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(54) **CONFORMATIONAL-SPECIFIC ANTIBODIES AGAINST A[BETA] OLIGOMERS**

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(57) **ABSTRACT**

The present disclosure relates to a conformational specific antibody molecule recognizing a conformational epitope of human trimeric or tetrameric oligomers, wherein said antibody molecule does not bind human monomeric Aβ 1-42 or human Ap 1-42 dimers; and pharmaceutical and diagnostic compositions comprising said antibody molecule as well as methods using same.

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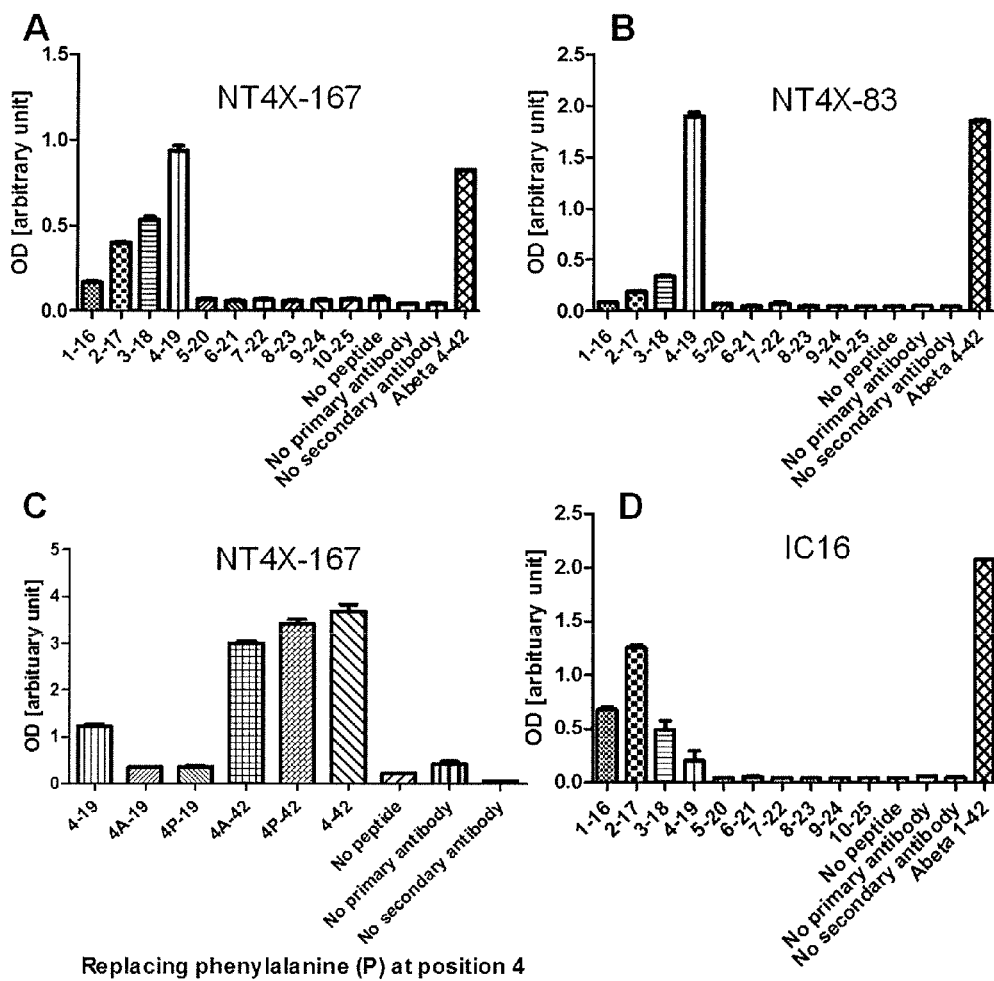
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§ 371 (c)(1),

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Figure 1



Replacing phenylalanine (P) at position 4

Figure 2

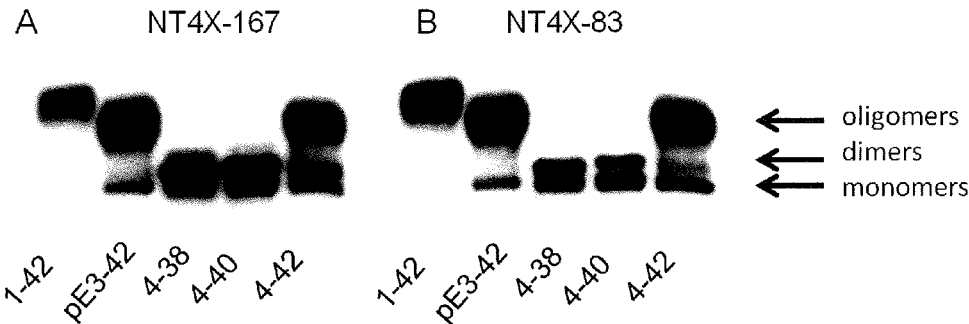


Figure 3

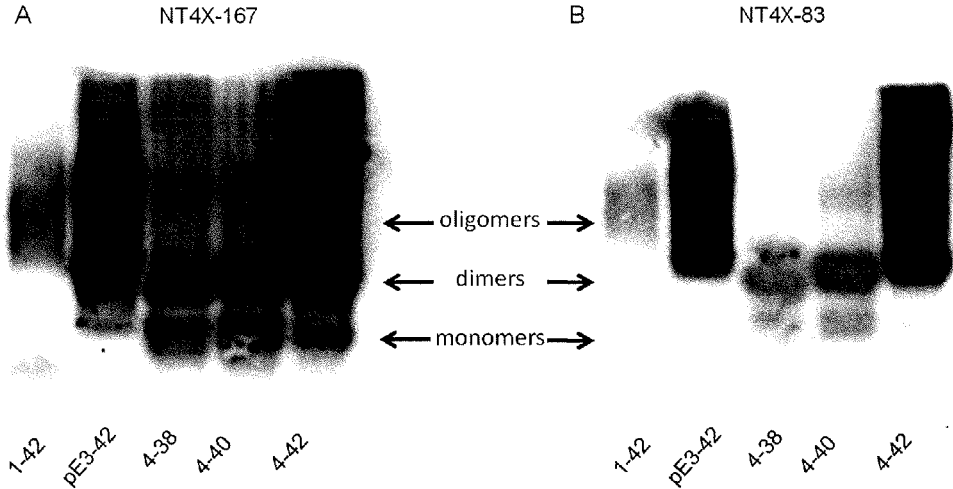


Figure 4

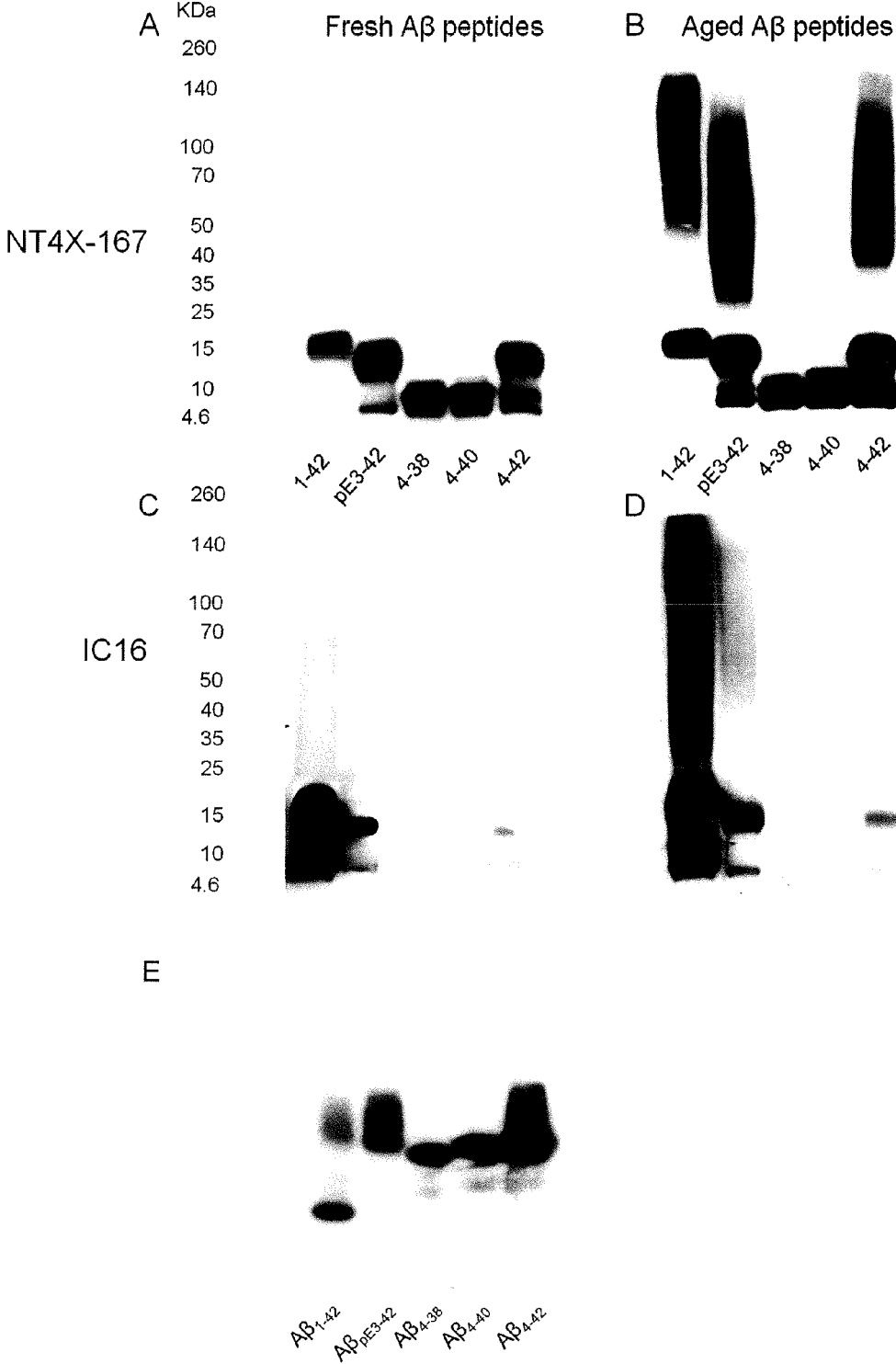


Figure 5

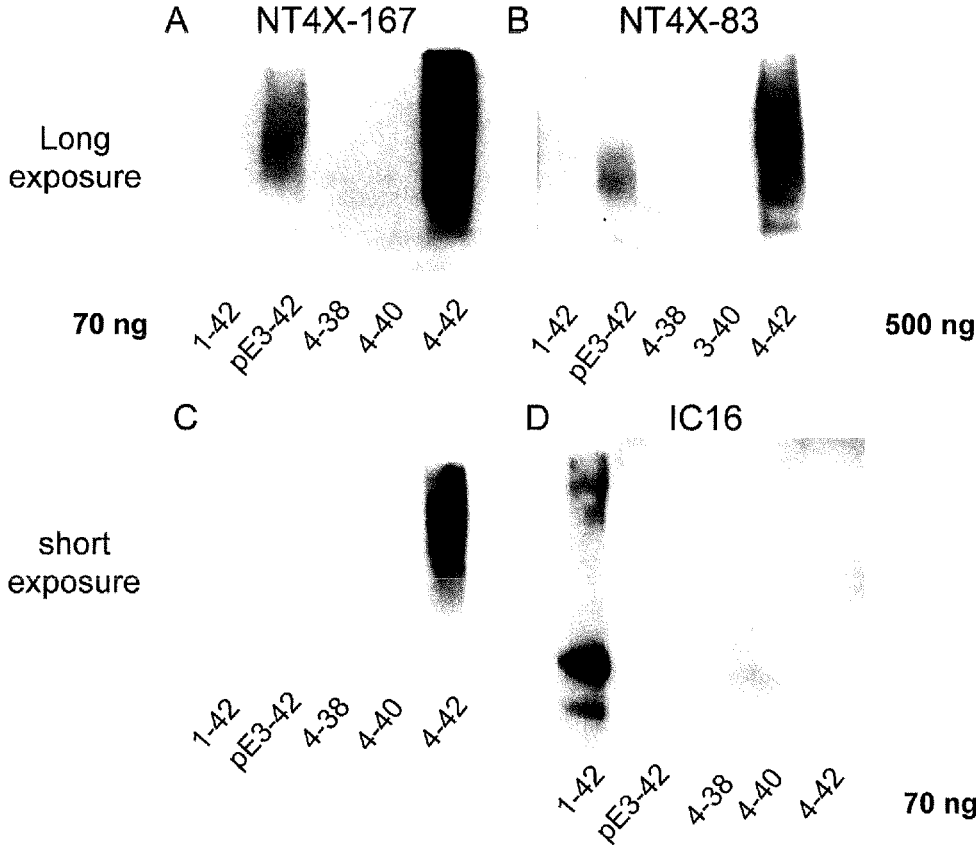


Figure 6

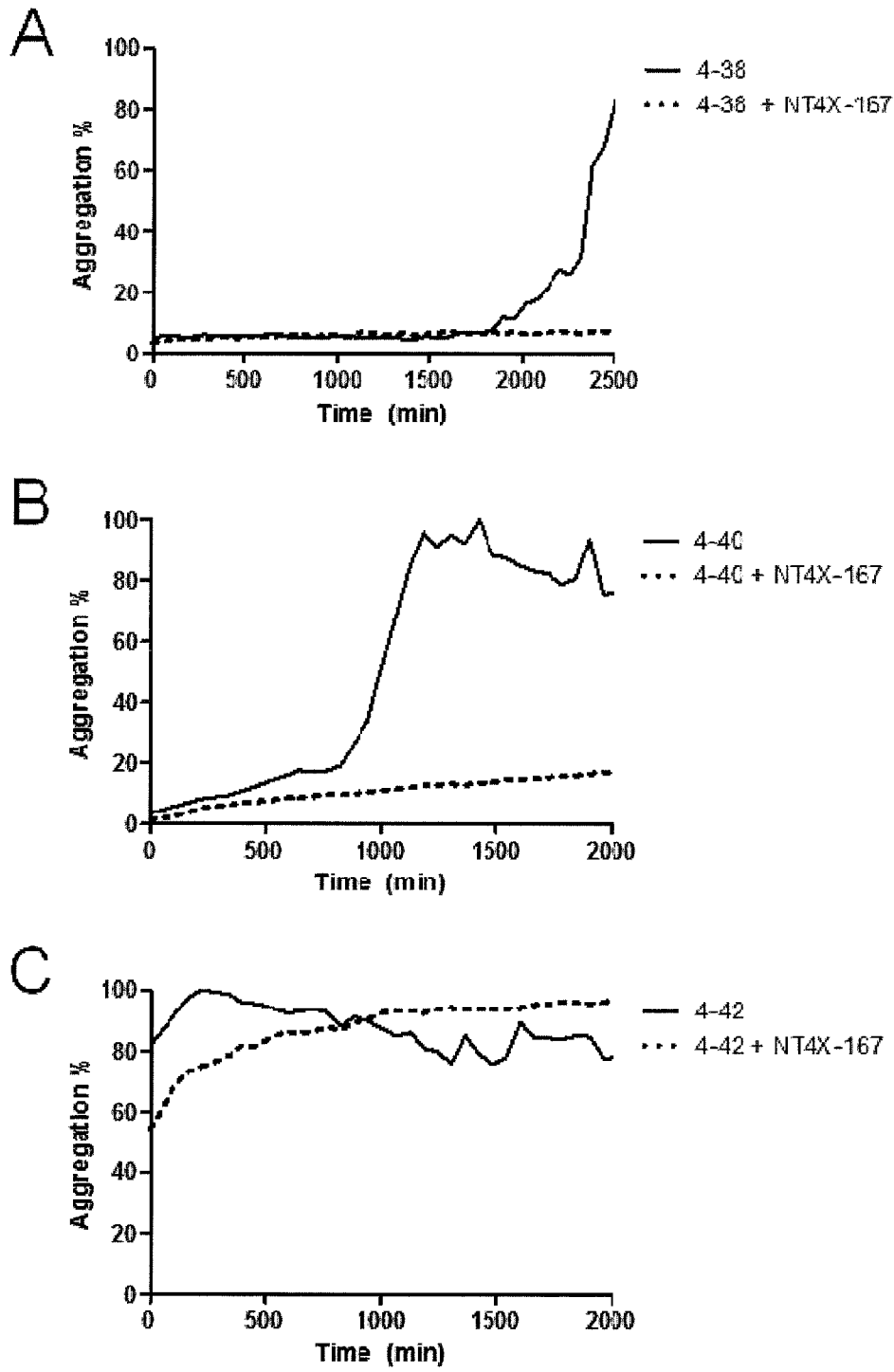


Figure 6 continued

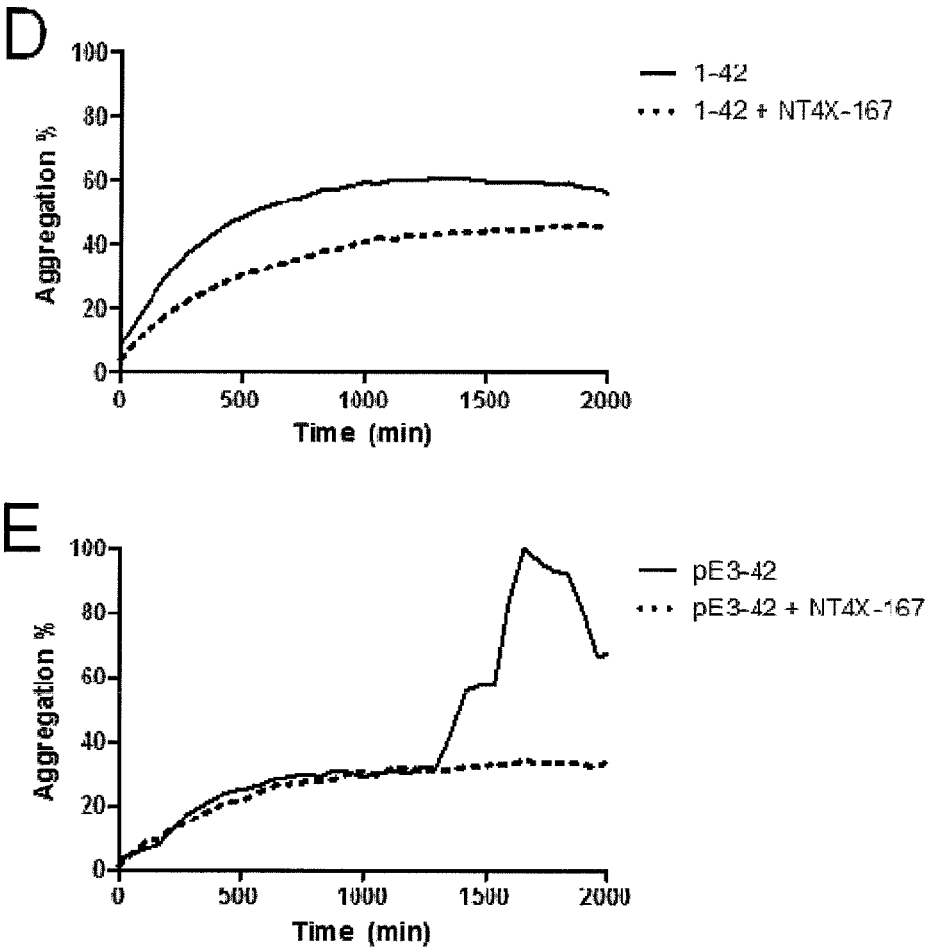


Figure 7

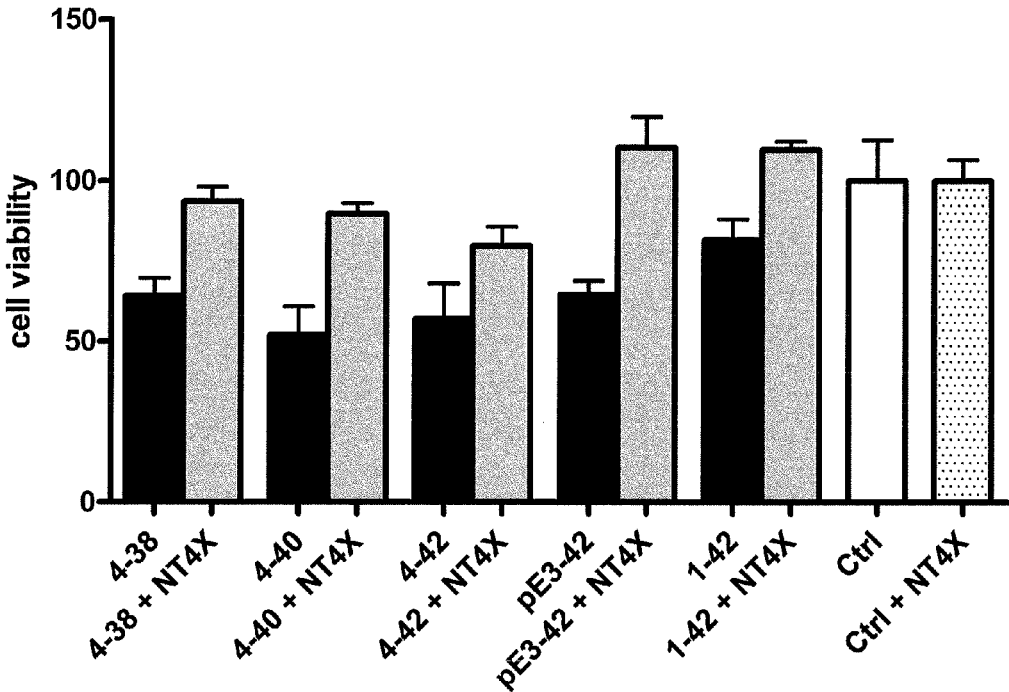




Figure 8

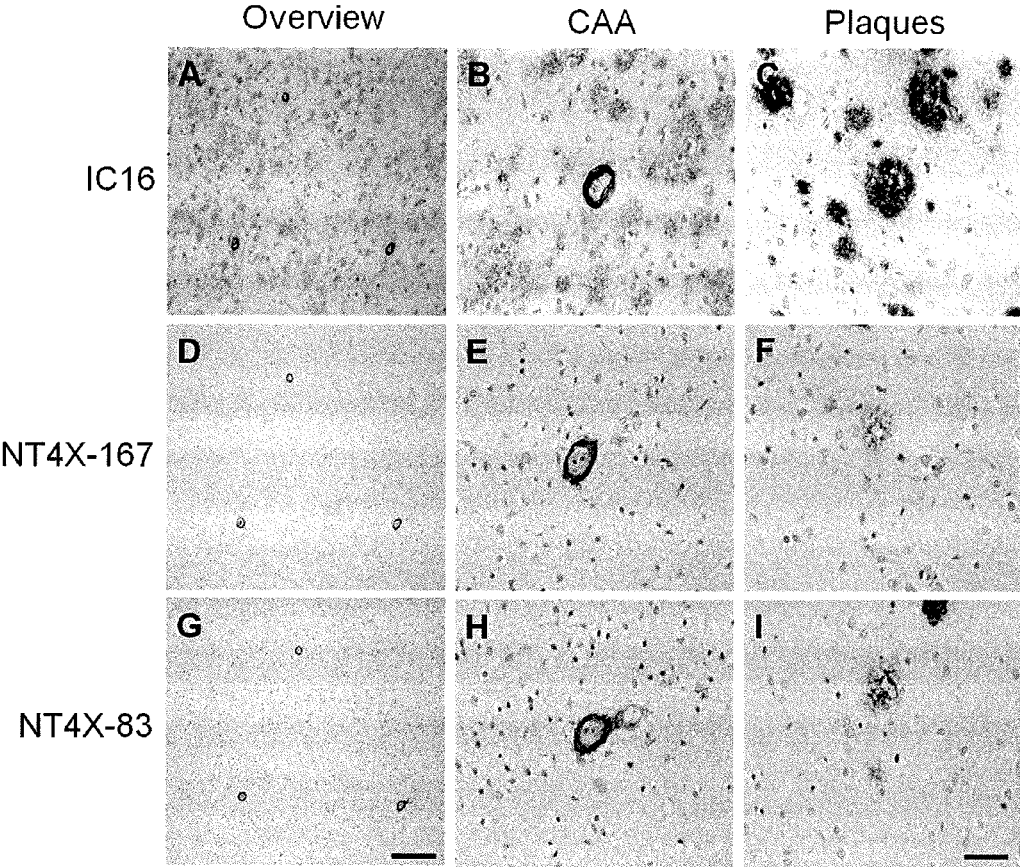


Figure 9

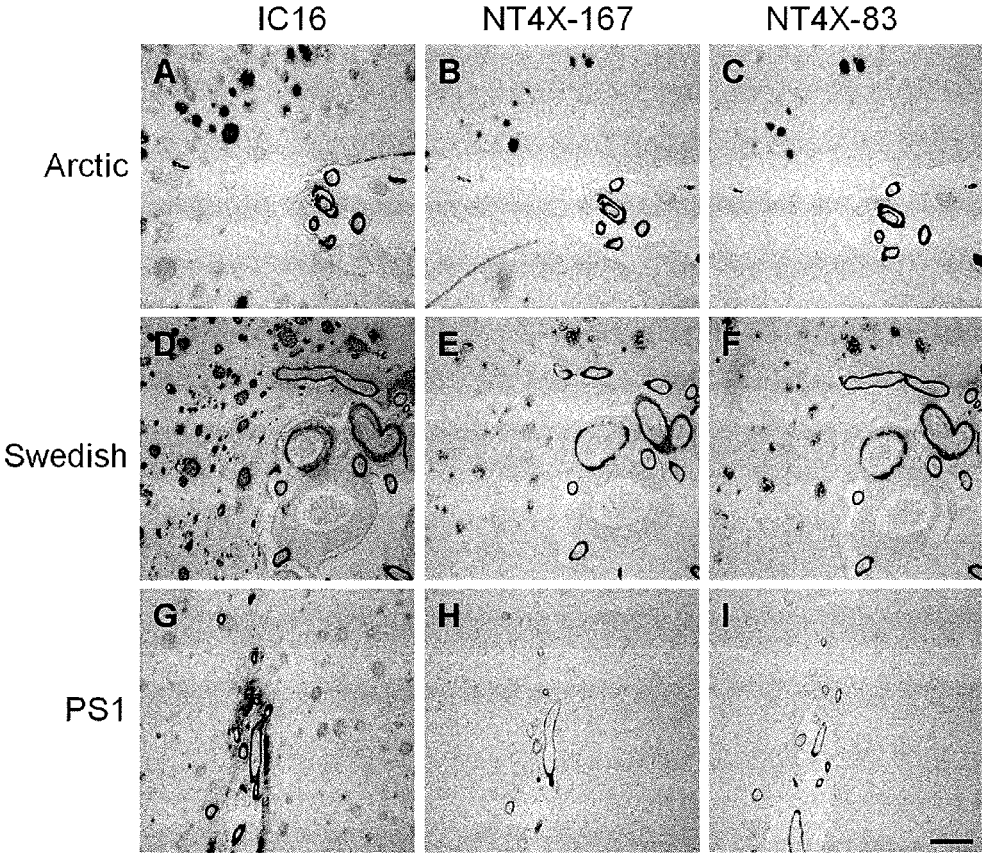


Figure 10

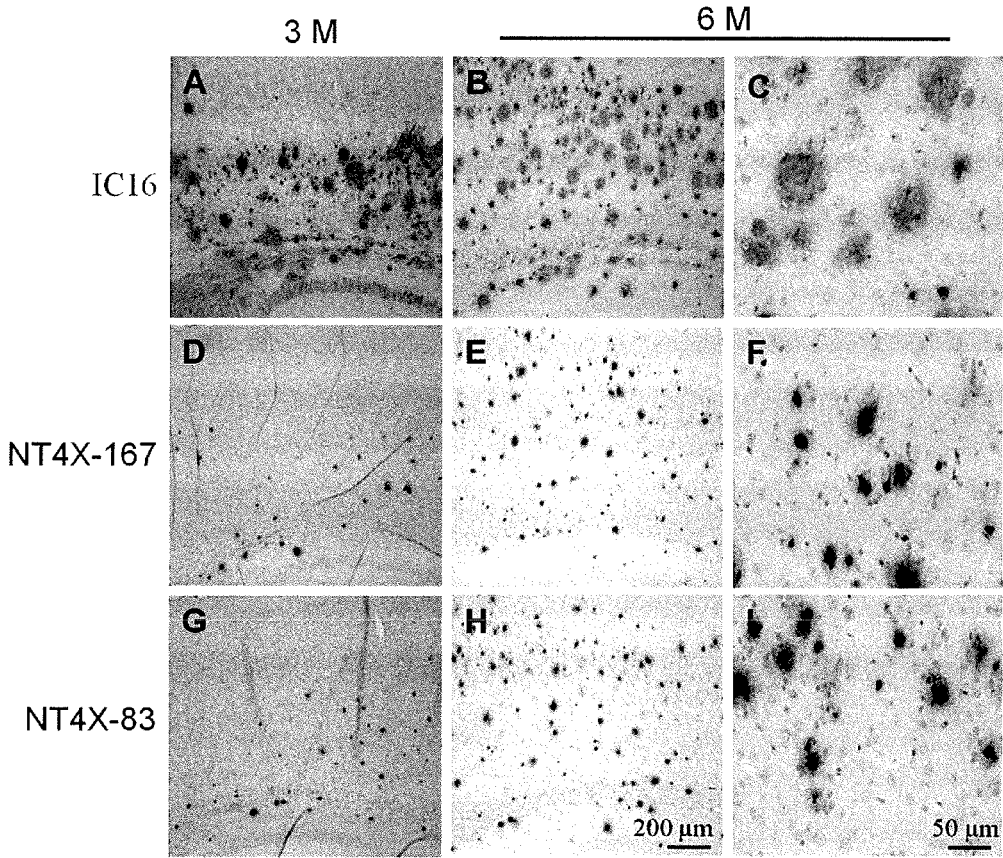


Figure 11

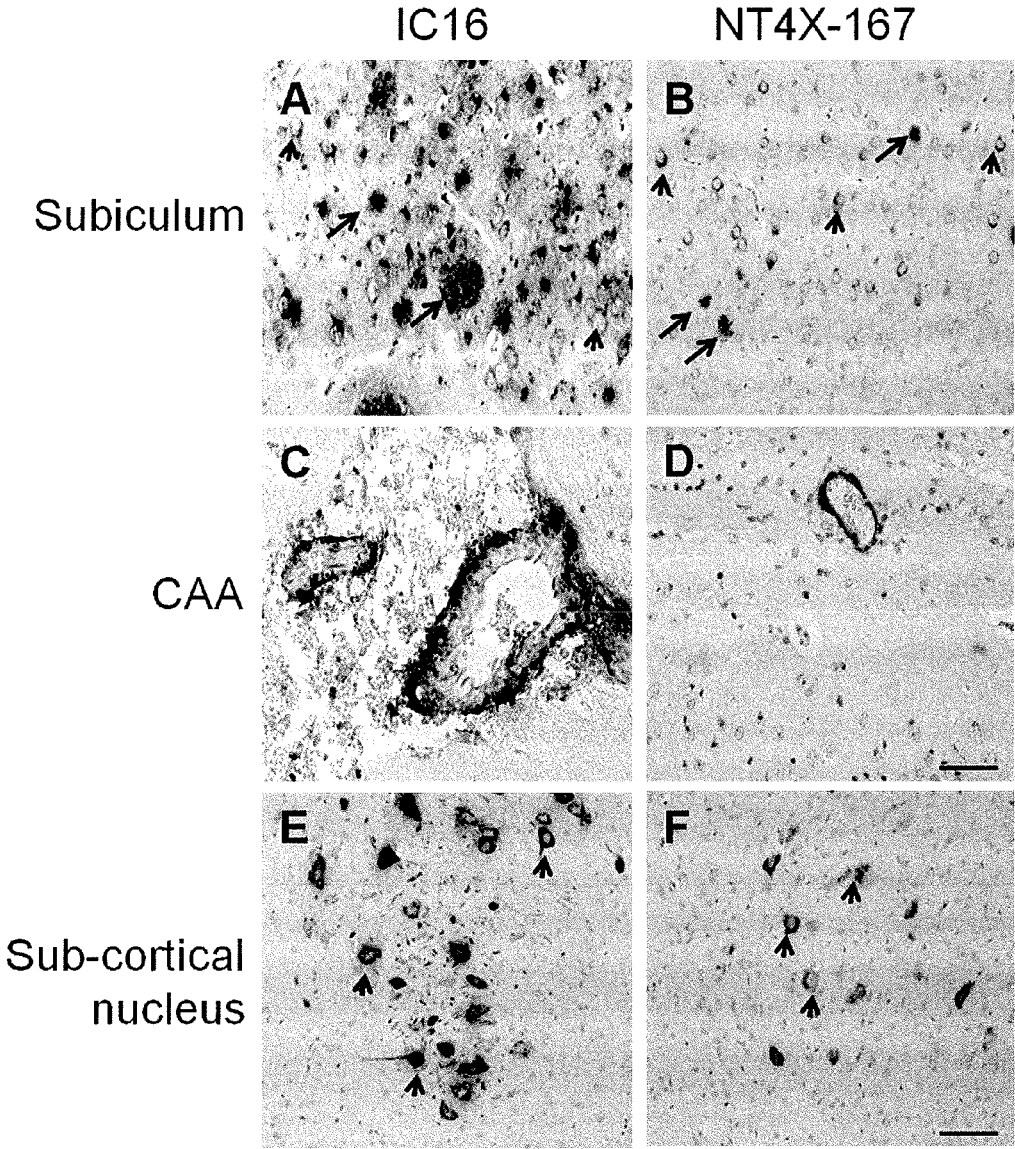


Figure 12

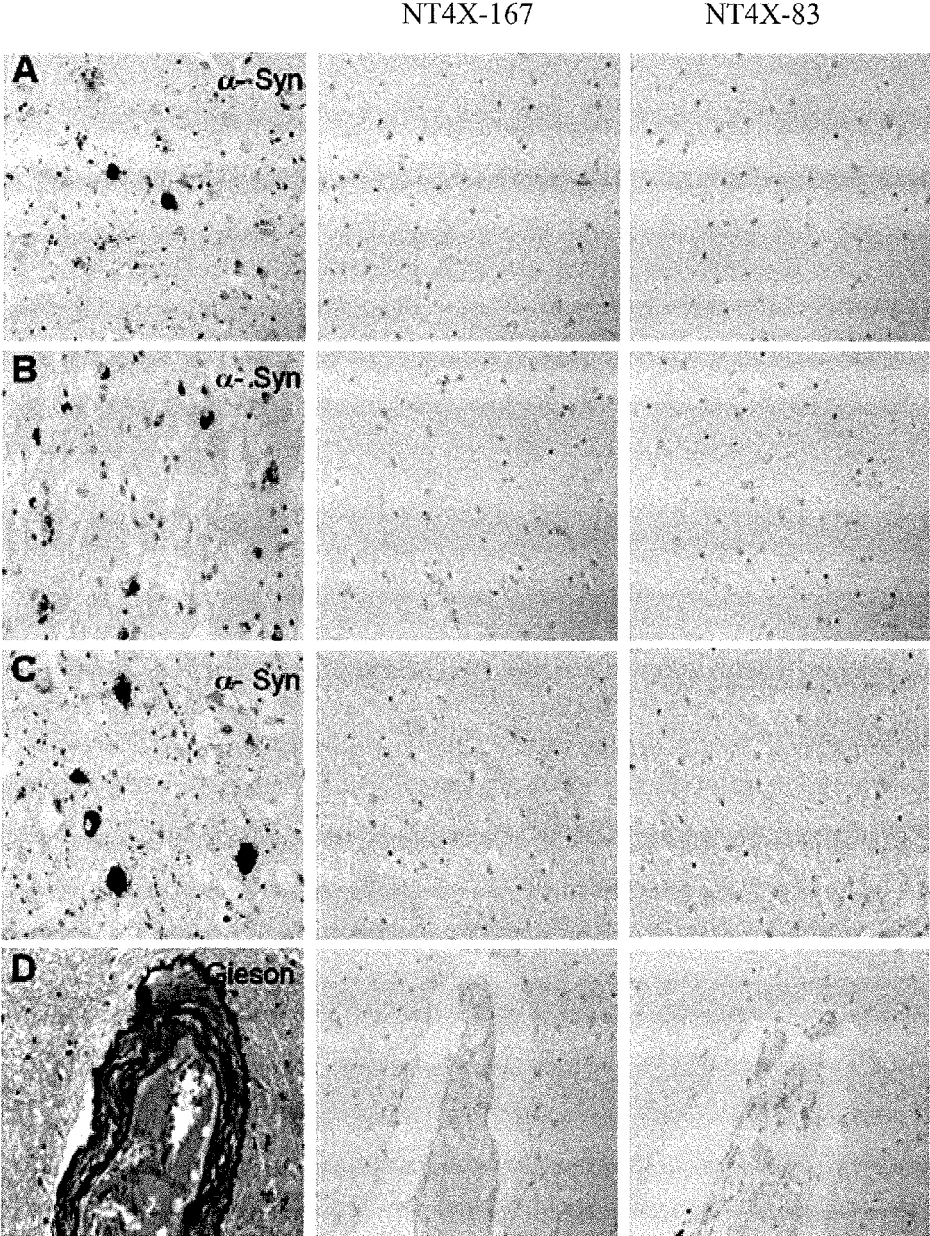


Figure 12 continued

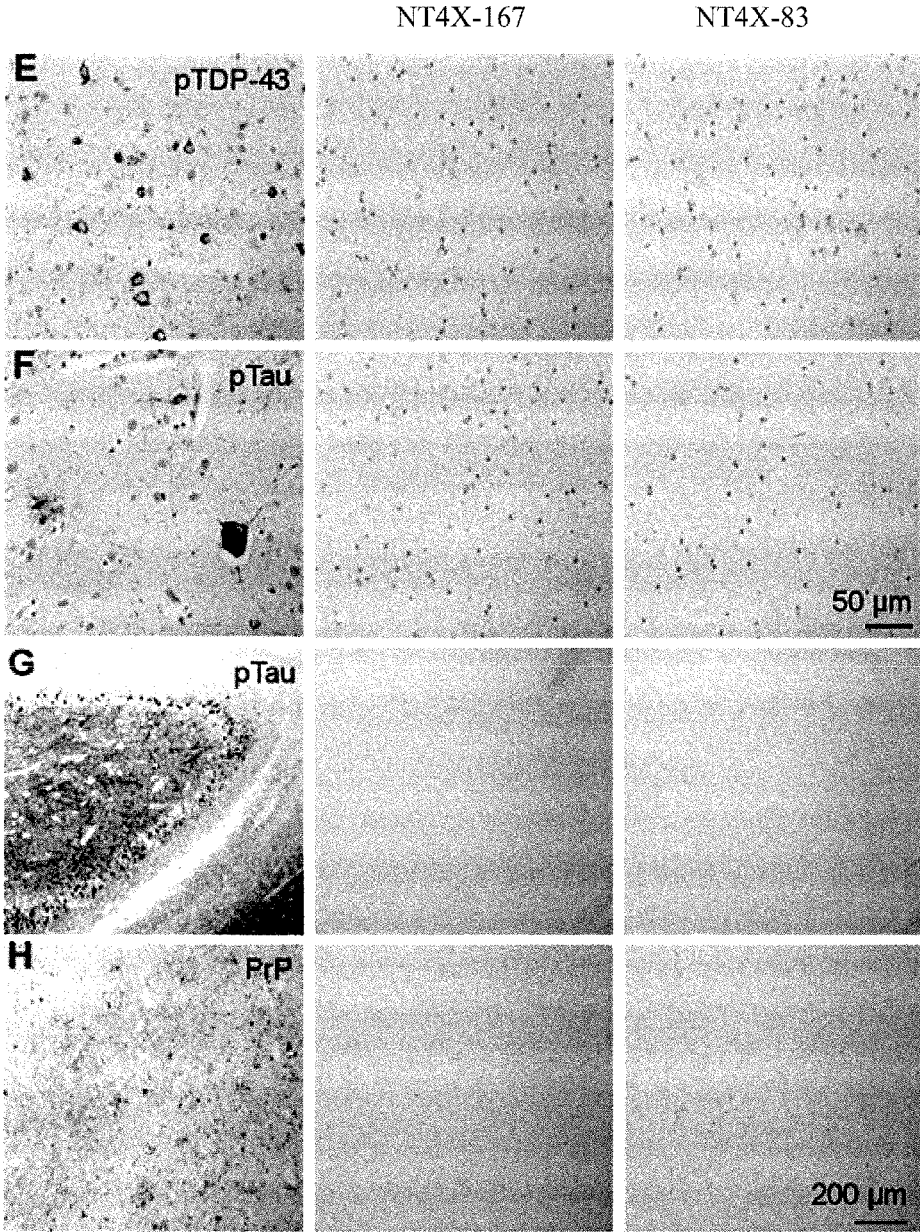


Figure 13

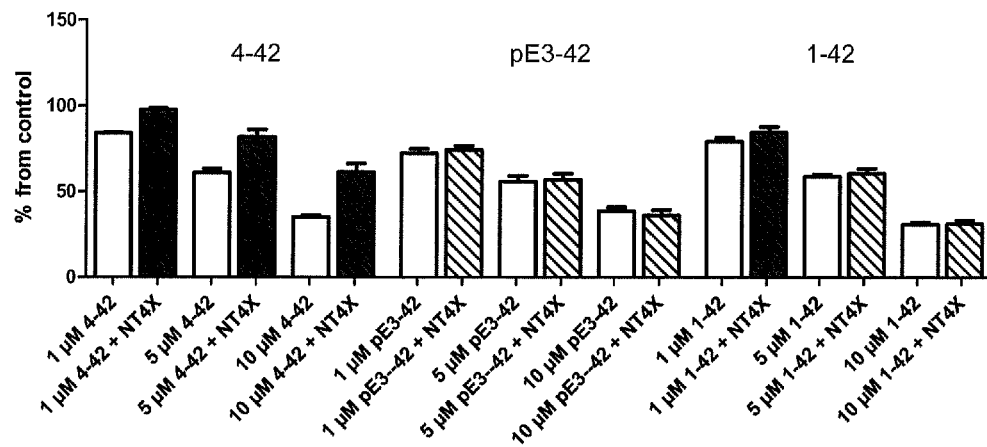
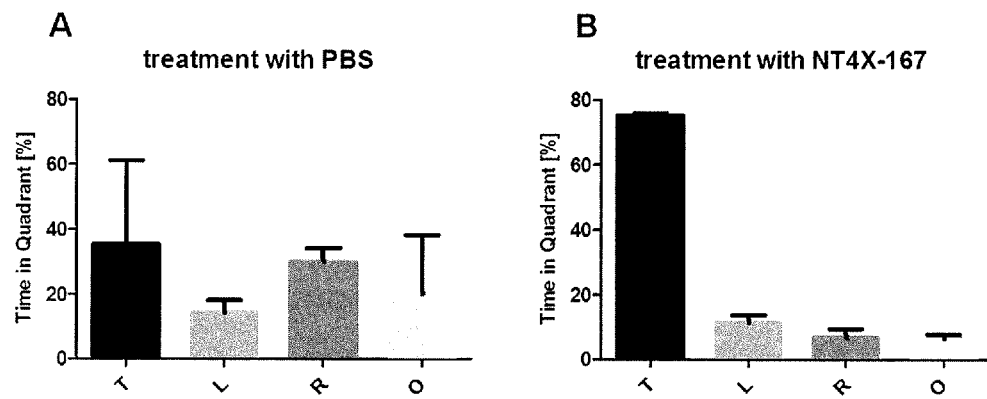


Figure 14



### CONFORMATIONAL-SPECIFIC ANTIBODIES AGAINST A[ $\beta$ ] OLIGOMERS

**[0001]** The present invention relates to a conformational specific antibody molecule directed against a conformational epitope formed by human trimeric or tetrameric A $\beta$  oligomers, wherein said antibody molecule does not bind human monomeric A $\beta$  1-42 or human A $\beta$  1-42 dimers; and pharmaceutical and diagnostic compositions comprising said antibody molecule as well as methods using same.

#### BACKGROUND OF THE INVENTION

**[0002]** AD represents a common progressive neurodegenerative disorder that is characterized neuropathologically by extracellular deposits composed of the amyloid- $\beta$  (A $\beta$ ) protein and intra-cellular accumulation of phosphorylated tau protein. The story of successful discoveries in modern AD research using novel molecular biological tools started with the biochemical analysis of  $\beta$ -amyloid containing blood vessels (CAA, cerebral amyloid angiopathy) and amyloid plaques consisting of A $\beta$ , which led to the isolation and sequencing of the gene encoding the larger amyloid precursor protein (APP) (Kang, et al. (1987) *Nature* 325, 733-736; Tanzi, et al. (1987) *Science* 235, 880-884).

**[0003]** For more than two decades, the amyloid hypothesis has been the central hypothesis in coining the molecular pathology of AD. According to this hypothesis, amyloid fibrils, which are large insoluble polymers of A $\beta$  found in senile plaques, are the major trigger of neuron loss and dementia typical for AD. Albeit there are convincing genetic, biochemical and cell biological data for a major role of A $\beta$  in AD, growing evidence points towards soluble A $\beta$  oligomers rather than A $\beta$  precipitated in plaques.

**[0004]** Haass and Selkoe have recently discussed the concept that soluble oligomers of A $\beta$  as diffusible assemblies are capable of interfering with synaptic function and integrity providing an important opening for understanding the basis of memory loss in AD. They argued that small soluble oligomers affect synaptic structure and plasticity, whereas large, insoluble plaque deposits might function as reservoirs of the pathological oligomers (Haass and Selkoe (2007) *Nat Rev Mol Cell Biol* 8, 101-112). Moreover, it has also been suggested that the extracellular formation of A $\beta$  plaques and other AD pathological events are preceded by intraneuronal A $\beta$  accumulation giving rise to a modified amyloid hypothesis.

**[0005]** A $\beta$ , composed of mostly 1-40 or 1-42 amino acids, is a critical component in the etiology of the neurodegenerative AD. A $\beta$  is prone to aggregate and forms amyloid fibrils progressively both in vitro and in vivo. To understand the process of amyloidogenesis, it is pivotal to examine the initial stages of the folding process. Recently, the equilibrium folding properties, assembly states, and stabilities of the early folding stages of A $\beta$ <sub>1-40</sub> and A $\beta$ <sub>1-42</sub> prior to fibril formation was examined. It was found that A $\beta$ <sub>1-40</sub> and A $\beta$ <sub>1-42</sub> have different conformations and assembly states upon refolding from their unfolded ensembles. A $\beta$ <sub>1-40</sub> is predominantly an unstable and collapsed monomeric species, whereas A $\beta$ <sub>1-42</sub> populates a stable structured trimeric or tetrameric species at concentrations above ~12.5  $\mu$ M. Thermodynamic analysis showed that the free energies of A $\beta$ <sub>1-40</sub> monomer and A $\beta$ <sub>1-42</sub> trimertetramer are ~1.1 and ~15/~22 kcal/M, respectively. The early aggregation stages of A $\beta$ <sub>1-40</sub> and A $\beta$ <sub>1-42</sub> contain different solvent-exposed hydrophobic surfaces that are

located at the sequences flanking its protease-resistant segment. It was concluded that the amyloidogenic folded structure of A $\beta$  is important for the formation of spherical  $\beta$  oligomeric species.

**[0006]** Ono and colleagues (Ono, et al. (2009) *Proc. Natl. Acad. Sci.* 106, 14745-14750) clarified the ill-defined term "oligomere" because many different ones have been reported and often existing in rapid equilibrium with monomers and higher-order assemblies. They reported results of studies in which specific oligomers have been stabilized structurally, fractionated in pure form, and then studied using different biochemical, microscopic and neurotoxicity assays. A $\beta$ <sub>1-42</sub> monomers were largely unstructured, but oligomers exhibited order-dependent increases in  $\beta$ -sheet content. Dimerization and subsequent monomer addition are processes in which significant and asymmetric monomer conformational changes occur. Oligomer secondary structure and order correlated directly with fibril nucleation activity. Neurotoxic activity increased disproportionately (order dependence >1) with oligomer order. Their results provided significant insights into the biophysical and pathobiological behavior of A $\beta$ , and importantly, into strategies for developing therapeutics for AD. The authors concluded that the "specific activity" of A $\beta$ <sub>1-42</sub> assemblies depends nonlinearly on oligomer order. In fact, A $\beta$ <sub>1-42</sub> dimers were reported to be 3-fold more toxic than monomers, and tetramers were 13-fold more toxic. The mechanism of toxicity is not well understood. Membrane permeabilization by amyloid oligomers may initiate a common group of downstream pathologic processes, including intracellular calcium dyshomeostasis, production of reactive oxygen species, altered signaling pathways, and mitochondrial dysfunction that represent key effectors of cellular dysfunction and cell death. Naturally secreted A $\beta$  oligomers may directly impair synaptic function as has been shown to block hippocampal long-term potentiation (reviewed in Wirths et al. (2004) *J Neurochem* 91, 513-520).

**[0007]** Moreover, in vitro and in vivo analysis of amyloid deposits in AD revealed various N- and C-terminal variants. Increased C-terminal length of A $\beta$  (from A $\beta$ <sub>x-40</sub> to A $\beta$ <sub>x-42</sub>) in AD enhanced aggregation, early deposition and promoted the toxicity of A $\beta$ . Beside A $\beta$  peptides, starting with aspartate as the first amino acid (A $\beta$ <sub>1</sub>), several N-truncated and modified A $\beta$  species have been described. Among A $\beta$  species present in AD plaques, it was reported that A $\beta$ <sub>4-42</sub> is a relatively abundant species in AD, aged controls and vascular dementia patients. Using immunoprecipitation in combination with mass spectrometry, Portelius and colleagues showed that A $\beta$ <sub>1-40</sub>, A $\beta$ <sub>1-42</sub>pyroglutamate A $\beta$ <sub>pE3-42</sub> and A $\beta$ <sub>4-42</sub> are the dominant fractions in the hippocampus and cortex of AD patients. Interestingly, it has been demonstrated that N-terminal deletions enhance A $\beta$  aggregation comparing A $\beta$ <sub>1-42</sub> with A $\beta$ <sub>1-42</sub>.

**[0008]** Intra- or extracellular deposition of proteins is a feature of neurodegenerative diseases and serves as a molecular pathologic basis for classification as proteinopathies. While immunore-activity for A $\beta$  or prion protein (PrP) is located predominantly extracellularly, proteins that deposit intracellularly include tau,  $\alpha$ -synuclein, or TAR DNA Binding Protein 43 (TDP-43). Tauopathies include progressive supranuclear palsy (PSP), corticobasal degeneration (CBD), and argyrophilic grain disease (AGD), all predominated by the 4R isoform of the tau protein, and Pick's disease (PiD), a 3R tauopathy. Alpha-synucleinopathies comprise Parkinson's disease (PD), dementia with Lewy bodies (DLB), and



multiple system atrophy (MSA). TDP-43 proteinopathies include a group of frontotemporal lobar degenerations (FTLD-TDP), as well as sporadic amyotrophic lateral sclerosis (ALS) cases, while the most frequent form of prion disease is sporadic Creutzfeldt-Jakob disease (CJD). Kayed et al. have reported that different types of soluble amyloid oligomers display a common conformation-dependent structure that is unique to soluble oligomers regardless of sequence suggesting that they share a common mechanism of toxicity. These conformational antibodies against A $\beta$  also recognized non-A $\beta$  oligomers (Kayed et al. (2003) *Science* 300, 486-489).

**[0009]** One of the major problems in the amyloid hypothesis is the weak correlation between the severity of dementia and the density and localization of amyloid plaques in the brain of AD patients. Memory impairment and pathological changes in many AD mouse models occur well before the onset of plaque deposition. Soluble oligomers are low molecular weight non-fibrillar structures, which are stable in aqueous solution and remain soluble even after high speed centrifugation. A $\beta$  oligomers develop preferentially intracellularly within neuronal processes and synapses rather than within the extracellular space. On the other side, A $\beta$  can be taken up by certain cells, and then concentrated into endosomes/lysosomes. At high concentrations, vesicular A $\beta$  aggregates form high molecular weight oligomers which are capable of seeding amyloid fibril growth. Results from several labs propose these oligomers to be the missing link in the amyloid hypothesis. While AR plaques are poor correlates for the clinical symptomatology in AD and Down syndrome patients, soluble oligomers are suggested to be good predictors for synaptic loss, neurofibrillary tangles and clinical phenotype. Just like in the human brain, studies using AD mouse models support the role of oligomers. In the Tg2576 mouse model, the appearance of A $\beta$  dodecamers coincided with the onset of spatial memory impairment. Interestingly, injection of these purified oligomers into the ventricle of wildtype rats caused a dramatic drop in spatial memory performance. With regard to short-term effects, oligomers have been shown to impair synaptic plasticity by blocking LTP and reinforcing LTD. Others generated APP transgenic mice expressing the E693 $\Delta$  mutation, which causes neuronal cell death and cognitive impairment by enhanced intracellular A $\beta$  oligomerization without plaque formation.

**[0010]** Therefore, oligomers are thought to be a good target for therapeutic antibodies especially that this pool represents a minor subset (around 1.4% of total A $\beta$ ) if compared to the plaque pool which occupies the whole parenchyma. Reports have shown that monoclonal antibodies raised against oligomers prevent oligomer-induced toxicity, production of reactive oxygen species, and their attachment to synapses in primary hippocampal cells. In addition, Klyubin et al. have shown that immunization against A $\beta$  oligomers neutralized LTP inhibition induced by A $\beta$  oligomers injected in the hippocampus of rats (Klyubin et al. (2005) *Nat Med* 11, 556-561). Cell surface receptor clustering near or within synapses might be a mechanism for oligomer toxicity in AD. A $\beta$  oligomers can exert their toxicity through binding at synapses. It has been demonstrated that oligomers undergo a progressive shift from an extrasynaptic, freely diffusive state toward the formation of static synaptic clusters acting as an extracellular scaffold for mGluR5.

**[0011]** A $\beta$  dodecamers have been described using synthetic A $\beta$  oligomers. Dimeric A $\beta$  was purified from AD brains and

also found to inhibit LTP in hippocampal slices and induce Tau hyperphosphorylation and other degenerative effects in cultured neurons. Selkoe and others reported that toxic A $\beta$  oligomers to be primarily dimers and trimers of A $\beta$  (Walsh et al. (2000) *Biochemistry* 39, 10831-10839; Walsh et al. (2002) *Nature* 416, 535-539; Klyubin, et al. (2008) *J. Neurosci.* 28, 4231-4237).

**[0012]** Reviewed in detail by Hampel et al. (Hampel et al. (2010) *Nat Rev Drug Discov* 9, 560-574) discussed that AD biomarkers are needed to monitor drug safety, to identify individuals who are most likely to respond to specific treatments, to stratify presymptomatic patients and to quantify the benefits of treatments. Cerebrospinal fluid markers (for example, increased phosphophorylated tau and decreased A $\beta$ <sub>1-42</sub> levels) are helpful as trait markers of AD that have high sensitivity and specificity. However, they explained that these biomarkers have yet to show value as markers of disease state. Of interest, it has been reported that levels of A $\beta$ <sub>1-42</sub> are already fully decreased at least 5 to 10 years before conversion to AD dementia, whereas T-tau and P-tau seem to be later markers. Additionally, A $\beta$  oligomers have also been studied as a potentially new biomarker in CSF. Analysis of HMW A $\beta$  oligomers derived from synthetic A $\beta$ <sub>1-42</sub> by size-exclusion chromatography, revealed that using an ELISA specifically detected HMW A $\beta$  oligomers of 40-200 kDa (Fukumoto et al. (2010) *Faseb J* 24, 2716-2726). The CSF levels of these HMW A $\beta$  oligomers showed a negative correlation with Mini-Mental State Examination scores in AD and mild cognitive patients, which are presymptomatic for AD.

**[0013]** Analysis of amyloid deposits in AD brains revealed various N- and C-terminal variants. The increased C-terminal length of A $\beta$  (from A $\beta$ <sub>x-40</sub> to A $\beta$ <sub>x-42</sub>) enhances its aggregation properties. Faster aggregation leads to earlier A $\beta$  deposition, which is believed to promote its toxicity. Recently, A $\beta$ <sub>1-43</sub> was discovered as a novel toxic peptide in AD. Besides A $\beta$  peptides starting with aspartate as the first amino acid (A $\beta$ 1), several N-truncated and modified A $\beta$  species have also been described. It was reported that A $\beta$ <sub>4-42</sub> is a relatively abundant species in AD, aged controls and vascular dementia patients. Using immunoprecipitation in combination with mass spectrometry, it was shown that A $\beta$ <sub>1-40</sub>, A $\beta$ <sub>1-42</sub>, A $\beta$ <sub>pE3-42</sub> and A $\beta$ <sub>4-42</sub> are the dominant fractions in the hippocampus and cortex of AD patients. Moreover, it has been demonstrated that N-terminal deletions enhance A $\beta$  aggregation comparing A $\beta$ <sub>4-42</sub> with A $\beta$ <sub>1-42</sub>. It was further demonstrated that A $\beta$ <sub>1-42</sub> and A $\beta$ <sub>pE3-42</sub> exhibited similar effects on neuronal cytotoxicity in primary cortical neurons and on memory impairment after intracerebroventricular injection in wildtype mice. A $\beta$ <sub>pE3-42</sub> is now an established factor contributing AD pathology and may even be aggravating the severity of the disease (Wittnam et al. (2012) *Journal of Biological Chemistry* 287, 8154-8162). While the possible role of A $\beta$ <sub>4-42</sub> although historically discovered first and found to be a major peptide in AD brain, its role as a biomarker or therapeutic target is not well explored.

**[0014]** The amyloid hypothesis in Alzheimer's disease (AD) considers amyloid  $\beta$  peptide (A $\beta$ ) deposition as the causative event triggering down-stream events like neurofibrillary tangles, cell loss, vascular damage and memory decline. A major complication of the amyloid hypothesis has been that so far none of the antibodies therapies directed against plaque-A $\beta$  has revealed the expected outcome in clinical trials. Although plaque load has been successfully reduced, the cognitive decline could not be influenced. More-

over, in everyday clinical practice,  $A\beta_{1-42}$  ELISAs are used that do not discriminate between the different aggregation states (monomers up to high molecular weight oligomers (HMW)) and are lowered during the development of AD.

**[0015]** In view of the above, it is desirable to target all  $A\beta_{x-42}$  low and high molecular weight oligomers specifically. The expected therapeutic benefit is to neutralize these oligomers by passive immunization as has been demonstrated recently for another oligomer-specific antibody 9D5 detecting only low molecular weight (LMW)  $A\beta_{pE3-42}$  (Wirhns et al. (2010) *J. Biol. Chem.* 285, 41517-41524; and WO 2011151076). Other antibodies specific for  $A\beta$ -peptides and certain oligomers are described in WO 2009056490, U.S. Pat. No. 7,763,249, EP 2 210 901, and EP 2 246 427.

**[0016]** WO 2009065054 discloses antibodies specific for the protofibril form of  $A\beta$ -peptides (~670 kDa). However, these antibodies differ from the presently described ones, since the antibodies of WO 2009065054 do not bind to trimeric or tetrameric human  $A\beta$ , i.e. low molecular weight forms of  $A\beta$ -oligomers (cf. FIGS. 1, 2A, and 3A of WO 2009065054). Wang et al. (Wang et al. (2009) *FEBS Letters* 583, 579-584) disclose scFv's W8, W9, W20, and WC2, which bind to  $A\beta$  trimers, tetramers and little bigger oligomers. However, the exact epitope, and/or the sequence of the variable regions of these antibody molecules are not described.

**[0017]** Kim et al. (Kim et al. (2004) *Neurobiology of Aging* 25, S145) describe the synthesis of conformationally-defined  $A\beta$  haptens, which may be useful for a future selection of antibodies that can selectively sequester these folding intermediates.

**[0018]** Presently, it is an object of the invention to provide novel antibodies, which can be advantageously used in the treatment and diagnosis of AD.

#### SUMMARY OF THE INVENTION

**[0019]** Together with full-length  $A\beta_{1-42}$ , N-truncated pyroglutamate  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  are major variants in AD brain. The inventor has identified antibodies that, due to its common epitope, react not only with HMW  $A\beta_{1-42}$ , but also with its major N-truncated variants  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  opening the potential to develop more sensitive biomarkers for diagnosing AD and monitoring drug efficacy in clinical trials. The present invention demonstrates the first successful attempt to identify an epitope that differentiates  $A\beta_{1-42}$  tri- or tetramers from  $A\beta_{1-42}$  mono- and dimers. It is known for several years that  $A\beta_{1-42}$  tri- or tetramers are the most toxic  $A\beta$  peptides at the beginning of the pathology of AD.

**[0020]** In particular, two novel conformational specific  $A\beta$  antibodies NT4X-83 and NT4X-167 preferentially reacting with N-truncated  $A\beta_{4-42}$  were developed having a unique binding pattern: NT4X recognized low and high molecular weight oligomers of  $A\beta_{1-42}$ ,  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$ , with the highest preferences for  $A\beta_{4-42}$ . NT4X detected only a minor portion of plaques in sporadic and familial AD cases and AD mouse models and did not cross-react with other aggregates typical for other common neurodegenerative diseases showing that NT4X staining is specific for AD. Importantly, NT4X inhibited aggregation and toxicity of  $A\beta_{1-42}$ ,  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$  implicating their potential for AD therapy.

**[0021]** Having two specific antibodies developed, it is now possible to target these oligomers specifically. The expected therapeutic benefit is to neutralize the toxic  $A\beta$ -oligomers by passive immunization for example.

**[0022]** Thus, in summary,

**[0023]** The antibodies of the invention recognize a conformational epitope formed by the amino acids in position four and five of  $A\beta$ , in particular phenylalanine at position four of  $A\beta$ .

**[0024]** The antibodies according to the invention differentiate  $A\beta_{1-42}$  LMW oligomers from  $A\beta_{1-42}$  mono- and dimers.

**[0025]** The antibodies according to the invention bind native LMW oligomers derived from  $A\beta_{pE3-42}$  and most efficiently  $A\beta_{4-42}$ .

**[0026]** Aged  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  develop HMW oligomers detected by the antibodies according to the invention.  $A\beta_{3-38}$  and  $A\beta_{4-40}$  did not aggregate into HMW oligomers.

**[0027]** The antibodies according to the invention rescued toxic effects of all  $A\beta$  peptides tested using cell viability and aggregation assays in vitro and detected only a minor fraction of plaques in brain from sporadic and familial AD patients and two transgenic mouse models of AD. As the amount of detected plaques by the antibodies according to the invention is minor, an adverse treatment effect by passive immunization by dissolving plaques is not likely and side effects are not expected. The antibodies according to the invention are therefore expected to be highly valuable in the treatment of AD.

**[0028]** The antibodies according to the invention specifically reacted with blood vessel walls and with intraneuronal  $A\beta$ . Hence, two assumptions can be made: 1. These oligomers are soluble. 2. These oligomers likely can be found in sufficient amounts in blood making a blood test for AD possible. Further, they did not cross-react with aggregates typical for other major neurodegenerative disorders implicating that the recognized aggregates are disease-specific for AD. Therefore, the antibodies according to the invention are highly valuable for diagnostic purposes.

**[0029]** Accordingly, in a first aspect, the invention relates to a monoclonal antibody molecule recognizing a conformational epitope formed by Phe at amino acid position 4 of human  $A\beta$  in trimeric or tetrameric  $A\beta$  oligomers, wherein said antibody molecule does not bind human monomeric  $A\beta_{1-42}$  or human  $A\beta_{1-42}$  dimers, as determined by native Western-blotting using 7  $\mu\text{g}$   $A\beta_{1-42}$  on a 18% SDS-free polyacrylamide gel and 0.45 nitrocellulose membranes and said antibody molecule as the primary antibody in a concentration of not to more than 10  $\mu\text{g}/\text{ml}$ .

**[0030]** The antibody molecule is particularly useful for use in medicine or veterinary medicine, in particular for use in the treatment and/or prevention of Alzheimer's disease, in particular wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease. Accordingly, the invention also relates to the use of an antibody molecule according to the first aspect in the manufacture of a medication.

**[0031]** In a related aspect, the invention further provides a method of treating or preventing Alzheimer's disease in a subject, comprising the step of administering an antibody molecule according to the first aspect to said subject.

**[0032]** In still another aspect, the invention relates to the use of the antibody molecule according to the first aspect in a method of diagnosing Alzheimer's disease, or in a method of identifying agents useful in the treatment and/or prevention of

Alzheimer's disease, in particular wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease.

**[0033]** Thus, in another aspect, the invention relates to a method of diagnosing Alzheimer's disease, comprising the step of determining the amount of low molecular weight oligomers of A $\beta$  in a sample of a subject to be diagnosed, using an antibody molecule according to the invention. In this context, the invention also pertains to a diagnostic composition comprising an antibody molecule according to the invention.

**[0034]** The invention further relates to the use of an antibody molecule according to the first aspect in a method of monitoring the efficacy of a treatment of Alzheimer's disease, in particular wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease. Accordingly, the invention also relates to a method of monitoring the efficacy of a treatment of Alzheimer's disease, comprising the step of determining the amount of low molecular weight oligomers of A $\beta$  in a sample of a subject to be diagnosed using an antibody molecule according to the invention.

**[0035]** In still another aspect, the invention also pertains to a hybridoma as deposited under DSM ACC3161 or hybridoma as deposited under DSM ACC3162, which produces an antibody according to the first aspect.

**[0036]** Likewise, the invention further provides a nucleic acid molecule encoding the antibody molecule according to the first aspect, a vector comprising said nucleic acid molecule, and a host cell comprising said nucleic acid molecule or said vector.

**[0037]** In a final aspect, the invention also relates to a method of producing an antibody molecule according to the first aspect comprising the step of (i) culturing the host cell or the hybridoma of the invention under conditions allowing synthesis of said antibody molecule and (ii) recovering said antibody molecule from said culture.

#### DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

**[0038]** In a first aspect, the invention relates to a monoclonal antibody molecule recognizing a conformational epitope formed by Phe at amino acid position 4 of trimeric or tetrameric human A $\beta$ , as determined by pepscan ELISA comprising the steps of precoating with 100 ng A $\beta$ <sub>4-19</sub> and A $\beta$ <sub>5-20</sub>, blocking, incubating with 7  $\mu$ g/ml of said antibody molecule, and detecting said antibody molecule with a labeled secondary antibody; wherein said antibody molecule does not bind human monomeric A $\beta$ <sub>1-42</sub> or human A $\beta$ <sub>1-42</sub> dimers, as determined by native

**[0039]** Western-blotting comprising the steps of applying 7  $\mu$ g A $\beta$ <sub>1-42</sub> on a 18% SDS-free polyacrylamide gel, transferring on a 0.45  $\mu$ m nitrocellulose membrane, blocking, incubating with said antibody molecule in a concentration of not more than 10  $\mu$ g/ml for 2 h at room temperature, and detecting said antibody molecule with a labeled secondary antibody.

**[0040]** The term "monoclonal antibody molecule" as used herein refers to an antibody molecule derived from a population of substantially homogeneous antibodies, i.e. the individual antibody molecules comprised in the population are identical except for possible naturally occurring mutations that may be present in minor amounts. Typically, monoclonal antibody molecules are highly specific and are directed against a single epitope. In addition to their specificity, the monoclonal antibody molecules may be advantageously syn-

thesized uncontaminated by other antibodies. The modifier "monoclonal" is not to be construed as requiring production of the antibody molecule by any particular method. However, a monoclonal antibody molecule as disclosed herein may be produced by the hybridoma method first described by Kohler, G. et al., Nature 256 (1975) 495, or by recombinant DNA methods (see, e.g., U.S. Pat. No. 4,816,567).

**[0041]** An example of the antibody molecule of the first aspect is an antibody produced by hybridoma as deposited under DSM ACC3161 or hybridoma as deposited under DSM ACC3162. Likewise, the antibody molecule may be an antibody molecule which competes for the same epitope with antibody NT4X-83, as deposited under DSM ACC3161, and/or antibody NT4X-167, as deposited under DSM ACC3162. Assays for determining competitive binding of two antibodies are known in the art. For example, a competitive assay may be based on the pepscan ELISA assay, as described below, wherein the human A $\beta$  1-42 is preincubated with NT4X-83 or NT4X-167, and subsequent binding of the competing antibody molecule is detected using a detection antibody, which is specific for said competing antibody molecule. Alternatively, the human A $\beta$  1-42 is preincubated with the competing antibody molecule, and subsequent binding of NT4X-83 or NT4X-167 is detected using a detection antibody, which is specific for NT4X-83 or NT4X-167.

**[0042]** The antibody molecule may be a recombinant full antibody (immunoglobulin), a F(ab)-fragment, a F(ab)<sub>2</sub>-fragment, a F(v)-fragment, a single-chain antibody, a chimeric antibody, a CDR-grafted antibody, a bivalent antibody-construct, a polyvalent antibody-construct, a synthetic antibody, a cross-cloned antibody, a fully-human antibody, a humanized antibody, nanobodies, and polyvalent nanobodies or diabodies.

**[0043]** Antibodies are also called immunoglobulins. They are gamma globulin proteins and generally consist in their natural form of two heavy chains and two light chains linked by disulfide bonds. Five types of mammalian Ig heavy chains are known:  $\alpha$ ,  $\delta$ ,  $\epsilon$ ,  $\gamma$ , and  $\mu$ , wherein the type of heavy chain defines the class (isotype) of the antibody, which are IgA, IgD, IgE, IgG, and IgM. The heavy chain contains two regions, the constant region and the variable region. The constant region shares high homology in all naturally occurring antibodies of the same isotype within the same species. Like the heavy chain, a light chain also consists of one constant domain and one variable domain. In mammals there are two types of immunoglobulin light chain, lambda ( $\lambda$ ) and kappa ( $\kappa$ ). The unique property or specificity of a given antibody is determined by the variable (V) regions. In particular, three variable loops in each the light (V<sub>L</sub>) and the heavy (V<sub>H</sub>) chain, are responsible for the antigen specificity.

**[0044]** The term "full antibody" is intended to refer to any antibody that has a typical overall domain structure of a naturally occurring antibody (i.e. comprising a heavy chain of three or four constant domains and a light chain of one constant domain as well as the respective variable domains). Nevertheless, each domain may comprise further modifications, such as mutations, deletions, or insertions, which do not change the overall domain structure. The antibody may be an IgA, IgD, IgE, IgG, or IgM antibody, including any subclass of these isotypes. In one preferred embodiment, the antibody is an IgG antibody, such as an IgG1 or IgG2 antibody. If recombinantly produced, the antibody may also comprise two different constant regions of heavy chains, e.g. one IgG1

and one IgG2 heavy chain, or heavy chains from different species. The heavy chains are, however, preferably from the same species.

**[0045]** A fragment of an antibody molecule contains at least one antigen binding fragment as defined above, and exhibits the same specificity as the complete antibody of which the fragment is derived from, e.g. an antibody fragment of NT4X-83 or NT4X-167 has the same specificity than the NT4X-83 or NT4X-167 antibody. Methods of producing antibody molecule fragments are known in the art. For example, Fab fragments may be generated from an immunoglobulin by using the enzyme papain. Likewise, F(ab)<sub>2</sub> fragments are formed by using the enzyme pepsin, which cleaves below the hinge region and, thus, below the disulfide bonds. Alternatively, the variable regions of the heavy and light chains can be fused together to form a single chain variable fragment (scFv). Thus, in the context of the this invention, an antibody molecule does also comprise variable and light regions, F(ab)-, F(ab)<sub>2</sub> fragments, CDR-regions, etc. of the antibody molecule as disclosed herein. The antibody molecule may also be a humanized or CDR-grafted antibody molecule, as well as genetically/recombinantly engineered. An example of an engineered antibody molecule, e.g. produced by CDR-grafting, is an antibody, in which at least one region of an immunoglobulin of one species is fused to another region of an immunoglobulin of another species in order to reduce its immunogenicity. CDR-grafting is well-known in the art. Alternatively, the antibody molecule may be a fully-human antibody. One variable region of the antibody molecule of the invention may also be combined with another variable region, which binds to another epitope, thereby generating bivalent or polyvalent antibody constructs. Also provided are derivatives of such antibody molecules, like single-chain antibodies, diabodies, bispecific single chain antibodies, nanobodies and polyvalent nanobodies.

**[0046]** The antibody molecule recognizes a conformational epitope, which is predominantly formed by Phe at amino acid position 4 of human A $\beta$  of each of the A $\beta$  forming trimeric and/or tetrameric oligomers. Generally, a conformational epitope is a formation of subunits, here human A $\beta$ , composing an antigen that is recognized by an antibody molecule. "Predominantly" is thereby intended to mean that said Phe is a prerequisite for detection with high intensity of binding to human A $\beta$ , as also demonstrated in the examples section, where the antibody molecule showed binding to A $\beta$ <sub>4-19</sub> but not to A $\beta$ <sub>5-20</sub>. However, in a preferred embodiment, the epitope further comprises the amino acids at position 2 and 3 of human A $\beta$ , as determined by pepscan ELISA. Pepscan is a procedure for mapping and characterizing epitopes involving the synthesis of overlapping peptides and analysis of the peptides in enzyme-linked immunosorbent assay (ELISAs).

**[0047]** A $\beta$  peptides with 16 amino acids in length may be used: A $\beta$ <sub>1-16</sub>, 2-17, 3-18, 4-19, 5-20, 6-21, 7-22, 8-23, 9-24 and 10-25. The experiment is performed using U form high binding 96 well-plate (microplates, Greiner bio-one) to coat the peptides. Samples are diluted in ice cold sterile-filtered 0.05 M carbonate buffer (2.4 g NaCO<sub>3</sub>×10H<sub>2</sub>O, 1.46 g NaHCO<sub>3</sub> added to 450 ml ddH<sub>2</sub>O, pH 9.6) and 50  $\mu$ l of the diluted samples are coated on the wells (the well-plate is kept on ice for some min before peptide application). For the peptide binding, the well-plate is incubated overnight at 4° C. The next day, the wells are washed three times by 0.01 M PBS (RT). The washing is done by applying 250  $\mu$ l PBS into each well and emptying it by inverting the plate and hit it hard on a spongy cloth. After

the washing step, three other steps are conducted (each step for 1 h at 25° C.): blocking (preventing unspecific antibody binding); incubation with primary antibody, i.e. the antibody molecule to be tested, at a concentration of 7  $\mu$ g/ml; incubation with the secondary antibody (e.g., HRP coupled goat anti-mouse IgG, Dianova, 1:10.000, cat no: 115-035-003; or a labeled secondary antibody recognizing the  $\kappa$  or  $\lambda$  light chain). The skilled person will know how to choose a suitable secondary antibody. After each mentioned step, three times washing with 0.01 M PBS is applied. For the blocking step, 200  $\mu$ l of the RT blocking solution containing 5% w/v Albumin Fraction V (Roth) in 0.01 M PBS, which is prepared, sterile filtered and kept at 4° C. a day before use. Both, primary and secondary antibodies, are diluted in the blocking solution and 100  $\mu$ l of each dilution is used for each well. Lastly, as a HRP enzyme substrate, 50  $\mu$ l of 3,3',5,5'-Tetramethylbenzidin (1-step Ultra TMB ELISA substrate, Thermo Scientific), that leads to the blue color production when oxidized, is added to each of the wells and the absorbance (OD) is measured within 5-10 min at 652 nm by a microplate reader (pQuant, B Tek Instrument INC) using Mikro Win 2000 software (Berthold Technologies). The reaction is stopped by adding 50  $\mu$ l of 2 M, H<sub>3</sub>PO<sub>4</sub> which leads to yellow color formation and the absorbance is measured at 450 nm.

**[0048]** The affinity constant for binding of an antibody molecule to the epitope formed by the trimeric or tetrameric A $\beta$  oligomers can span a wide range, extending from about 10<sup>5</sup> mol<sup>-1</sup> to about 10<sup>12</sup> mol<sup>-1</sup>, and the affinity constant is preferably at least 10<sup>6</sup> mol<sup>-1</sup>, more preferably at least 10<sup>7</sup> mol<sup>-1</sup>, even more preferably at least 10<sup>8</sup> mol<sup>-1</sup>, most preferably at least 10<sup>9</sup> mol<sup>-1</sup>, and even most preferably at least 10<sup>10</sup> mol<sup>-1</sup>, such as at least 10<sup>11</sup> mol<sup>-1</sup>. Procedures for determining an affinity constant of an antibody molecule are known in the art.

**[0049]** The antibody molecule recognizes a conformational epitope formed in trimeric or tetrameric A $\beta$  oligomers. The A $\beta$  oligomers may be composed of a single type of A $\beta$ , or of a mixture of one, two, three, or more types of A $\beta$ . In a preferred embodiment, the trimeric or tetrameric A $\beta$  oligomers are oligomers of A $\beta$  1-X, A $\beta$  4-X, and/or pyro-Glu-A $\beta$  3-X, wherein X is 42, 40 or 38, preferably wherein X is 42. In one preferred embodiment, the A $\beta$  oligomers are oligomers of A $\beta$  1-42. In another preferred embodiment, the A $\beta$  oligomers are oligomers of A $\beta$  1-40. In another preferred embodiment, the A $\beta$  oligomers are oligomers of A $\beta$  1-38. In another preferred embodiment, the A $\beta$  oligomers are oligomers of A $\beta$  4-42. In another preferred embodiment, the A $\beta$  oligomers are oligomers of A $\beta$  4-40. In another preferred embodiment, the A $\beta$  oligomers are oligomers of A $\beta$  4-38. In another preferred embodiment, the A $\beta$  oligomers are oligomers of pyro-Glu-A $\beta$  3-42. In another preferred embodiment, the A $\beta$  oligomers are oligomers of pyro-Glu-A $\beta$  3-40. In another preferred embodiment, the A $\beta$  oligomers are oligomers of pyro-Glu-A $\beta$  3-38.

**[0050]** As discussed in the background of the invention section, Ono, et al. reported results of studies in which specific oligomers have been stabilized structurally, fractionated in pure form, and then studied using different biochemical, microscopic and neurotoxicity assays. It could be demonstrated that A $\beta$ <sub>1-42</sub> monomers were largely unstructured, but oligomers exhibited order-dependent increases in  $\beta$ -sheet content. Dimerization and subsequent monomer addition are processes in which significant and asymmetric monomer conformational changes occur, thereby generating an epitope

which is specifically bound by the antibody molecule of the invention under native conditions. Ono et al. reported that  $A\beta_{1-42}$  dimers were 3-fold more toxic than monomers, and tetramers were even 13-fold more toxic. In fact, it is known for several years that  $A\beta_{1-42}$  tri- or tetramers are the most toxic  $A\beta$  peptides at the beginning of the pathology of AD. Accordingly, the amount of trimers and tetramers of  $A\beta$  in a sample are considered being predictive of disease state and disease progression. The antibody molecule according to the invention is capable of differentiating between  $A\beta_{1-42}$  monomers and dimers on the one hand, and  $A\beta_{1-42}$  trimers and tetramers on the other hand. Accordingly, due to its specificity for the specific epitope identified herein, the antibody is not adsorbed to the less toxic monomers and dimers, and even shows very low binding to plaques. Due to these surprising characteristics, the antibody molecule according to the invention is particularly useful in diagnosis, treatment and prophylaxis of diseases associated with human  $A\beta$ , in particular  $A\beta_{1-42}$ , such as Alzheimer's disease.

**[0051]** In order to determine whether an antibody molecule binds  $A\beta_{1-42}$  tri- or tetramers but not the respective monomers and dimers, one may conduct native Western-blotting comprising the steps of applying 7  $\mu\text{g}$   $A\beta_{1-42}$  on a 18% SDS-free polyacrylamide gel, transferring on a 0.45  $\mu\text{m}$  nitrocellulose membrane, blocking, incubating with said antibody molecule in a concentration of not more than 10  $\mu\text{g}/\text{ml}$ , preferably 5-10  $\mu\text{g}/\text{ml}$ , such as 6-9  $\mu\text{g}/\text{ml}$ , e.g. 7 or 8  $\mu\text{g}/\text{ml}$ , for 2 h at room temperature, and detecting said antibody molecule with a labeled secondary antibody. The secondary antibody may be any suitable secondary antibody, which is capable of recognizing the antibody molecule of the invention, e.g. by binding to its Fc-portion or to the light chain, or to another polypeptide, which was introduced into the antibody molecule by way of a fusion polypeptide in order to provide a specific antigenic epitope. The secondary antibody is usually labeled with a marker, dye or enzyme, which allows a quantitative detection of a signal which correlates with the binding of the secondary antibody to the antibody molecule, and thus with the binding of the antibody molecule to the conformational epitope.

**[0052]** More specifically, the peptides are mixed with the dye (Tris-Glycin Native Probenpuffer, Anamed) in the ration of 1:1 for 5 min at RT. Then, the sample is loaded on 18% SDS-free gels (Progel Tris Glycin 18%, 1.0 mm, Anamed). Running the proteins on the gel is done using the SDS-free running buffer (Tris-Glycin Native Laufpuffer Anamed). The proteins are then transferred on 0.45  $\mu\text{m}$  nitrocellulose membrane (GE Healthcare), using the semi-dry method (BRAD, Trans-Blot® SD Semi-Dry): The membrane, gel and two thick blotting papers are wetted into the transfer buffer (48 mM Tris-HCL buffer, 39 mM glycine (5% methanol) pH 9.2). Wet thick papers are placed around the membrane and the gel (the membrane was lying under the gel) and the protein transfer is performed at 25 V for 1 h (Trans-Blot®SD semi-dry Transfer cell, BIO-RAD). Upon completion of protein transfer, the membrane is washed two times each for 10 min in 1 $\times$  TBS/T (0.1 M Tris, 1.5 M NaCl and 0.5% Tween 20) on a rotator (Heidolph, DOUMAX 1030) with speed of 40 rpm. For  $A\beta$  peptides, membranes are boiled in 0.1 M PBS for about 1 min in microwave (ShARP) on 800 Watt power. After the start of boiling, the power is decreased to 80 Watt and the boiling is continued for 4 more minutes. Membranes are briefly washed in 1 $\times$  TBS/T and blocked with 10% non-fat dry milk (Roth) in 1 $\times$  TBS/T for 1 h at room temperature (RT) (40 rpm shaking). The membrane is incubated with following

primary antibodies overnight at 4° C. or 2 h at RT (40 rpm shaking). This step is followed by two times of washing with 1 $\times$  TBS/T each for 10 min (40 rpm shaking) and the membrane is incubated with secondary antibody (e.g. Polyclonal goat anti-mouse, Dianova diluted 1:4000) conjugated with horseradish peroxidase, for 2 h at RT. The skilled person will know how to choose an appropriate secondary antibody. After washing for two times, each for 10 min, for development, the membrane is incubated in the horseradish peroxidase substrate solution containing 8 ml solution A (0.25 mg/ml luminol (Roth) in 0.1 M Tris/HCl), 800  $\mu\text{l}$  solution B (1.1 mg/ml para-coumaric (Roth) acid in DMSO (Sigma) and 2.5  $\mu\text{l}$   $\text{H}_2\text{O}_2$  (Roth) for 2 min. After slightly drying the membrane on a tissue paper, it is exposed to x-ray films (Hyperfilm EC, Amersham Biosciences) for different time points and developed in a CUR1x60 (AGFA). Further guidance can be found in the examples section.

**[0053]** The inventor observed a striking difference between the binding of NT4X with  $A\beta$  peptides under reducing and native conditions. While under reducing conditions, LMW oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  were detected, a strong signal was seen for monomers of  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$ , but not  $A\beta_{1-42}$ . Under reducing conditions,  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  formed stable LMW oligomers after dissolving the peptides. Aging of the peptides resulted in a shift of the SDS-stable signal  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  to HMW oligomers. No difference was observed between under reducing and native conditions for  $A\beta_{4-38}$  and  $A\beta_{4-40}$ , which apparently only formed monomers and dimers. Accordingly, as shown in the examples, the antibody molecule in accordance with the invention may not only bind oligomers of low order, but also "aged  $A\beta$  polypeptides", i.e. HMW oligomers of  $A\beta$  1-42, pyro-Glu- $A\beta$ 3-42 and/or  $A\beta$ 4-42, as determined by native Western-blotting using 18% SDS-free polyacrylamide gel and 0.45 nitrocellulose membranes and said antibody molecule as the primary antibody in a concentration of not more than 10  $\mu\text{g}/\text{ml}$ , preferably 5-10  $\mu\text{g}/\text{ml}$ , such as 6-9  $\mu\text{g}/\text{ml}$ , e.g. 7 or 8  $\mu\text{g}/\text{ml}$ . The native western blot assay may be conducted as described above.

**[0054]** As discussed above, the antibody molecule recognizes aggregates that are specific for Alzheimer's disease. For example, the antibody molecule recognizes aggregates in immunohistochemistry of brain tissue sections, in particular wherein the Alzheimer's disease is familial Alzheimer's disease. The immunohistochemistry may be carried out as follows: 4  $\mu\text{m}$  brain sections are obtained by cutting paraffin-embedded brain tissue using a microtome. Deparaffinization is performed by incubation in xylol (Carl Roth GmbH) two times each for 5 min and followed by series of decreasing ethanol (Chemie-vertrieb) concentration (100%, 95% and 70% for 10, 5 and 3 min respectively) to rehydrate the sections. This stage is followed by 1 min washing the sections with deionized  $\text{H}_2\text{O}$ . Endogenous peroxidase blocking is achieved by incubation of the sections for 30 min in the solution of 200 ml 0.01 M PBS and 2 ml 30%  $\text{H}_2\text{O}_2$ . Sections are washed by deionized  $\text{H}_2\text{O}$  for 1 min. The antigen retrieval is performed by boiling the sections in 10 M Citrate buffer (4.2 g Citric acid-Monohydrate added to 2000 ml  $\text{H}_2\text{O}$ ; pH=6) for 2 min in a microwave at 800 Watt power. After 2 min the power is decreased to 80 Watt and the boiling is continued for additional 8 min. After cooling down the sections for 15 min they are washed with deionized  $\text{H}_2\text{O}$  for 1 min followed by incubation in 0.01 M PBS+0.1% Triton for 15 min and for 1 min incubation in 0.01 M PBS as washing

step. Incubation with freshly prepared 88% formic acid for 3 min followed. Sections are washed twice by 0.01 M PBS for 1 and 5 min. Unspecific blocking is done by 1 h in 4% milk powder (Roth), 10% fetal cow serum (FCS) in 0.01 M PBS in wet a chamber at RT. Primary antibodies, diluted in 0.01 M PBS and 10% FCS, are applied on the sections and are incubated at RT overnight (in a wet chamber). On the next day, sections are washed three times, each for 5 min by 0.01 M PBS+0.1% Triton and for 1 min by 0.01 PBS. Then, secondary antibody (Rabbit anti-mouse, Dako, cat. no. E0354), conjugated with biotin is diluted 1:200 in 0.01 PBS and 10% FCS and is applied on the sections in the wet chamber and incubated for 1 h at 37° C. This step is followed by Avidin-Biotin complex (ABC) (Vectastain® Elite ABC Kit, Vector Laboratories; cat. No. PK6100) incubation at 37° C. for 1.5 h which is prepared and kept at 4° C. at least 30 min before its application. ABC solution is prepared as followed: 0.01 M PBS+10% FCS+1:100 solution A+1:100 solution B. Afterwards, the sections are washed three times with 0.01 PBS each time for 5 min, and the sections are exposed to diaminobenzidine (DAB) solution (5 ml 50 mM Tris/HCl PH 7.5+100 µl DAB stock solution (25 mg/ml 5-(4-Dimethylamino-Benzylidene) Rhodanine (sigma) in 0.05 M Tris/HCL, PH 7.4)+2.5 µl 30% H<sub>2</sub>O<sub>2</sub> (Roth) (added right before use) for a few minutes until staining was observed. The sections are washed with 0.01 PBS, three times each time for 5 min, and counterstaining is performed using filtered hematoxylin, for 40 s. The sections are dipped into deionized H<sub>2</sub>O and then washed in running tap water for 5 min. As the last step in staining, sections are dehydrated by series of increasing ethanol concentration (1 min 70%, 5 min 95% and 10 min 100%) and in the end two times in xylol, each for 5 min. The sections, then, are embedded using eukitt quick hardening mounting medium (Carl Roth GmbH (Roti®-Histokit)). However, the antibody molecule may also be used in other types of assays, such as in a quantitative assay for determining the amount of trimeric and tetrameric oligomers of human Aβ in e.g. body fluids such as blood, serum or spinal fluid, for example by way of an ELISA.

**[0055]** In another preferred embodiment, the antibody molecule inhibits further aggregation of trimeric or tetrameric Aβ oligomers of Aβ 1-X, Aβ 4-X, and/or pyro-Glu-Aβ 3-X, wherein X is 42, 40 or 38, preferably wherein X is 42, when tested in a concentration of 10 µM in 50 mM sodium phosphate buffer, 50 mM NaCl, 20 µM thioflavin T and 0,01% sodium azide, pH 7.4, at 37° C. in a peltier adapter with stirring in a thioflavin T aggregation assay. In one preferred embodiment, the Aβ oligomers are oligomers of Aβ 1-42. In another preferred embodiment, the Aβ oligomers are oligomers of Aβ 1-40. In another preferred embodiment, the Aβ oligomers are oligomers of Aβ 1-38. In another preferred embodiment, the Aβ oligomers are oligomers of Aβ 4-42. In another preferred embodiment, the Aβ oligomers are oligomers of Aβ 4-40. In another preferred embodiment, the Aβ oligomers are oligomers of Aβ 4-38. In another preferred embodiment, the Aβ oligomers are oligomers of pyro-Glu-Aβ 3-42. In another preferred embodiment, the Aβ oligomers are oligomers of pyro-Glu-Aβ 3-40. In another preferred embodiment, the Aβ oligomers are oligomers of pyro-Glu-Aβ 3-38.

**[0056]** Aβ peptides are solubilized in 10 mM NaOH at a concentration of 1 mg/ml, sonicated for 5 min, frozen in liquid nitrogen, and stored at -80° C. until use. Aggregation of Aβ peptides is investigated online using ThT aggregation assay (Varian fluorescence spectrophotometer) is using an

excitation wavelength of 446 nm and emission wavelength of 482 nm. Samples contain 10 µM of Aβ, 50 mM sodium phosphate buffer (pH 7.4), 50 mM NaCl, 20 µM ThT and 0.01% sodium azide. The samples are incubated at 37° C. in a peltier adapter with stirring. Data points are recorded every 10 min during the assay and plotted in a diagram time [min] vs. fluorescence [a.u.]. If the antibody molecule is capable of inhibiting Aβ-oligomerization, the graph will reach a plateau phase, in which the oligomers are “stabilized” by the binding molecule, and do not further oligomerize and aggregate. In the context of the present invention, an antibody molecule is capable of inhibiting Aβ-oligomerization, if the resulting graph shows no inflection point after 250 min, preferably after 300 min, more preferably after 350 min, even more preferably after 400 min, such as after 450 min, and most preferably after 500 min incubation time.

**[0057]** An example of an antibody molecule in accordance with the present invention is the NT4X-83 and NT4X-167 antibody. Said antibodies are producible from and deposited as hybridomas. Thus, the invention also relates to a hybridoma as deposited under DSM ACC3161 or hybridoma as deposited under DSM ACC3162. Methods for producing the NT4X-83 and NT4X-167 antibody are also described below.

**[0058]** The sequences of the CDRs and the variable regions of the light and heavy chains of the NT4X-83 and NT4X-167 antibodies are provided below under the section sequences. Based on the provided sequences and on the antibodies as obtainable and deposited as a hybridoma under DSM ACC3161 and DSM ACC3162, the skilled person can readily design further antibody molecules having the desired properties described above. Accordingly, in a preferred embodiment, the antibody molecule comprises a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 1, 2 and 3, and a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 4, 5 and 6 (NT4X-83). In another preferred embodiment, the antibody molecule comprises a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 9, 10 and 11, and a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 12, 13 and 14 (NT4X-167). However, useful antibody molecules may also be obtained, which comprise a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 1, 2 and 3 (NT4X-83). In still another preferred embodiment, the antibody molecule comprises a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 9, 10 and 11 (NT4X-167). Likewise, the antibody molecule may comprise a variable region that comprises a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 4, 5 and 6 (NT4X-83). In still another preferred embodiment, the antibody molecule comprises a variable region that comprises a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 12, 13 and 14 (NT4X-167).

**[0059]** The antibody molecule may also be defined by reference to a sequence of a complete variable region. It is most preferred that such variable regions comprise the CDRs and/or combination of CDRs given above. Alternatively, a particular combination of CDRs set forth above differs not more than in ten amino acid residues, such as 9 amino acid residues, preferably 8 amino acid residues, such as 7 amino acid residues, more preferably 6 amino acid residues, such as 5 amino acid residues, even more preferably 4 amino acid residues, such as 3 amino acid residues, and most preferably in 2 amino

acid residues, such as 1 amino acid residue from the combination of sequences taken together.

**[0060]** Hence, in one preferred embodiment, the antibody molecule comprises a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 7, and comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 8 (NT4X-83).

**[0061]** In another preferred embodiment, the antibody molecule comprises a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 15, and comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 16 (NT4X-167).

**[0062]** In still another preferred embodiment, the antibody molecule comprises a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 7 (VH NT4X-83).

**[0063]** In another preferred embodiment, the antibody molecule comprises a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 15 (VH NT4X-167).

**[0064]** In another preferred embodiment, the antibody molecule comprises a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 8 (VL NT4X-83).

**[0065]** In another preferred embodiment, the antibody molecule comprises a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 16 (VL NT4X-167).

**[0066]** The person skilled in the art is readily in a position to verify the sequence identity of sequences, e.g. by simply comparing said sequences over the whole length of the sequence provided herein. For example, such an alignment can be performed using publicly available computer homology programs such as the "BLAST" program provided at the NCBI homepage at <http://www.ncbi.nlm.nih.gov/blast/blast.cgi>, using the default settings provided therein. Further methods of calculating sequence identity percentages of sets of amino acid sequences are known in the art.

**[0067]** Therefore, one aspect of the invention relates to an antibody molecule comprising

**[0068]** (a) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 1, 2 and 3, and a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 4, 5 and 6 (NT4X-83); or

**[0069]** (b) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 9, 10

and 11, and a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 12, 13 and 14 (NT4X-167); or

**[0070]** (c) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 1, 2 and 3 (NT4X-83); or

**[0071]** (d) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 9, 10 and 11 (NT4X-167); or

**[0072]** (e) a variable region that comprises a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 4, 5 and 6 (NT4X-83); or

**[0073]** (f) a variable region that comprises a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 12, 13 and 14 (NT4X-167).

**[0074]** In a preferred embodiment, said antibody molecule comprises

**[0075]** (a) a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 7, and comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 8 (NT4X-83); or

**[0076]** (b) a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 15, and comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 16 (NT4X-167); or

**[0077]** (c) a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 7 (VH NT4X-83); or

**[0078]** (d) a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 15 (VH NT4X-167); or

**[0079]** (e) a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 8 (VL NT4X-83); or

**[0080]** (f) a variable region comprising an amino acid sequence which is at least 80%, preferably 85%, more preferably 90%, even more preferably 95%, most preferably 97%, such as 100% identical to the amino acid sequence as shown in SEQ ID NO: 16 (VL NT4X-167).

**[0081]** Preferably, said antibody molecule is selected from a recombinant full antibody (immunoglobulin), a F(ab)-fragment, a F(ab)<sub>2</sub>-fragment, a F(v)-fragment, a single-chain antibody, a chimeric antibody, a CDR-grafted antibody, a bivalent antibody-construct, a polyvalent antibody-construct, a synthetic antibody, a cross-cloned antibody, a fully-human antibody, a humanized antibody, nanobodies, and diabodies, polyvalent nanobodies or diabodies.

**[0082]** In a related aspect, the present invention also relates to a nucleic acid molecule encoding the antibody molecule according to the invention. Preferably, the nucleic acid mol-

ecule comprises the antibody molecule encoding sequence in functional linkage with additional regulatory sequences, which direct the expression and translation of the coding sequence. Examples for regulatory sequences are a promoter, e.g. a regulatory promoter, an enhancer, a poly-A signal sequence, and a signal peptide encoding region. The choice of the regulatory sequences will highly depend on the host cell to be used for expression of the nucleic acid molecule. Alternatively, the nucleic acid molecule may not contain any further regulatory sequences.

**[0083]** The nucleic acid molecule may be comprised in a vector. Generally, a vector is any vehicle used to integrate foreign nucleic acid material into a cell or cell genome, and typically contains elements that are capable of introducing, maintaining, and/or expressing nucleic acid sequences into a cell or, integrating nucleic acid sequences into the genome of a cell or of a host organism. Preferably, a suitable vector according to the invention further comprises an element selected from the group consisting of an origin of replication, a selectable marker, a transposon, a polytropic transposon, a retrovirus, an element capable of homologous recombination, and an element capable of non-homologous recombination. Generally, a wide variety of suitable vectors are known in the art and available to the skilled person. Accordingly, the invention also relates to a host cell comprising the nucleic acid molecule or the vector described above. Generally, the host cell may be any suitable prokaryotic or eukaryotic cell which is susceptible for transformation or transfection with the nucleic acid molecule or vector described herein. For example, such host cell may be a bacterial cell, e.g. a cell of *E. coli* or *B. subtilis*, a yeast cell, e.g. a *Saccharomyces* cell, a fungal cell, e.g. an *Aspergillus* cell, or a eukaryotic cell, e.g. a mammalian cell, a rodent cell, e.g. a CHO cell, a human cell or an insect cell. An overview of examples of different corresponding expression systems to be used for generating the host cell, is described, for example, in Bitter et al. (Methods in Enzymology 153 (1987), 516-544), in Sawers et al. (Applied Microbiology and Biotechnology 46 (1996), 1-9), Billman-Jacobe (Current Opinion in Biotechnology 7 (1996), 500-4), Hockney (Trends in Biotechnology 12 (1994), 456-463), Griffiths et al., (Methods in Molecular Biology 75 (1997), 427-440), Colonna et al. (1992) J. Imm. Methods 152: 89-104, Gurtu et al. (1996) Biochem. Biophys. Res. Comm. 229: 295-298. The transformation or genetically engineering of the host cell with a nucleic acid molecule or vector can be carried out by standard methods, as for instance described in Sambrook and Russell (2001), Molecular Cloning: A Laboratory Manual, CSH Press, Cold Spring Harbor, N.Y., USA; Methods in Yeast Genetics, A Laboratory Course Manual, Cold Spring Harbor Laboratory Press, 1990.

**[0084]** In particular, such a host cell is useful in producing the antibody molecule. In this context, the invention further provides a method of producing an antibody molecule according to the invention comprising the step of (i) culturing the host cell or the hybridoma of the invention under conditions allowing synthesis of said antibody molecule and (ii) recovering said antibody molecule from said culture. Culturing will depend on the host cell in question, and includes fermentation in small scale and in large scale in laboratory and industrial fermenters, under conditions which are conducive for the production of the antibody molecule. Usually culturing is performed in liquid media comprising, for example, a carbon source, a nitrogen source, trace minerals and metals, as well as growth factors. Suitable culture media

are known in the art. If the antibody molecule is secreted into the culture medium, it can be directly isolated from the culture. Otherwise, it might become necessary to disrupt the cells first in order to release the antibody molecule from the periplasma or cell into the culture medium. Methods for recovering an antibody molecule from a culture are also known in the art and may include precipitation, chromatographic procedures, such as affinity chromatography (e.g. Protein A or Protein G chromatography), hydroxylapatite chromatography, hydrophobicity chromatography, ion exchange chromatography, size exclusion chromatography, and/or electrophoretic procedures. As apparent from the above, these purification steps may be combined, e.g. by first conducting a protein A chromatography and subsequently a hydroxylapatite chromatography. Depending on the purification conditions chosen, the antibody molecule may be separated in the fractions or in the flow through.

**[0085]** As discussed above, it is believed that the antibody molecule described herein is useful and may be used in medicine and veterinary medicine, i.e. it may be used in the manufacture of a medicament, e.g. a medicament for the treatment of Alzheimer's disease. Accordingly, the antibody molecule described herein may be used in the treatment and/or prevention of Alzheimer's disease, in particular wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease. In a preferred embodiment, the Alzheimer's disease is sporadic Alzheimer's disease. In another preferred embodiment, the Alzheimer's disease is familial Alzheimer's disease, including those caused by mutations in APP (arctic and Swedish mutation) and Presenilin-1 (PS1).

**[0086]** For this purpose, the antibody molecule is preferably formulated in a pharmaceutical composition comprising a pharmaceutically acceptable carrier, excipient and/or diluent. The carrier may be chosen dependent on the route of administration as well as on the concentration of the antibody molecule. The pharmaceutical composition may be in the form of a lyophilised composition or an aqueous solution, in particular a stabilized aqueous solution. Generally, an appropriate amount of a pharmaceutically acceptable salt is used in the carrier to render the composition isotonic. For example, the carrier may include but is not limited to phosphate buffered saline, Ringer's solution, dextrose solution, water, emulsions, such as oil/water emulsions, various types of wetting agents, sterile solutions, etc. The acceptable excipients, carriers, or stabilisers are non-toxic at the dosages and concentrations employed, including buffers such as citrate, phosphate, and other suitable organic acids; salt-forming counter-ions, e.g. sodium and potassium; low molecular weight (>10 amino acid residues) polypeptides; proteins, e.g. gelatine, or serum albumin; hydrophilic polymers, such as polyvinylpyrrolidone; amino acids, e.g. histidine, glutamine, lysine, asparagine, arginine, or glycine; carbohydrates including glucose, mannose, dextrans, mono- and/or disaccharides, e.g. sucrose, mannitol, trehalose or sorbitol; complexing agents, e.g. EDTA; non-ionic surfactants, such as Pluronic, Tween, or polyethylene glycol. Preservatives may also be present such as, for example, antimicrobials, antioxidants, chelating agents, and inert gases and the like. Examples for antioxidants and/or preservatives are, e.g. methionine, ascorbic acid, tocopherol, octadecyldimethylbenzyl ammonium chloride, hexamethonium chloride, benzalkonium chloride, benzethonium chloride, phenol, butyl or benzyl alcohol, alkyl parabens such as methyl or propyl paraben, catechol, resorcinol, cyclohexanol, 3-pentanol, or



m-cresol. Suitable carriers and their formulations are described in greater detail in Remington's Pharmaceutical Sciences, 17th ed., 1985, Mack Publishing Co.

**[0087]** Accordingly, in another aspect, the invention pertains to a method of treating or preventing Alzheimer's disease in a subject, comprising the step of administering an antibody molecule according to the invention to said subject.

**[0088]** The subject may be a non-human animal, preferably a mammal such as a horse, cow, pig, mouse, rat, guinea pig, cat, dog, goat, sheep, non-human primate, or a human. The subject may be a healthy subject, a subject having/suffering from Alzheimer's disease, or subjects showing/having susceptibility for the development of Alzheimer's disease (e.g. carrying a genomic mutation which correlates with the occurrence of Alzheimer's disease, such as Swedish mutation, arctic mutation, etc.). Thus, in one preferred embodiment, the Alzheimer's disease is sporadic Alzheimer's disease. In another preferred embodiment, the Alzheimer's disease is familial Alzheimer's disease. Preferably, the antibody molecule is formulated in a pharmaceutical composition comprising a pharmaceutically acceptable carrier, to excipient and/or diluent, as defined above.

**[0089]** In one embodiment, the pharmaceutical composition may comprise the antibody molecule in an amount of about 10 ng/kg to about 5 mg/kg or about 10 mg/kg, and the pharmaceutical composition may be administered at a suitable dose, i.e. about 1 ng/kg body weight to about 100 mg/kg body weight of a subject, preferably at a dose of about 10 ng/kg to about 10 mg/kg, more preferably at a dose of about 10 ng/kg to about 5 mg/kg per body weight. However, the dosage regimen will be determined by an attending physician and depend upon many factors, including the patient's size and condition, body surface area, age, sex, time and route of administration, and on other drugs being administered concurrently. Administration will preferably be intravenously but may also be subcutaneously, intramuscularly, intraperitoneally, intracranially or directly into the cerebral fluid or selected brain regions. The compositions comprising an antibody molecule as described and provided herein may also be administered directly, e.g., by biolistic delivery to an internal or external target site or by catheter to a site in an artery or a vein.

**[0090]** The antibody molecule, preferably formulated in a pharmaceutical composition, may be administered, e.g., enterally, orally (e.g., formulated as a pill, tablet (buccal, sublingual, orally, disintegrating, capsule, thin film, liquid solution or suspension, powder, solid crystals or liquid), rectally (e.g., as a suppository, enema), via injection (e.g., intravenously, subcutaneously, intramuscularly, intraperitoneally, intradermally) via inhalation (e.g., intrabronchially), topically, vaginally, epicutaneously, or intranasally. Preferably, the composition is administered via injection, in particular intravenously, subcutaneously, intramuscularly, intraperitoneally, or intradermally; more preferably the composition is administered intravenously. It is also contemplated to administer the antibody molecule or the pharmaceutical composition directly into cerebral fluid or selected brain regions, i.e. intracranially.

**[0091]** Preparations for parenteral administration include sterile aqueous solutions, e.g., water, alcoholic/aqueous solutions, including saline and buffered media, including include sodium chloride solution, Ringer's dextrose, dextrose and sodium chloride solution, lactated Ringer's solution; or non-aqueous solutions, e.g., propylene glycol, polyethylene gly-

col, vegetable oils (such as olive oil), and injectable organic esters (such as ethyl oleate), suspensions, and emulsions. Intravenous vehicles include fluid and nutrient replenishers, electrolyte replenishers (such as those based on Ringer's dextrose), and the like.

**[0092]** However, the antibody molecule disclosed herein may also be formulated and comprised in a diagnostic composition. In this case, the antibody molecule may be linked with a marker or label, such as a dye, e.g. a fluorescence dye, a marker enzyme, or a compound which is detectable in routine diagnostic methods and medical scanning procedures. Accordingly, the diagnostic composition may be formulated for in vivo or in vitro use.

**[0093]** Hence, also contemplated is the use of an antibody molecule in accordance with the invention in a method of diagnosing Alzheimer's disease, and a method of diagnosing Alzheimer's disease, comprising the step of determining the amount of low molecular weight oligomers, in particular trimeric and tetrameric oligomers of A $\beta$  in a sample of a subject to be diagnosed, using an antibody molecule according to the invention. The sample may be a blood sample, a cerebrospinal fluid sample, a serum sample, or a brain tissue sample. In a preferred embodiment, the step of determining the amount of low molecular weight oligomers of A $\beta$  is conducted by means of an ELISA, e.g. as described above. Alternatively, the step of determining the amount of low molecular weight oligomers of A $\beta$  is conducted by means of immunohistochemistry.

**[0094]** The antibody molecule, e.g. as formulated in a diagnostic composition, may also be used in a method of identifying agents useful in the treatment and/or prevention of Alzheimer's disease, in particular wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease. For example, the antibody molecule may be used as a positive control, or to detect the amount of trimeric or tetrameric A $\beta$  in a immunohistochemistry sample, as described above, in for example a test animal treated with the candidate agent. Accordingly, the antibody molecule disclosed herein may also be used in a method of monitoring the efficacy of a treatment of Alzheimer's disease, in particular wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease.

**[0095]** Thus, also provided is a method of monitoring the efficacy of a treatment of Alzheimer's disease, comprising the step of determining the amount of low molecular weight oligomers of A $\beta$  in a sample of a subject to be diagnosed using an antibody molecule as disclosed herein. Similar to the method of diagnosing Alzheimer's disease, the step of determining the amount of low molecular weight oligomers of A $\beta$  may be conducted by means of an ELISA; and the method may be conducted on a blood sample, a cerebrospinal fluid sample, a serum sample, or a brain tissue sample, as outlined above.

**[0096]** In the following, the present invention is illustrated by reference to figures, sequences and examples which are not intended to limit the scope of the present invention.

#### DESCRIPTION OF THE FIGURES

**[0097]** FIG. 1 Pepsan analysis using ELISA of (A, C) NT4X-167 and (B) NT4X-83 as well as IC16 (D) antibodies. Linear peptides with 16 amino acids in length were probed and incubated with NT4X antibody. NT4X-67 recognized amino acid position 1-4 with phenylalanin at position 4 with the highest intensity. No signals above background were seen

with no peptide, no primary nor secondary antibody. As expected the positive control  $A\beta_{4-42}$  reacted with the antibody. (C) NT4X-167 signal was completely abolished by replacing phenylalanine with alanine (A) or proline (P).

**[0098]** FIG. 2 SDS-PAGE Western blot analysis of NT4X antibodies. Freshly dissolved synthetic Abeta (7  $\mu$ g) variants were probed to a membrane and incubated with (A) NT4X-167 and (B) NT4X-83 antibodies. The antibodies recognized low molecular weight tetrameric oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  as well as monomers and dimers of  $A\beta_{pE3-42}$ ,  $A\beta_{4-42}$ ,  $A\beta_{4-38}$  and  $A\beta_{4-40}$ . This experiment demonstrates that the NT4X antibodies have a preferential binding to low molecular weight oligomers of the most abundant  $A\beta_{x-42}$  variants in the brain of AD patients, but lack binding to  $A\beta_{1-42}$  monomers and dimers.

**[0099]** FIG. 3 Native Western blot of NT4X antibodies. Freshly dissolved  $A\beta$  (7  $\mu$ g) variants were probed to a membrane and incubated with (A) NT4X-167 and (B) NT4X-83 antibodies. The antibodies recognized abundant LMW oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  as well as monomers and dimers of  $A\beta_{pE3-42}$ ,  $A\beta_{4-42}$ ,  $A\beta_{4-38}$  and  $A\beta_{4-40}$ . Of note,  $A\beta_{1-42}$  monomers or dimers were not observed with NT4X antibodies. This experiment demonstrates that the NT4X antibodies have a preferential binding to LMW oligomers of the most abundant  $A\beta_{x-42}$  variants in the brain of AD patients. There was no difference between NT4X-167 and NT4X-83 specificity.

**[0100]** FIG. 4 Western blot of freshly dissolved comparing with aged  $A\beta$  peptides under reducing conditions.  $A\beta$  variants (7  $\mu$ g) were probed to a membrane and incubated with NT4X-167 and IC16 antibodies. (A) Using freshly dissolved peptides NT4X-167 recognized low molecular weight (LMW) oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  as well as monomers and dimers of  $A\beta_{pE3-42}$ ,  $A\beta_{4-42}$ ,  $A\beta_{4-38}$  and  $A\beta_{4-40}$ . It did not recognize monomers and dimers of  $A\beta_{1-42}$ . (B) Using aged peptides NT4X-167 also recognized high molecular weight (HMW) oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$ . (C) Using freshly dissolved peptides IC16 only recognized monomers, dimers and LMW oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$ . Of note, IC16 strongly reacted with  $A\beta_{1-42}$  aggregates followed by  $A\beta_{pE3-42}$ . (D) Aged peptides showed the same profile with a much stronger reactivity of HMW oligomers of  $A\beta_{1-42}$ , which was weaker for  $A\beta_{pE3-42}$ . (E) Under native conditions freshly dissolved AR peptides displayed formation of abundant oligomers of different sizes.  $A\beta_{1-42}$  run mostly as low molecular weight oligomers, while the N-truncated  $A\beta$  peptides displayed prominent bands a higher molecular weight. Native Western blot of  $A\beta$  peptides using the pan- $A\beta$  polyclonal antiserum 24311.

**[0101]** FIG. 5 Sensitivity of NT4X-167 using native Western blotting of freshly dissolved  $A\beta$  peptides. NT4X antibodies recognized  $A\beta_{4-42}$  oligomers with the highest sensitivity followed by  $A\beta_{pE3-42}$  in the  $\mu$ M range with (A) NT4X-167 being more sensitive as 70 ng of peptides having similar signal intensity as 500 ng detected with (B) NT4X-83. (C) represent a shorter exposure compared to A. (D) IC16 detected monomers and oligomers of full length 1-42, but none of the N-truncated peptides.

**[0102]** FIG. 6 Aggregation of  $A\beta$  peptides using ThT fluorescence assay. All  $A\beta$  peptides exhibited a typical aggregation profile. Co-incubation with NT4X-167 inhibited further aggregation: It clearly inhibited aggregation of (A)  $A\beta_{4-38}$ , (B)  $A\beta_{4-40}$  (D)  $A\beta_{1-42}$  and (E)  $A\beta_{pE3-42}$ . (C) The elevated aggregation signal of  $A\beta_{4-42}$  at the beginning of the experi-

ment indicates a poor level of monomers. Incubation with NT4X-167 lowered the aggregation signal demonstrating an inhibitory effect of the antibody treatment.

**[0103]** FIG. 7 Viability assay using SY5Y neuroblastoma cells. NT4X-167 antibody treatment rescued cell viability of neuroblastoma cells SY5Y. The monomerized 10  $\mu$ M peptides were freshly dissolved and incubated together with the antibody for 20 h at 37° C., followed by incubation for 96 h at 4° C. without shaking. Aged peptides with or without antibody were applied to SY5Y cells for 12 h and the viability was measured by MTS. One-way analysis of variance demonstrated a significant difference within the groups of  $p < 0.0001$  ( $F = 6.418$ ,  $df = 11$ ) followed by Dunnett's multiple comparison test between the groups. The cell viability was significantly reduced by all peptides compared to untreated control cells ( $P < 0.01$ ) and was rescued by NT4X-167 treatment in all cases.

**[0104]** FIG. 8 Immunohistochemical staining pattern in superior temporalis gyrus of sporadic AD brain. Upper row shows the staining pattern in a sporadic AD case using IC16 antibody demonstrating different magnifications of cerebral amyloid angiopathy (CAA) and plaques (A-C). Second and the third row show the staining pattern of the same region as IC 16 using two novel antibodies, NT4X-167 and NT4X-83, respectively (D-F) and (G-I). Comparison of figure A with D and G clearly shows that occasionally NT4X-83 and -167 can recognize CAA rather than plaques even when abundant plaques are recognized by IC16. Both NT4X antibodies show a comparable staining pattern. Scale bar: A, D, G: 200  $\mu$ m and B, C, E, F, H, I: 50  $\mu$ m.

**[0105]** FIG. 9 Immunohistochemistry staining of cases with familial AD. Panel A-C and D-F illustrate brain sections from patients with an Arctic and Swedish mutation in the APP gene respectively. Panel 3 shows brain sections from a patient with a PS1 mutation with (G-I). While, shown in all the figures, much CAA is recognized by the tested antibodies (IC16, NT4X-167 and NT4X-83), plaques can be recognized by NT4X only in the APP Arctic and Swedish cases but not in the PS1 Familial AD case. Scale bar: 200  $\mu$ m.

**[0106]** FIG. 10 Immunohistochemical staining of cortical sections of 3 and 6 months old 5XFAD transgenic mice. (A-C) Immunostaining with IC16 demonstrating age-dependent increase in plaque load. (D-F) Immunostaining with NT4X-167 showing age-dependent increase in plaque load however the amount of positives plaques was less pronounced compared to IC16. (G-I) The pattern for NT4X-83 was similar to NT4X-167. Scale bar: A, B, D, E, G and H: 200  $\mu$ m and in image C, F and I: 50  $\mu$ m.

**[0107]** FIG. 11 Immunohistochemical staining of APP/PS1KI transgenic mouse brain. (A-B)

**[0108]** Immunostaining with IC16 showing abundant intraneuronal (small arrow) and plaque (large arrow)  $A\beta$  in the hippocampal formation (subiculum) at 1.5 months of age using IC16 (A) and to a lesser amount using NT4X-167 (B). Blood vessel staining (cerebral amyloid angiopathy, CAA) with IC16 (C) and NT4X-167 (D). Abundant intraneuronal  $A\beta$  staining at 1.5 months of age in sub-cortical region with IC16 (E) and NT4X-167 (F). Scale bars: A-D: 50  $\mu$ m and E-F: 100  $\mu$ m.

**[0109]** FIG. 12 No cross-reactivity of NTX4 antibodies with other aggregates of major neurodegenerative diseases. Alpha-synuclein ( $\alpha$ -syn) positive aggregates in (A) Lewy bodies and Lewy neurites in Dementia with Lewy bodies (DLBD) and (B) Parkinson disease (PD) and (C) glial inclu-

sions in Multiple system atrophy (MSA). (D) Pathological vessels in Binswanger encephalopathy. (E) Phospho-TDP-43 immunoreactive neuronal cytoplasmic in a case with Frontotemporal lobar degeneration (FTLD). Phospho-Tau (pTau) immunoreactive aggregates in (F) Progressive supranuclear palsy (PSP) and (G) in Pick's disease (PiD). (H) Prion protein (PrP) immunopositive synaptic deposits in sporadic Creutzfeldt-Jakob disease (CJD). Scale bar in A-F 50  $\mu\text{m}$ , G-H, 200  $\mu\text{m}$ .

**[0110]** FIG. 13 NT4X treatment rescued toxic effect of freshly dissolved  $\text{A}\beta_{4-42}$  but not of pyroglutamate  $\text{A}\beta\text{pE}_{3-42}$  and only weakly of  $\text{A}\beta_{1-42}$  in vitro. In rat primary cortical neurons, all  $\text{A}\beta$  peptides induced significant cellular toxicity compared to cultures without peptide. NT4X treatment rescued the toxic effects with high potency only of cultures treated with  $\text{A}\beta_{4-42}$ . While NT4X treatment rescued toxicity of 1  $\mu\text{M}$  of  $\text{A}\beta_{1-42}$ , 5 and 10  $\mu\text{M}$  could not be rescued. The toxicity of  $\text{A}\beta\text{DE}_{3-42}$  could not be rescued by NT4X at all concentrations tested.

**[0111]** FIG. 14 Passive immunization with NT4X-167 antibody rescued memory deficits in aged 5XFAD mice. Twenty-four hours after the last acquisition trial, a probe trial was performed to assess spatial reference memory between placebo-treated (PBS) versus NT4X-167 immunized 5XFAD mice. A: At 30 weeks of age, PBS-treated 5XFAD mice displayed no significant preference for the target quadrant, as indicated by the percentage time spent in different quadrants of the pool. B: 5XFAD mice, which received intraperitoneal injections of 10 mg/kg NT4X-167 antibody starting at the age of 20 weeks (10 weekly injections between week 20 and 30) showed a significant preference for the target quadrant indicating that their spatial reference memory was restored. Abbreviations: T=target quadrant; L=left quadrant; R=right quadrant; O=opposite quadrant; m=age in months.

DESCRIPTION OF THE SEQUENCES
NT4X-83 HCDR1 (SEQ ID NO: 1): GFNIRDY
NT4X-83 HCDR2 (SEQ ID NO: 2): VDPANGNT
NT4X-83 HCDR3 (SEQ ID NO: 3): ARRIYYGYALPAY
NT4X-83 LCDR1 (SEQ ID NO: 4): QSLNSGNQKNY
NT4X-83 LCDR2 (SEQ ID NO: 5): WAS
NT4X-83 LCDR3 (SEQ ID NO: 6): QNDYSYPLTY
NT4X-83 VH (SEQ ID NO: 7): EVHLQQSGAELVKPGASVKLSCTASGFNIRDYIHVWVKRPEQGLEWIGRVD PANGNTKYDKPKQKATITADTSSDTAYLQLSSLTAEDTAVYFCARRIYYGYALPAYWQGQTLVTVSA
NT4X-83 VL /SEQ ID NO: 8): DIVMTQSPSSSLTVTAGEKVTMSCKSSQSLNSGNQKNYLTWYQQKPGQPPKLLIYWASTRESGVPDRFPGSGGTDFTLTISVQAEDLAVYCYCQNDYSYPLTFGAGTKLELK
NT4X-167 HCDR1 (SEQ ID NO: 9): GFSLTSYG

-continued

DESCRIPTION OF THE SEQUENCES
NT4X-167 HCDR2 (SEQ ID NO: 10): MWSGGIT
NT4X-167 HCDR3 (SEQ ID NO: 11): ARGSRYALDY
NT4X-167 LCDR1 (SEQ ID NO: 12): QDISNY
NT4X-167 LCDR2 (SEQ ID NO: 13): YTS
NT4X-167 LCDR3 (SEQ ID NO: 14): QQGNTLPPT
NT4X-167 VH (SEQ ID NO: 15): QVQLKQSGPGLVQPSQSLISITCTVSGFSLTSYGIHWVRQSPGKGLEWLGVMWSGGITDFYAAFISRLSISRDISKSKQVFFKMNLSLQADDTAIYYCARGRSRYALDYWQGQTSVSVSS
NT4X-167 VL (SEQ ID NO: 16): DIQMTQTTSSLSASLGLDRVTISCRASQDISNYLNWYQQKPDGTVKLLIYYTSRHLHSGVPSRFRSGSGSDYSLTISNLEQEDIATYFCQQGNTLPPTFGGGKLEIK
Human $\text{A}\beta_{1-42}$ (SEQ ID NO: 17): DAEFRHDSGYEVHHQKLVFPAEDVGSNKGAIIGLMVGGVVIA

## DEPOSITED BIOLOGICAL MATERIAL

**[0112]** The following samples have been deposited in accordance with the Budapest treaty on the international recognition of the deposits of microorganisms for the purposes of patent procedure at the DSMZ, German Collection of Microorganisms and Cell Cultures in Braunschweig, Germany:

Reference given by the Depositor	Date of deposit	Accession number
NT4X-83	7 Mar. 2012	DSM ACC3161
NT4X-167	7 Mar. 2012	DSM ACC3162

## EXAMPLES

**[0113]** Statistical differences were evaluated using one-way ANOVA followed by Dunnett's multiple comparison test or unpaired t-test as indicated. All data are given as mean  $\pm$  standard error of the mean (SEM). All statistics were calculated using GraphPad Prism V5.00 software (USA).

## Example 1

**[0114]** Generation and Screening of Antibodies specific for  $\text{A}\beta_{1-42}$  Trimer/tetramers

**[0115]** The novel oligomeric  $\text{A}\beta$  specific antibodies NT4X-83 (IgG2a; official name of cell line  $\text{A}\beta_{4-40}$  NT4X-83; DSM ACC3161) and NT4X-167 (IgG1; official name of cell line  $\text{A}\beta_{4-40}$  NT4X-167; DSM ACC3162) were generated by immunizing three Balb/c mice with unconjugated  $\text{A}\beta_{4-40}$ . After preparation of the lymph nodes they were fused with the myeloma cell line P3-X63-Ag8 for generation of the hybridoma cells. The hybridoma supernatants of mixed clones were screened by ELISA and subcloned.

**[0116]** The monoclonal antibodies were selected by enzyme-linked immune-absorbent assay (ELISA) and immunohistochemistry. The idea behind the generation of novel oligomeric antibodies was that A $\beta_{4-40}$  peptides are forming dimers in solution that can be used as a stable epitope for antibodies that bind specifically at the N-terminus of A $\beta_{4-40}$ . Therefore A $\beta_{4-40}$  was used for immunizing mice and positive clones screened in four steps. After fusion, the hybridoma cells were screened by an ELISA for antibody production that (1) bind A $\beta_{4-10}$  and (2) A $\beta_{4-40}$ , but (3) not A $\beta_{36-40}$ . Positive antibody clones were further screened by immunohistochemical staining of human brain sections. (4) The last step of the screening procedure was that they should not preferentially bind to amyloid plaques. Two monoclonal antibodies were identified NT4X-83 and NT4X-167 and further characterized.

#### Example 2

**[0117]** Epitope Mapping of NT4X-83 and NT4X-167

**[0118]** Pepsin assays (FIG. 1) were performed in order to identify the epitope binding of NT4X-167 and NT4X-83 as well as IC16. Pepsin is a procedure for mapping and characterizing epitopes involving the synthesis of overlapping peptides and analysis of the peptides in enzyme-linked immunosorbent assay (ELISAs). A $\beta$  peptides with 16 amino acids in length were used: A $\beta_{1-16}$ , 2-17, 3-18, 4-19, 5-20, 6-21, 7-22, 8-23, 9-24 and 10-25. The experiment was performed using U form high binding 96 well-plate (microplates, Greiner bio-one) to coat the peptides. Samples were diluted in ice cold sterile-filtered 0.05 M carbonate buffer (2.4 g NaCO<sub>3</sub> × 10 H<sub>2</sub>O, 1.46 g NaHCO<sub>3</sub> added to 450 ml ddH<sub>2</sub>O, pH 9.6) and 50  $\mu$ l of the diluted samples were coated on the wells (the well-plate was kept on ice for some min before peptide application). For the peptide binding, the well-plate was incubated overnight at 4° C. The next day, the wells were washed three times by 0.01 M PBS (RT). The washing was done by applying 250  $\mu$ l PBS into each well and emptying it by inverting the plate and hit it hard on a spongy cloth.

**[0119]** After the washing step, three other steps were followed (each step for 1 h at 25° C.):

**[0120]** Blocking (preventing unspecific antibody binding).

**[0121]** Incubation with primary antibody ((NT4-X (167) 1:300, 7  $\mu$ g/ml, NT4-X (83) 1:100, 7  $\mu$ g/ml and IC16 1:750).

**[0122]** Incubation with the secondary antibody (I-IRP coupled goat anti-mouse IgG, Dianova, 1:10.000, cat no: 115-035-003).

**[0123]** After each mentioned step, three times washing with 0.01 M PBS was applied.

**[0124]** For the blocking step, 200  $\mu$ l of the RT blocking solution containing 5% w/v Albumin Fraction

**[0125]** V (Roth) in 0.01 M PBS, which was prepared, sterile filtered and kept at 4° C. a day before the use was used.

**[0126]** Both, primary and secondary antibodies, were diluted in the blocking solution and 100  $\mu$ l of each dilution was used for each well.

**[0127]** Lastly, as a HRP enzyme substrate, 50  $\mu$ l of 3,3',5,5'-Tetramethylbenzidine (1-step Ultra TMB

**[0128]** ELISA substrate, Thermo Scientific), that leads to the blue color production when oxidized, was added to each of the wells and the absorbance (OD) was measured within 5-10 min at 652 nm by a microplate reader (pQuant, B Tek Instrument INC) using Mikro Win 2000 software (Berthold

Technologies). The reaction was stopped by adding 50  $\mu$ l of 2 M, H<sub>3</sub>PO<sub>4</sub> which led to the yellow color formation and the absorbance was measured at 450 nm.

**[0129]** Using pepsin ELISA for signal detection, revealed that the binding site of NT4X-167 and NT4X-83 ranged between N-truncated A $\beta_{2-4}$  with the highest signal for N-truncated A $\beta_4$  starting with phenylalanine at position four. No major difference in the binding specificity between both NT4X antibodies was detected. Mutational analysis of A $\beta_{4-19}$  replacing phenylalanine with alanine (A $\beta_{4A-19}$ ) or proline (A $\beta_{4P-19}$ ) inhibited binding of NT4X-167 antibody. Therefore phenylalanine at position four of A $\beta$  appears to be essentially required for antibody binding. IC16 reacted with amino acids 1-4 and is therefore specific for the N-terminus of A $\beta$ .

#### Example 3

**[0130]** Binding Characteristics of NT4X-83 and NT4X-167

**[0131]** For Western blot analysis under reducing conditions, peptides were loaded on 4-12% SDS VarioGels Tris-Tricin (Anamed; cat. no#VG41210), transferred to 0.45  $\mu$ m nitrocellulose membranes and detected using the primary antibodies IC16 (1  $\mu$ g/ml) and NT4X (7  $\mu$ g/ml). Stock solutions of synthetic A $\beta$  peptides (1 mg/ml in 10 mM NaOH) were prepared, sonicated for 5 min in water bath, quickly frozen in liquid nitrogen and stored at -80° C. Using 7  $\mu$ g of freshly dissolved peptides from stock solutions NT4X-167 recognized LMW oligomers of A $\beta_{1-42}$ , A $\beta_{pE3-42}$  and A $\beta_{4-42}$  as well as monomers and dimers of A $\beta_{pE3-42}$ , A $\beta_{4-42}$ , A $\beta_{4-38}$  and A $\beta_{4-40}$ . It did not recognize monomers and dimers of A $\beta_{1-42}$  (exposure time 10 seconds).

**[0132]** For the experiments, desired amount of the purred synthetic full length A $\beta$  peptides were mixed with loading buffer (VarGel SDS Probenpuffer, Anamed) in the ratio of 1:1 and the peptides were denatured for 5-10 min at 95° C. (Uno-Thermoblock, Biometra). The denatured proteins were loaded on 4-12% Tris-Tricin gels with 1 mm of thickness. 5  $\mu$ l of Spectra™ Multicolor low and broad range protein ladders (Thermo and Fermentas) were also loaded on different wells. The proteins were run by VarGel Tricin Running buffer (Anamed) and separated due to electrophoresis starting with 60 V for 15-20 min and continuing with 150 V for about 1 h. After the completion of running, proteins were transferred from the gel to 0.45  $\mu$ m nitrocellulose membrane (GE Healthcare). For transferring the proteins, semi-dry method (BRAD, Trans-Blot® SD Semi-Dry) was used, as followed:

**[0133]** The membrane, gel and also two thick blotting papers were wetted into the transfer buffer. Wet thick papers were placed around the membrane and the gel (the membrane was lying under the gel) and the protein transfer was performed at 25 V for 40 min (Trans-Blot®SD semi-dry Transfer cell, BIO-RAD).

**[0134]** Upon the completion of protein transferring, the membrane was washed two times each for 10 min in 1× TBS/T (0.1 M Tris, 1.5 M NaCl and 0.5% Tween 20) on a rotator (Heidolph, DOUMAX 1030) with speed of 40 rpm. For A $\beta$  peptides, membranes were boiled in 0.1 M PBS for about 1 min in microwave (ShARP) on 800 Watt power. After the start of boiling, the power was decreased to 80 Watt and the boiling was continued for 4 more minutes. Membranes were briefly washed in 1× TBS/T and blocked with 10% non-fat dry milk (Roth) in 1× TBS/T for 1 h at room temperature (RT) (40 rpm shaking). The membrane was incubated with following primary antibodies overnight at 4° C. or 2 h at

RT (40 rpm shaking). This step was followed by two times of washing with  $1 \times$  TBS/T each for 10 min (40 rpm shaking) and the membrane was incubated with secondary antibody (Polyclonal goat anti-mouse, Dianova diluted 1:4000) conjugated with horseradish peroxidase, for 2 h at RT. After washing for two times, each for 10 min, for development, the membrane was incubated in the horseradish peroxidase substrate solution containing 8 ml solution A (0.25 mg/ml luminal (Roth) in 0.1 M Tris/HCl), 800  $\mu$ l solution B (1.1 mg/ml para-coumaric (Roth) acid in DMSO (Sigma) and 2.5  $\mu$ l  $H_2O_2$  (Roth) for 2 min. After slightly drying the membrane on a tissue paper, it was exposed to x-ray films (Hyperfilm EC, Amersham Biosciences) for different time points and developed in a CUR1x60 (AGFA).

**[0135]** For Western blotting under native conditions 18% SDS-free Tris-Glycine Progels (Anamed) [TG18110] were used. NT4X-167 is more sensitive as 70 ng of AB peptides having similar signal intensity as 500 ng detected with NT4X-83. Using these conditions, no signal for  $A\beta_{1-42}$ ,  $A\beta_{4-38}$  and  $A\beta_{4-40}$  was detected. Only LMW of  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$ .

**[0136]** In brief, the peptides were mixed with the dye (Tris-Glycine Native Probenpuffer, Anamed) in the ration of 1:1 for 5 min at RT. Then, the desirable amount of the protein and dye was loaded on the 18% SDS-free gels (Progel Tris Glycine 18%, 1.0 mm, Anamed). Running the proteins on the gel was done using the SDS-free running buffer (Tris-Glycine Native Laufpuffer Anamed). The same nitrocellulose membranes as SDS-reduced Western blotting were used to which the proteins were transferred and the difference in the transferring was the content of the transfer buffer used. In the latter method, instead of 20% methanol, 5% methanol was used. Transferring the proteins to the membrane was accomplished using 25 V for 1 h. All the not mentioned conditions were the same as SDS-reduced Western blotting.

**[0137]** Buffers Used:

**[0138]** SDS-reduced western blotting transfer buffer:

**[0139]** 48 mM Tris-HCL buffer, 39 mM glycine (20% methanol) pH 9.2

**[0140]** Native western blotting transfer buffer:

**[0141]** 48 mM Tris-HCL buffer, 39 mM glycine (5% methanol) pH 9.2

**[0142]** Monomerization of  $A\beta$  Peptides

**[0143]** Synthetic  $A\beta$  peptides (PSL, GmbH, Heidelberg) were dissolved in a final concentration of 1 mg/ml in 10 mM NaOH. Next, the dissolved peptides were sonicated for 5 min with the instrument frequency of 35 KHz and the maximum power of 160 W and frozen in liquid nitrogen and stored at  $-80^\circ$  C. till until further use.

**[0144]** Aging of  $A\beta$  Peptides

**[0145]** Monomerized peptides (1 mg/ml in 10 mM NaOH) were incubated for 20 h in a  $37^\circ$  C. incubator with shaking (60 rpm). Thereafter, they were transferred to  $4^\circ$  C. and were incubated for 4 days (96 h) without shaking.

**[0146]** 1. Preference of NT4X binding for  $A\beta$  tetramers under denaturing conditions

**[0147]** Freshly dissolved  $A\beta$  peptides were subjected to SDS-PAGE to dissect the binding specificity of NT4X antibodies (FIG. 2). Under denaturing conditions NT4X antibodies detected only low molecular weight (LMW) oligomers of  $A\beta_{1-42}$  representing likely tetramers and no monomers or dimers. NT4X detected LMW oligomers derived from  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  with lower reactivity for monomers or dimers. NT4X antibodies detected only monomers and dimers of  $A\beta_{4-38}$  and  $A\beta_{4-40}$ .

**[0148]** 2. Preference of NT4X antibodies for  $A\beta$  low molecular weight oligomers under native conditions

**[0149]** Under native conditions (FIG. 3), freshly dissolved  $A\beta$  peptides ending at amino acid 42 ( $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$ ,  $A\beta_{4-42}$ ) immediately formed oligomers, a tendency not observed for  $A\beta_{4-38}$  and  $A\beta_{4-40}$ . As expected, the oligomers are running as less distinct bands under native conditions as compared to SDS PAGE. Although equal amounts of peptides were loaded, the strongest signal with NT4X antibodies were detected with oligomers of  $A\beta_{4-42}$  followed by  $A\beta_{pE3-42}$  and lastly by  $A\beta_{1-42}$ . NT4X detected only monomers and dimers of  $A\beta_{4-40}$  and  $A\beta_{4-38}$ . The sensitivity for NT4X-167 to detect  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$  was in the low  $\mu$ M range and was more sensitive compared to NT4X-83 (FIG. 5).

**[0150]** 3. Aging of  $A\beta$  induced SDS-resistant high molecular weight oligomers detected by NT4X

**[0151]** A comparison of SDS-PAGE Western blot of freshly dissolved peptides with aged  $A\beta$  peptides (20 h at  $37^\circ$  C. with gentle shaking followed by 96 h at  $4^\circ$  C.) (FIG. 4) demonstrated that NT4X-167 reacted with high molecular weight (HMW) oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE-42}$  and  $A\beta_{4-42}$ . In contrast, IC16 only reacted with  $A\beta_{1-42}$  HMW oligomers.  $A\beta_{4-38}$  and  $A\beta_{4-40}$  did not generate HMW oligomers under reducing condition. These findings argue for a unique and selective binding of NT4X to a conformational epitope formed by aged HMW  $A\beta_{pE-42}$  and  $A\beta_{pE-42}$  oligomers, which is not detectable by IC16 having its epitope between  $A\beta_{1-3}$ . Interestingly, HMW oligomers of  $A\beta_{1-42}$  are also detected by NT4X.

**[0152]** Haass and Selkoe argued that small molecules that can specifically inhibit the formation of  $A\beta$  oligomers and/or prevent their binding to and stabilization on neuronal membranes is at the top of the pathological cascade in AD and will also be beneficial for down-stream events like tangle formation, synaptic and neuron loss and eventually cognitive decline (Haass and Selkoe (2007) *Nat Rev Mol Cell Biol* 8, 101-112). More recently, the group around De Strooper (Benilova et al. (2012) *Nat Neurosci advance online publication*) discussed that it is likely that several of the identified oligomeric species (mostly derived from  $A\beta_{1-42}$ ) have similar or overlapping properties. Moreover, they state that coexistence of several populations that do or do not propagate into fibrils is possible. Despite the differences in structure, stability and concentration, all oligomers may contribute to  $A\beta$  toxicity. They further discussed some technical issues defining oligomers like the apparent 'SDS resistance' (Benilova et al.; supra). It was demonstrated that SDS can induce artificially oligomerization of  $A\beta$ . Monomers, trimers and tetramers as major bands derived from  $A\beta$  oligomers,  $A\beta$  fibrils and  $A\beta$  monomers were isolated after SDS-PAGE separation.

**[0153]** Therefore other techniques like applying native Western blotting is a useful way characterizing  $A\beta$  oligomers. In the present report, we observed a striking difference between the binding of NT4X with  $A\beta$  peptides under reducing and native conditions. While under reducing conditions, LMW oligomers of  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  were detected, a strong signal was seen for monomers of  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$ , but not  $A\beta_{1-42}$ . Under reducing conditions,  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$  formed stable LMW oligomers after dissolving the peptides. Aging of the peptides resulted in a shift of the SDS-stable signal  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and AB to HMW  $A\beta_{4-42}$  to HMW oligomers. No difference was observed between under reducing and native conditions for  $A\beta_{4-38}$  and  $A\beta_{4-40}$ , which apparently only formed monomers and dimers.

## Example 4

**[0154]** Functional Characteristics of NT4X-83 and NT4X-167

**[0155]** Monomerization of synthetic peptides: Stock solutions of synthetic peptides (1 mg/ml in 10 mM NaOH; PSL, Heidelberg) were prepared, sonicated for 5 min in water bath (Sonorex RK 100H, Bandelin electronic), quickly frozen in liquid nitrogen and stored at  $-80^{\circ}\text{C}$ .

**[0156]** Aging of peptides: Aged A $\beta$  peptides were generated by incubation for 20 h at  $37^{\circ}\text{C}$ . with gentle agitation (60 r/min) followed by 96 h at  $4^{\circ}\text{C}$ . without shaking.

**[0157]** 1. Antibody NT4X inhibited A $\beta$  aggregation in vitro

**[0158]** Aggregation of A $\beta$  peptides was investigated online using ThT aggregation assay (Varian fluorescence spectrophotometer) using an excitation wavelength of 446 nm and emission wavelength of 482 nm. Samples contained 10  $\mu\text{M}$  of A $\beta$ , 50 mM sodium phosphate buffer (pH 7.4), 50 mM NaCl, 20  $\mu\text{M}$  ThT and 0.01% sodium azide. The samples were incubated at  $37^{\circ}\text{C}$ . in a peltier adapter with stirring. Data points were recorded every 10 min during the assay. As shown in FIG. 6, all peptides aggregated over time. However, the aggregation of A $\beta_{4-38}$  started significantly later. Of note, the aggregation rate of A $\beta_{4-42}$  differed from all other peptides as they already started aggregation from a higher level. Notably, co-incubation with antibody NT4X-167 efficiently inhibited and/or delayed the aggregation of all A $\beta$  peptides (ratio of 5:1; antibody:A $\beta$ ). This observation argues for a beneficial treatment effect of both NT4X antibodies as they did not show any differences in recognition pattern.

**[0159]** 2. Antibody NT4X rescued A $\beta$ -induced cell toxicity

**[0160]** Next, the toxicity of aged A $\beta$  peptides on SH-SY5Y neuroblastoma cells was studied (FIG. 7). To determine whether the toxic effect of A $\beta$  can be neutralized by NT4X-167 antibody, neuroblastoma cells were co-incubated either with peptides alone or with peptides plus NT4X-167: SH-SY5Y neuroblastoma cells were routinely cultured. 5500 cells were cultured in the wells containing 200  $\mu\text{l}$  of fresh pre-warm DMEM-F12 (PANIT<sup>TM</sup> BIOTECH GmbH) medium for 24 h (at  $37^{\circ}\text{C}$ ., 5%  $\text{CO}_2$ ) and thereafter the cells were starved for 24 h using serum-free pre-warmed medium. After 24 h, medium was replaced by medium containing aged peptides in the presence or absence of NT4X-167 antibody and incubated for 24 h. Cell viability was determined using MTS assay (Promega), according to the manufacturer's instructions compared to vehicle treated control cells. The percentage of cell viability was calculated using the following formula:  $[(A_{490} \text{ sample} - A_{490} \text{ background}) / (A_{490} \text{ control} - A_{490} \text{ background})] \times 100$ .

**[0161]** Using the aged peptide protocol a significant effect on cell viability with all A $\beta$  peptides was observed with the N-truncated peptides having the strongest effect compared to A $\beta_{1-42}$ . Dunnett's multiple comparison test gave the following significant results for the toxicity of 10  $\mu\text{M}$  aged A $\beta$  peptides: control versus A $\beta_{4-38}$  ( $p < 0.05$ ); control versus A $\beta_{4-40}$  ( $p < 0.01$ );

**[0162]** control versus A $\beta_{4-42}$  ( $p < 0.01$ ); control versus A $\beta_{pE3-42}$  ( $p < 0.05$ ). Therapeutic effect of NT4X-167 was demonstrated by increased cell viability comparing application of A $\beta$  versus A $\beta$  and antibody: A $\beta_{4-38}$  ( $p < 0.01$ ); A $\beta_{4-40}$  ( $p < 0.01$ ); A $\beta_{4-42}$  ( $p < 0.01$ ); A $\beta_{pE3-42}$  ( $p < 0.05$ ); A $\beta_{1-42}$  ( $p < 0.01$ ). This observation argues again for a beneficial treatment

effect of both NT4X antibodies as they did not show any differences in recognition pattern.

**[0163]** In view of the above data, the antibodies of the invention are believed to be highly valuable in a therapeutic treatment of AD.

## Example 5

**[0164]** Diagnostic Properties of NT4X-83 and NT4X-167

**[0165]** Human and mouse tissue was processed as described previously (Wirhth et al. (2010) *J. Neural Transm.* 117, 85-96). In brief, 4  $\mu\text{m}$  paraffin sections were pretreated with 30%  $\text{H}_2\text{O}_2$  in PBS to block endogenous peroxidases and antigen retrieval was achieved by boiling sections in 0.01 M citrate buffer pH 6.0, followed by 3 min incubation in 88% formic acid. Primary antibodies were incubated overnight, followed by incubation with biotinylated secondary antibodies (DAKO) before staining was visualized using the ABC method with Vectastain kit (Vector Laboratories) and diaminobenzidine as chromogen.

**[0166]** In brief, 4  $\mu\text{m}$  brain sections were obtained by cutting paraffin-embedded brain tissue using a microtome. Deparaffinization was performed by incubation in xylol (Carl Roth GmbH) two times each for 5 min and followed by series of decreasing ethanol (Chemie-vertrieb) concentration (100%, 95% and 70% for 10, 5 and 3 min respectively) to rehydrate the sections. This stage was followed by 1 min washing the sections with deionized  $\text{H}_2\text{O}$ . Endogenous peroxidase blocking was achieved by incubation of the sections for 30 min in the solution of 200 ml 0.01 M PBS and 2 ml 30%  $\text{H}_2\text{O}_2$ . Sections were washed by deionized  $\text{H}_2\text{O}$  for 1 min. The antigen retrieval was performed by boiling the sections in 10 M Citrate buffer (4.2 g Citric acid-Monohydrate added to 2000 ml  $\text{H}_2\text{O}$ ; pH=6) for 2 min in a microwave at 800 Watt power. After 2 min the power was decreased to 80 Watt and the boiling was continued for 8 more min. After cooling down the sections for 15 min they were washed with deionized  $\text{H}_2\text{O}$  for 1 min followed by incubation in 0.01 M PBS+0.1% Triton for 15 min and for 1 min incubation in 0.01 M PBS as washing step. Incubation with freshly prepared 88% formic acid for 3 min followed. Sections were washed twice by 0.01 M PBS for 1 and 5 min. Unspecific blocking was done by 1 h in 4% milk powder (Roth), 10% fetal cow serum (FCS) in 0.01 M PBS in wet a chamber at RT. Primary antibodies, diluted in 0.01 M PBS and 10% FCS, were applied on the sections and were incubated at RT overnight (in a wet chamber). On the next day, sections were washed three times, each for 5 min by 0.01 M PBS+0.1% Triton and for 1 min by 0.01 PBS. Then, secondary antibody (Rabbit anti-mouse, Dako, cat. no. E0354), conjugated with biotin was diluted 1:200 in 0.01 PBS and 10% FCS and was applied on the sections in the wet chamber and incubated for 1 h at  $37^{\circ}\text{C}$ . This step was followed by Avidin-Biotin complex (ABC) (Vectastain<sup>®</sup> Elite ABC Kit, Vector Laboratories; cat. No. PK6100) incubation at  $37^{\circ}\text{C}$ . for 1.5 h which was prepared and kept at  $4^{\circ}\text{C}$ . at least 30 min before its application. ABC solution was prepared as followed: 0.01 M PBS+10% FCS+1:100 solution A +1:100 solution B. Afterwards, three times washing with 0.01 PBS each time for 5 min, was done and the sections were exposed to diaminobenzidine (DAB) solution (5 ml 50 mM Tris/HCl PH 7.5+100  $\mu\text{l}$  DAB stock solution (25 mg/ml 5-(4-Dimethylamino-Benzylidene)Rhodanine (sigma) in 0.05 M Tris/HCL, PH 7.4)+2.5  $\mu\text{l}$  30%  $\text{H}_2\text{O}_2$  (Roth) (added right before use) for a few minutes until staining was observed. The sections were washed with 0.01 PBS, three times each time for 5

min, counterstaining was performed using filtered hematoxylin, for 40 s. The sections were dipped into deionized H<sub>2</sub>O and then washed in running tap water for 5 min. As the last step in staining, sections were dehydrated by series of increasing ethanol concentration (1 min 70%, 5 min 95% and 10 min 100%) and in the end two times in xylol, each for 5 min. The sections, then, were embedded using eukitt quick hardening mounting medium (Carl Roth GmbH (Roti®-Histokit)).

**[0167]** In order to characterize the staining pattern of the two NT4X antibodies in AD patients, cortical tissue sections with sporadic (Table 1, FIG. 8) and familial AD (Table 2, FIG. 9) were analyzed. Human brain samples were obtained from the Netherlands Brain Bank (NBB) and the Department of Pathology, University of Helsinki, Finland (a generous gift of Dr. Hannu Kalimo) and were approved by the local Ethical Committees. Compared to the IC16, both NT4X antibodies recognized only a minor portion of plaques in brain tissue of AD patients. Cerebral amyloid angiopathy (CAA) staining of blood vessel walls was seen with all antibodies. In familial AD cases, the amount of NT4X-positive plaques was almost absent in patients with a mutation in presenilin-1 gene (PS1), and much weaker in cases with an APP mutation compared to IC16 staining.

**[0168]** Table 1 lists the demographic data of sporadic AD patients and non-demented controls and the staining profile of the antibodies. Of note, none of the controls showed NT4X staining of plaques although 3 of them were positive with IC16 demonstrating a clear differentiation between AD and control cases. The amount of NT4X-positive plaques in sporadic AD brain was low. Abbreviations: No, number of cases; M, male; F, female; ApoE4, number of cases with at least one ApoE4 allele; CM, cerebral amyloid angiopathy

	No	Age		Sex M/F	Braak stage	ApoE4	Plaques (IC16)	CAA (IC16)	Plaques (NT4X)	CAA (NT4X)
		Mean ± SEM								
Sporadic AD	13	76 ± 3		3/10	4-6	7/13	13/13	13/13	3/13	13/13
Controls	10	80 ± 2		6/4	0-1	2/10	5/10	3/10	0/10	3/10

**[0169]** Table 2 lists the demographic data and staining profile of the antibodies in familial AD cases. NT4X did not show any plaques in PS1 cases, while CM and plaques were observable in cases with Arctic and Swedish mutations in the APP gene. Abbreviations: M, male; F, female; CM, cerebral amyloid angiopathy.

Gene	Mutation	Sex	Age	IC 16		NT 4-X	
				Plaques	CAA	Plaques	CAA
APP	Arctic	M	64	+	+	+	+
	Swedish	F	61	+	+	+	+
PS1	Δ Exon 9	M	61	+	+	-	+
		M	64	+	+	-	+
		M	69	+	+	-	+

**[0170]** Next, the staining pattern in two transgenic AD mouse models 5XFAD and APP/PS1KI was analyzed. APP/PS1KI and 5XFAD mice have been described previously (Casas et al. (2004) *Am. J. Pathol.* 165, 1289-1300; Jawhar et al. (2012) *Neurobiol. Aging* 33). All mice were backcrossed for more than 10 generations on a C57BL/6J genetic background and housed at a 12-h day/12-h night cycle with free

access to food and water. All animals were handled according to German guidelines for animal care and studies were approved by the local legal authorities (LAVES). In 5XFAD, IC16 already detected abundant plaques at 3 months of age NT4X antibodies reacted only with a fraction and stained preferentially the central core of plaques (FIG. 10). In APP/PS1KI mice, a model with an earlier and more robust pathology compared to 5XFAD, NT4X detected early aggregates starting at 1.5 months of age preferentially within neurons in the hippocampal formation and subcortical areas, blood vessel walls and some plaques (FIG. 11). These findings argue that NT4X antibodies detect early pathological aggregates in like intraneuronal Aβ aggregation, which has been demonstrated to induce neuron loss in this model.

**[0171]** In order to study a potential cross-reactivity with other disease-typical aggregates, brain tissue sections were stained with disease-specific markers and compared with NT4X reactivity. NT4X antibodies did not cross-react with other aggregated deposits of non-AD neurodegenerative disorders. The following pathological structures showed no immunoreactivity (Table 3; FIG. 12): (i) Phospho-Tau immunoreactive structures, including tufted astrocytes (PSP) and Pick bodies in PiD. (ii) α-Synuclein immunopositive Lewy bodies (brainstem and cortex) and Lewy neurites in PD and DLB, and glial cytoplasmic inclusions in MSA. (iii) Phospho-TDP-43 immunoreactive neuronal cytoplasmic and neuritic deposits in FTLT-DTP, ALS, and AD with limbic TDP-43 deposits. (iv) PrP immunopositive amyloid plaques, synaptic, plaque-like, and perineuronal deposits. In addition, there was no immunoreactivity associated with small vessel disease in Binswanger disease.

TABLE 3

Demographic data and examined anatomical regions of the cases for cases other neurodegenerative disorders.				
No	Case/ Disease	Age	Sex	Examined regions
1	DLBD	81	F	Temporal Cx
2	PD	62	M	Mesencephalon (SN)
3	MSA	52	M	Pons
4	PSP	69	M	Basal Ganglia
5	PiD	70	F	Hippocampus + Ent Cx + Temp Cx
6	FTLD	62	F	Hippocampus + Ent Cx + Temp Cx
7	CJD MV	72	F	Hippocampus + Ent Cx + Temp Cx + Cbl
8	Binswanger disease	49	F	Basal Ganglia

Abbreviations: DLBD, dementia with Lewy bodies; PD, Parkinson's disease; MSA, multiple system atrophy; PSP, progressive supranuclear palsy; PiD, Pick's disease; FTLT, frontotemporal lobar degeneration; CJD, Creutzfeldt-Jakob disease; M, male; F, female; Ent Cx: entorhinal cortex; Temp Cx, temporal cortex, SN: substantia nigra, Cbl: cerebellum.

**[0172]** In view of the above data, it is believed that the antibodies according to the invention are highly valuable for diagnosing AD.

## Example 6

**[0173]** NT4X-167 Rescues A $\beta$ <sub>1-42</sub> and A $\beta$ <sub>4-42</sub> Toxicity In Vitro

**[0174]** To determine whether the toxic effect of A $\beta$  can be neutralized by NT4X-167 antibody, primary cell cultures were co-incubated either with peptides alone or with peptides plus NT4X-167.

**[0175]** Cortical neurons from embryonic day 16-17 Wistar rat fetuses were prepared as previously described (Pillot et al., 1999). In brief, dissociated cortical cells were plated at 50,000 cells/well in 48-well plates precoated with 1.5 mg/mL polyornithine (Sigma). Cells were cultured in a chemically defined Dulbecco's Modified Eagle's/F12 medium free of serum (Gibco) and supplemented with hormones, proteins and salts. Cultures were kept at 35° C. in a humidified 5% CO<sub>2</sub> atmosphere, and at 6-7 DIV, cortical population is determined to be at least 97% neurons. At 6 DIV, the medium was removed and cortical neurons were incubated for 24 h with vehicle or Ab oligomers at the indicated concentrations.

**[0176]** Following a 24-h incubation of primary cortical neurons with A $\beta$  oligomers, cell viability and in vitro toxicity was determined using a calcein-AM assay (Invitrogen, Molecular Probes) (see also Youssef et al., 2008). Briefly, cells were washed twice with PBS and incubated protected from light for 30 min at room temperature in the presence of 2 mM calcein-AM solution prepared in PBS. Cells were then washed twice with PBS and incubated for 15 min at room temperature in PBS containing 1% Triton X-100 (v/v). The level of calcein fluorescence was monitored by fluorescence emission at 530 nm after exciting at 485 nm, using a Fluostar microplate reader (BMG-Labtechnologies, France).

**[0177]** NT4X treatment rescued toxic effect of freshly dissolved A $\beta$ <sub>4-42</sub>, but not of pyroglutamate A $\beta$ pE<sub>3-42</sub> and only weakly of A $\beta$ <sub>1-42</sub> in vitro (cf. FIG. 13). In rat primary cortical neurons, all A $\beta$  peptides induced significant cellular toxicity compared to cultures without peptide. NT4X treatment rescued the toxic effects with high potency only of cultures treated with A $\beta$ <sub>4-42</sub>. While NT4X treatment rescued toxicity of 1  $\mu$ M of A $\beta$ <sub>1-42</sub>, 5 and 10  $\mu$ M could not be rescued. The toxicity of A $\beta$ pE<sub>3-42</sub> could not be rescued by NT4X at all concentrations tested.

## Example 7

**[0178]** Spatial Reference Memory by Morris Water Maze

**[0179]** Spatial reference memory in 5XFAD mice (Oakley et al., 2006) was evaluated using the Morris water maze (Morris, 1984). Thereby, mice learn to use spatial cues to locate a hidden, circular platform (10 cm) in a circular pool (110 cm diameter) filled with tap water. The water was made opaque by adding non-toxic white paint and maintained at 20° C. for the test duration. The pool was divided into four virtual quadrants that were defined based on their spatial relationship to the platform: left, right, opposite and target quadrant, which contains the goal platform. ANY-Maze video tracking software (Stoelting Co., Wood Dale, USA) was used to record escape latency, path length, swimming speed and quadrant preference. In order to test whether the groups differed regarding their memory for the former location of the platform in the probe trial, we calculated for each mouse a platform quadrant preference ratios follows: Time spent in Target Quadrant (time spent in Target Quadrant+Time spent in Opposite Quadrant). Preference ratios close to 1 indicate well, whereas ratios close to 0 indicate poor spatial memory.

**[0180]** Female 5XFAD mice were tested at the age of 30 weeks. Each individual mouse was tested at one age only using the cued trials followed by the acquisition training and finalized by the probe trial. After the probe trial the mice were sacrificed. Testing began with three days of cued training. For these trials the platform was marked with a triangular flag. Mice were introduced into the water at the edge of the pool facing the wall. They were then given one minute to find the submerged platform. Mice that failed to find the platform in 60 sec were gently guided to it. All mice were allowed to sit on the platform for 10 sec before being removed from the pool. To prevent hypothermia, all mice were kept in front of a heat lamp for 3 min before being returned to their home cage. Each mouse received four training trials per day with an average inter-trial interval of 15 min. Both the location of the platform and the position at which mice were introduced into the pool changed between trials.

**[0181]** Twenty-four hours after the last day of cued training, mice performed five days of acquisition training. For this part of testing, the flag was removed from the platform. In addition to the distal cues existing in the room proximal visual cues were attached to the outside of the pool. The platform location remained stationary for each mouse throughout training. At the start of every trial, mice were introduced into the pool from one of four predefined entry points. The order in which these entry points were used varied between training days (Vorhees and Williams, 2006). To avoid quadrant bias, the experimental cohorts were randomly split and trained to find one of two different platform locations. Trials were conducted as during the cued training phase.

**[0182]** Twenty-four hours after the last acquisition trial, a probe test was performed to assess spatial reference memory. The platform was removed from the pool, and mice were introduced into the water from a novel entry point. Mice were then allowed to swim freely for one minute while their swimming path was recorded.

**[0183]** Twenty-four hours after the last acquisition trial, a probe trial was performed to assess spatial reference memory between placebo-treated (PBS) versus NT4X-167 immunized 5XFAD mice. At 30 weeks of age, PBS-treated 5XFAD mice displayed no significant preference for the target quadrant, as indicated by the percentage time spent in different quadrants of the pool (cf. FIG. 14 A). 5XFAD mice, which received intraperitoneal injections of 10 mg/kg NT4X-167 antibody starting at the age of 20 weeks (10 weekly injections between week 20 and 30) showed a significant preference for the target quadrant indicating that their spatial reference memory was restored (cf. FIG. 14 B).

**[0184]** All references cited herein are incorporated by reference.

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**[0186]** EP 2 246 427  
**[0187]** WO 2009056490  
**[0188]** WO 2009065054  
**[0189]** WO 2011151076  
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**1-15.** (canceled)

**16.** A monoclonal antibody molecule recognizing a conformational epitope formed by Phe at amino acid position 4 of trimeric or tetrameric human A $\beta$ ,

as determined by pepscan ELISA comprising the steps of precoating with 100 ng A $\beta$ <sub>4-19</sub> and A $\beta$ <sub>5-20</sub>, blocking, incubating with 7  $\mu$ g/ml of said antibody molecule, and detecting said antibody molecule with a labeled secondary antibody;

wherein said antibody molecule does not bind human monomeric A $\beta$ <sub>1-42</sub> or human A $\beta$ <sub>1-42</sub> dimers,

as determined by native Western-blotting comprising the steps of applying 7  $\mu$ g A $\beta$ <sub>1-42</sub> on a 18% SDS-free polyacrylamide gel, transferring on a 0.45  $\mu$ m nitrocellulose membrane, blocking, incubating with said antibody molecule in a concentration of not more than 10  $\mu$ g/ml for 2 h at room temperature, and detecting said antibody molecule with a labeled secondary antibody.

**17.** The antibody molecule of claim **16**, wherein the epitope further comprises the amino acids at position 2 and 3 of human A $\beta$ , as determined by pepscan ELISA.

**18.** The antibody molecule of claim **16**, wherein the trimeric or tetrameric A $\beta$  oligomers are oligomers of A $\beta$  1-X, A $\beta$  4-X, pyro-Glu-A $\beta$  3-X, or combinations thereof, wherein X is 42, 40 or 38.

**19.** The antibody molecule of claim **18**, wherein X is 42.

**20.** The antibody molecule of claim **16**, wherein said antibody molecule comprises

- (a) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 1, 2 and 3, and a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 4, 5 and 6 (NT4X-83); or
- (b) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 9, 10 and 11, and a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 12, 13 and 14 (NT4X-167); or
- (c) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 1, 2 and 3 (NT4X-83); or
- (d) a variable region that comprises a H-CDR1, a H-CDR2 and a H-CDR3 as shown in SEQ ID NOs: 9, 10 and 11 (NT4X-167); or
- (e) a variable region that comprises a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 4, 5 and 6 (NT4X-83); or

(f) a variable region that comprises a L-CDR1, a L-CDR2 and a L-CDR3 as shown in SEQ ID NOs: 12, 13 and 14 (NT4X-167).

**21.** The antibody molecule of claim **16**, wherein said antibody molecule comprises

- (a) a variable region comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 7, and comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 8 (NT4X-83); or
- (b) a variable region comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 15, and comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 16 (NT4X-167); or
- (c) a variable region comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 7 (V14 NT4X-83); or
- (d) a variable region comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 15 (VH NT4X-167); or
- (e) a variable region comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 8 (VL NT4X-83); or
- (f) a variable region comprising an amino acid sequence which is at least 80% identical to the amino acid sequence as shown in SEQ ID NO: 16 (VL NT4X-167).

**22.** The antibody molecule of claim **16**, wherein said antibody molecule also binds HMW oligomers of A $\beta$  1-42, pyro-Glu-A $\beta$  3-42, A $\beta$  4-42, or combinations thereof, as determined by native Western-blotting comprising the steps of applying 7  $\mu$ g A $\beta$ <sub>1-42</sub> on a 18% SDS-free polyacrylamide gel, transferring on a 0.45  $\mu$ m nitrocellulose membrane, blocking, incubating with said antibody molecule in a concentration of not more than 10  $\mu$ g/ml for 2 h at room temperature, and detecting said antibody molecule with a labeled secondary antibody.

**23.** The antibody molecule of claim **16**, wherein said antibody molecule recognizes aggregates that are specific for Alzheimer's disease, as determined by immunohistochemistry of brain tissue sections.

**24.** The antibody molecule of claim **23**, wherein the Alzheimer's disease is familial Alzheimer's disease.

**25.** The antibody molecule of claim **16**, wherein said antibody molecule inhibits further aggregation of trimeric or tetrameric A $\beta$  oligomers of A $\beta$  1-X, A $\beta$  4-X, pyro-Glu-A $\beta$  3-X, or combinations thereof, wherein X is 42, 40 or 38, when tested in a concentration of 10  $\mu$ M in 50 mM sodium phosphate buffer, 50 mM NaCl, 20  $\mu$ M thioflavin T and 0.01% sodium azide, pH 7.4, at 37° C. in a peltier adapter with stirring in a thioflavin T aggregation assay.

**26.** The antibody molecule of claim **25**, wherein X is 42.

**27.** The antibody molecule of claim **16**, wherein said antibody molecule competes for the same epitope with an antibody selected from the group of antibodies consisting of NT4X-83, as deposited under DSM ACC3161, and NT4X-167, as deposited under DSM ACC3162.

**28.** The antibody molecule according to claim **16**, wherein said antibody molecule is selected from a recombinant full antibody (immunoglobulin), a F(ab)-fragment, a F(ab)<sub>2</sub>-fragment, a F(v)-fragment, a single-chain antibody, a chimeric antibody, a CDR-grafted antibody, a bivalent antibody-construct, a polyvalent antibody-construct, a synthetic antibody, a cross-cloned antibody, a fully-human antibody, a humanized antibody, nanobodies, and diabodies, polyvalent nanobodies or diabodies.

**29.** The antibody molecule according to claim **16**, wherein said antibody molecule is an antibody produced by hybridoma as deposited under DSM ACC3161 or hybridoma as deposited under DSM ACC3162.

**30.** The antibody molecule according to claim **16**, wherein pepscan ELISA is carried out using a 96 well-plate precoated overnight with 100 ng A $\beta$ <sub>4-19</sub> and A $\beta$ <sub>5-20</sub> in 0.05 M carbonate buffer, pH 9.6, at 4° C., which has been subsequently washed three times with 250  $\mu$ l 0.01 M PBS, blocked with, 200  $\mu$ l 5% w/v Albumin Fraction V in 0.01 M PBS for 1 h at 25° C., followed by three times washing with 250  $\mu$ l 0.01 M PBS, and incubating with a concentration of the antibody molecule of 7  $\mu$ g/ml in 100  $\mu$ l of 5% w/v Albumin Fraction V in 0.01 M PBS for 1 h at 25° C., followed by three times washing with 250  $\mu$ l 0.01 M PBS, and incubation for 1 h at 25° C. with a horseradish peroxidase conjugated secondary antibody recognizing the antibody molecule in 100  $\mu$ l of 5% w/v Albumin Fraction V in 0.01 M PBS, followed by three times washing with 250  $\mu$ l 0.01 M PBS, and addition of 50  $\mu$ l of 3,3',5,5'-Tetramethylbenzidine.

**31.** The antibody molecule according to claim **16**, wherein native Western-blotting is carried out using 7  $\mu$ g A $\beta$ <sub>1-42</sub> on a 18% SDS-free polyacrylamide gel, semi-dry blotting transfer on a 0.45  $\mu$ m nitrocellulose membrane, two times washing for 10 min in 1 TBS/T (0.1 M Tris, 1.5 M NaCl and 0.5% Tween 20) on a rotator with 40 rpm, blocking with 10% non-fat dry milk in 1 $\times$ TBS/T for 1 h at room temperature on a shaker with 40 rpm, incubation with said antibody molecule in a concentration of not more than 10  $\mu$ g/ml for 2 h at room temperature on a shaker with 40 rpm, two times of washing with 1 $\times$ TBS/T each for 10 min with 40 rpm shaking, incubation with a secondary antibody conjugated with horseradish peroxidase and recognizing the antibody molecule, for 2 h at RT, washing for two times for 10 min with 1 $\times$ TBS/T, and incubation for 2 min with a solution of 8 ml 0.25 mg/ml luminal in 0.1 M Tris/HCl, comprising 800 ml 1.1 mg/ml paracoumaric acid in DMSO and 2.5  $\mu$ l H<sub>2</sub>O<sub>2</sub>, and exposition to x-ray film.

**32.** A method of medical or veterinary treatment, comprising the step of administering the antibody molecule according to claim **16** to a subject.

**33.** A method of treating or preventing Alzheimer's disease, comprising the step of administering an antibody molecule according to claim **16** to a subject.

**34.** The method of claim **33**, wherein the Alzheimer's disease is sporadic Alzheimer's disease or familial Alzheimer's disease.

**35.** A method of diagnosing Alzheimer's disease, comprising the step of determining the amount of trimeric and tetrameric oligomers of A $\beta$  in a sample of a subject to be diagnosed using an antibody molecule according to claim **16**.

**36.** A method of identifying agents useful in the treatment and/or prevention of Alzheimer's disease, comprising the steps of treating a test animal treated with the candidate agent and detecting the amount of trimeric and tetrameric oligomers of A $\beta$  in an immunohistochemistry sample of said animal using an antibody according to claim **16**.

**37.** A method of monitoring the efficacy of a treatment of Alzheimer's disease, comprising the step of determining the amount of trimeric and tetrameric oligomers of A $\beta$  in a sample of a subject to be diagnosed using an antibody molecule according to claim **16**.

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