.82 (14).

We genotyped 430 women and 330 men. Of these, 126 of the women and 52 of the men had a DSM-III-R diagnosis of definite lifetime MD.

Genotyping. Using the SSCP technique (17) we identified a common single nucleotide polymorphism, A→T 1890 in the 3' UTR of the CHRM2 gene based on accession # M16404. The upstream primer was 5'-ACAAAACGTG CAATTCAGGA G-3' [SEQ ID NO:2]. The downstream primer was 5'-CAGAGACTGA TAAATTTGTA G-3' [SEQ ID NO:3]. The PCR reaction consisted of QIAGEN 10x PCR buffer 2.0  $\mu$ l, 0.4  $\mu$ l of 10 mM dNTP, 0.4  $\mu$ l of 10  $\mu$ M each primer, 0.5 units of Taq polymerase, ddH<sub>2</sub>O 16.1  $\mu$ l and 50 ng DNA in total volume of 20  $\mu$ l. Amplification consisted of 30 cycles with denaturation at 95° C. for 30 seconds, annealing temperature 54° C. for 1 minute, and extension at 72° C. for 1 minute. This produced a 208 bp product, which was digested with Dpn II restriction endonuclease using 1  $\mu$ l 10x New England Biological (NEB) enzyme buffer, 0.3  $\mu$ l of Dpn II, 3.7  $\mu$ l of double distilled H<sub>2</sub>O, and 5  $\mu$ l PCR product incubated at 37° C. for 3 hours. The product were visualized by electrophoresis in 10% acrylamide gel. The A allele gave 81 and 127 bp products. The T allele gave 58+23 and 127 bp products.

Statistics. The frequency of the 11, 12 and 22 genotypes in male and female subjects with and without a diagnosis of MD was compared by Chi square analysis. A Bonferroni corrected  $\alpha$  of 0.025 was used. To determine the percent of the variance of MD, accounted for by the CHMR2 gene, we scored those with the 11 genotype as 1, and those with the remaining two genotypes as 0. Those without MD were scored as 0, and those with MD as 1. The presence or absence of MD was the dependent variable and the gene score was the independent variable in a linear regression analysis. The SPSS statistical package was used (SPSS, Inc, Chicago, Ill.).

#### Results

The results are shown in Table 1. For the women without MD, 25.7% carried the 11 genotype compared to 43.7% for with MD. There was a proportionate decrease in the frequency of the other two genotypes ( $\chi^2=13.53$ , d.f.=1.,

$p=0.001$ ). By contrast for men, while the frequency of the 11 genotype in those without MD was similar to that for the women (27.7%), there was no increase in frequency for the 11 genotype for those with MD (26.9%,  $\chi^2=1.48$ , d.f.=1.,  $p=0.47$ ). The CHRM2 alleles were in Hardy-Weinberg equilibrium for those without MD. Regression analysis showed that in women  $r^2=0.030$ ,  $F=13.37$ ,  $p=0.0003$ . By contrast, in men  $r^2=0.00001$ ,  $F=0.002$ ,  $p=0.96$ .

#### Discussion

These results suggest that CHRM2 may be a gene associated with MD in women but not in men. This association with MD is consistent with the postulated role of enhanced or hypersensitive cholinergic systems in depression (8, 9). The association of cholinergic systems with REM sleep (10, 18), and the disturbance of REM sleep in individuals with depression are consistent with the presence of genetic defects in the cholinergic system in MD.

In women, the CHRM2 gene accounted for 3 percent of the variance of MD while in men it accounted for virtually none of the variance. The increased frequency of MD in females, and some twin studies (5) suggesting a greater heritability of MD in women, raise the possibility that the CHRM2 is a gender specific gene for MD. While the mechanism by which a non-X-linked gene would have this effect is not known, there are several possibilities. These include hormone responsive promoters or enhancers, affecting either CHRM2 other genes that interact with CHRM2. A second possibility is that if women are more sensitive to stress expressed through the cholinergic stress pathway (11, 12, 13), they would also be more likely to show an association between the CHRM2 gene and depression.

Since the A/T polymorphism was in the 3' region of the CHRM2 gene we assume it was in linkage disequilibrium with alleles affecting gene function, possibly microsatellites (19). The fact that even though the CHRM2 gene accounted for only 3 percent of the variance of MD in women, even though the association was significant, is an expected characteristic of a complex polygenic disorder where multiple genes are involved and each accounts for only a small percent of the variance. In our experience, 3 percent is a higher than the average  $r^2$  for most gene-phenotype associations for behavioral disorders (20, 21, 22).

Table 1. Association of the CHRM2 Gene with Major depression in women and men in the Minnesota Twin and Family Study

TABLE 1

Association of the CHRM2 Gene with Major depression in women and men in the Minnesota Twin and Family Study						
CHRM2 genotypes						
	N	11	12	22	X <sup>2</sup>	p
<b>Women</b>						
No MD	304	78 (25.7)	159 (52.3)	67 (22.0)		
Definite MD	126	55 (43.7)	49 (38.9)	22 (17.5)	13.53	.001
Total	430					
<b>Men</b>						
No MD	278	77 (27.7)	122 (43.9)	79 (28.4)		
Definite MD	52	14 (26.9)	27 (51.9)	11 (21.2)	1.48	.47
Total	330					

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SEQUENCE LISTING

<160> NUMBER OF SEQ ID NOS: 3

<210> SEQ ID NO 1  
 <211> LENGTH: 2210  
 <212> TYPE: DNA  
 <213> ORGANISM: Human

<400> SEQUENCE: 1

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ttgttaattt tattctattt ccttcaggt taaatggtt atttgctact tggctactga      180
ttagagaacg caaaatgaat aactcaaaa actcctctaa caatagcctg gctcttacia      240
gtccttataa gacattttaa gtggtgttta ttgtcctggt ggctggatcc ctcagtttgg      300
tgaccattat cgggaacatc ctagtcatgg tttccattaa agtcaaccgc cacctccaga      360
cogtcaaaa ttacttttta ttcagcttgg cctgtgctga ccttatcata ggtgttttct      420
ccatgaactt gtacaccctc tacactgtga ttggttactg gcctttggga cctgtggtgt      480
    
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-continued

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What is claimed is:

- 1. A method for screening a female patient to determine whether such patient is at increased risk for developing major depression, said method comprising analyzing a sample of said patient's genetic material to determine the patient's genotype with respect to the cholinergic muscarinic receptor 2 (CHRM2) gene, wherein the presence of a T at nucleotide number 1890 in the 3' untranslated region (3' UTR) of said gene [SEQ ID NO:1] correlates With an increased risk for developing depression.
- 2. The method of claim 1, wherein the analysis utilizes an amplification reaction.
- 3. The method of claim 2, wherein the amplification reaction utilizes PCR.
- 4. The method of claim 3, wherein the PCR reaction utilizes a primer selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:3.
- 5. A method for diagnosing major depression in a female patient, said method comprising analyzing a sample of a female patient's genetic material to determine the patient's genotype with respect to the cholinergic muscarinic receptor 2 (CHRM2) gene, wherein the presence of a T at nucleotide number 1890 in the 3' untranslated region (3' UTR) of said gene [SEQ ID NO: 1] is indicative of the presence of major depression.
- 6. The method of claim 5, wherein the analysis utilizes an amplification reaction.
- 7. The method of claim 6, wherein the amplification reaction utilizes PCR.
- 8. The method of claim 7, wherein the PCR reaction utilizes a primer selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:3.

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- 9. A method for testing a sample of female human genetic material, said method comprising the step of determining whether a T is present at nucleotide number 1890 in the 3' untranslated region (3' UTR) of the cholinergic muscarinic receptor 2 (CHRM2) gene [SEQ ID NO: 1].
- 10. The method of claim 9, wherein the determination utilizes an amplification reaction.
- 11. The method of claim 10, wherein the amplification reaction utilizes PCR.
- 12. The method of claim 11, wherein the PCR reaction utilizes a primer selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:3.
- 13. A kit suitable for screening a female subject to determine whether such subject is at increased risk for having or developing major depressive disorder associated with the presence of a T at nucleotide number 1890 in the 3' untranslated region (3' UTR) of the cholinergic muscarinic receptor 2 (CHRM2) gene [SEQ ID NO:1], said kit comprising
  - a) material for determining the subject's genotype with respect to said gene;
  - b) suitable packaging material; and optionally
  - c) instructional material for use of said kit wherein item a) comprises at least one PCR primer selected from the group consisting of SEQ ID NO: 2, SEQ ID NO: 3, and combinations thereof.

\* \* \* \* \*

UNITED STATES PATENT AND TRADEMARK OFFICE  
**CERTIFICATE OF CORRECTION**

PATENT NO. : 6,743,589 B2  
DATED : June 1, 2004  
INVENTOR(S) : David E. Comings et al.

Page 1 of 1

It is certified that error appears in the above-identified patent and that said Letters Patent is hereby corrected as shown below:

Title page,

Item [73], Assignee, please insert -- City of Hope, Duarte, California; --

Item [56], **References Cited**, U.S. PATENT DOCUMENTS, "Comings et l" should be -- Comings et al. --;

Column 8,

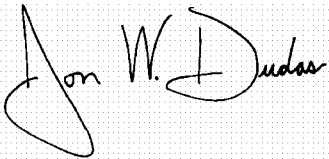
Line 8, "Weiner" should be -- Welner --;

Column 12,

Line 26, insert a comma after "said kit".

Signed and Sealed this

Fifteenth Day of March, 2005

A handwritten signature in black ink on a light gray dotted background. The signature reads "Jon W. Dudas" in a cursive style. The "J" is large and loops around the "on". The "W" and "D" are also prominent.

JON W. DUDAS

*Director of the United States Patent and Trademark Office*