



## N-Truncated Aβ Starting at Position Four—Biochemical Features, Preclinical Models, and Potential as Drug Target in Alzheimer's Disease

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The discussion of whether amyloid plaque  $A\beta$  is a valid drug target to fight Alzheimer's disease (AD) has been a matter of scientific dispute for decades. This question can only be settled by successful clinical trials and the approval of disease-modifying drugs. However, many clinical trials with antibodies against different regions of the amyloid  $A\beta$  peptide have been discontinued, as they did not meet the clinical endpoints required. Recently, passive immunization of AD patients with Donanemab, an antibody directed against the N-terminus of pyroglutamate  $A\beta$ , showed beneficial effects in a phase II trial, supporting the concept that N-truncated  $A\beta$  is a relevant target for AD therapy. There is long-standing evidence that N-truncated  $A\beta$  variants are the main variants found in amyloid plaques besides full-length  $A\beta_{1-42}$ , t, therefore their role in triggering AD pathology and as targets for drug development are of interest. While the contribution of pyroglutamate  $A\beta_{3-42}$  to AD pathology has been well studied in the past, the potential role of  $A\beta_{4-42}$  has been largely neglected. The present review will therefore focus on  $A\beta_{4-42}$  as a possible drug target based on human and mouse pathology, *in vitro* and *in vivo* toxicity, and anti- $A\beta_{4-x}$  therapeutic effects in preclinical models.

Keywords: N-truncated A $\beta$ , Tg<sub>4-42</sub>, transgenic mouse model, immunotherapy, neuron loss, PET, *in vivo* imaging, memory decline

## INTRODUCTION

Even though the field of Alzheimer's disease (AD) research has rapidly developed over the last decade, there is still a lack of disease-modifying therapies. Passive immunization with Donanemab a pyroglutamate A $\beta$  (A $\beta_{pE3}$ ) specific antibody showed disease-modification on cognition and for the ability to perform the activities of daily living (Mintun et al., 2021). No biomarkers are yet available based on N-truncated A $\beta$  although autoantibodies were identified in plasma (Marcello et al., 2011). Trieb et al. (1996) investigated whether amyloid- $\beta$  peptides may be relevant targets for the immune system using peripheral blood lymphocytes from healthy blood donors and patients with AD. While healthy donors elicited normal proliferative responsiveness after stimulation, a significant reduction was observed using lymphocytes from AD patients. Meanwhile, lower levels of naturally occurring anti-A $\beta$  auto-antibodies have also been reported in the CSF (Du et al., 2001) and sera (Weksler et al., 2002) of AD patients, and elevated serum levels were also reported (Nath et al., 2003).

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Bayer TA (2021) N-Truncated Aβ Starting at Position Four—Biochemical Features, Preclinical Models, and Potential as Drug Target in Alzheimer's Disease. Front. Aging Neurosci. 13:710579. doi: 10.3389/fnagi.2021.710579 It is also of note that in plasma of patients with mild cognitive impairment (MCI) and AD reduced pools of autoantibodies of the IgM class directed against pyroglutamate  $A\beta_{3-X}$  ( $A\beta_{pE3-X}$ ) have been reported (Marcello et al., 2009). In MCI patients, the level of the autoantibodies correlated with cognitive performance as evaluated by mini-mental state examination. Nand C-terminally truncated  $A\beta$  variants, their potential function and toxicity as well as their potential as drug targets were discussed recently (Bayer and Wirths, 2014; Dunys et al., 2018; Wirths and Zampar, 2019). The current mini-review, discusses the potential role of N-truncated  $A\beta$  in AD, focussing on N-truncated  $A\beta$  starting with position four  $A\beta_{4-42}$ .

### DISCOVERY OF N-TRUNCATED $A\beta_{4-42}$ AND PREVALENCE IN THE HUMAN BRAIN

As revealed by a study by Portelius et al. (2010), the relative prevalence of full-length and N-truncated AB is of significant interest within the Alzheimer field. The authors used A $\beta$  antibodies binding to A $\beta_{4-9}$  and A $\beta_{8-22}$  for immunoprecipitation. This was followed by mass spectrometry for identification of all A $\beta$  variants in post-mortem tissue from patients with sporadic AD, familial AD with mutations in the presenilin-1 (PS-1; PSEN-1), or amyloid precursor protein (APP) genes. The authors demonstrated that the dominating A $\beta$  isoforms are A $\beta_{1-42}$ , A $\beta_{pE3-42}$ , A $\beta_{4-42}$ , and A $\beta_{1-40}$ . The most prevalent variants in the hippocampus and the cortex were  $A\beta_{1-42}$  and  $A\beta_{4-42}$ . The importance of  $A\beta_{4-42}$  did not receive appropriate attention in the past, although N-truncated  $A\beta_{4-X}$ has been discovered with the first sequencing endeavors of  $A\beta$ peptides isolated from plaque cores. This surprising finding puzzled Masters et al. (1985) as the most abundant variant of AB in the formic acid soluble fraction of plaque cores and subsequent peptide sequencing started with phenylalanine at position four  $(A\beta_{4-X})$  and not with the full-length  $A\beta_{1-X}$  they had hoped.

Glenner and Wong (Glenner and Wong, 1984) published ground-breaking work showing the full-length sequence of Ab peptides derived from the vasculature of AD patients. A number of reports have discussed the dominant presence of N-truncated AB variants within amyloid plaques in AD and Down syndrome patients (Harigaya et al., 2000; Tekirian, 2001; Miravalle et al., 2005; Piccini et al., 2005; Jawhar et al., 2011; Bayer and Wirths, 2014), while others suggest that full-length Aβ<sub>1-42</sub> is pathologically relevant (Haass et al., 1992; Näslund et al., 1994; Selkoe, 2001; Walsh et al., 2002). Haass et al. (1993) discovered that AB is produced as a normal physiological process in APP transfected cell lines and cultured cells. The cells were analyzed by epitope mapping and radiosequencing of secreting A $\beta$  variants. They observed mainly full-length A $\beta_{1-42}$  (aspartate-1), but also two other A $\beta$  peptides starting with the amino acid phenylalanine at position four and glutamate at position 11.

Although different methodologies for extracting and solubilizing aggregated A $\beta$  have the potential for over- or underestimating the relative prevalence of the various A $\beta$  pools within amyloid plaques, there is general agreement that the C-terminus mostly ends with A $\beta_{X-42}$  (alanine-42) and less abundantly with  $A\beta_{X-40}$  (arginin-40). For example, Ancolio et al. (1999) have demonstrated a large elevation of N-truncated  $A\beta_{x-42}$  in familial AD, postulating that all  $A\beta_{x-42}$  variants are the main factors driving AD pathology.

The idea that N-truncated Aß may represent a potential drug target to fight AD has been largely neglected but was brought into attention recently (Jawhar et al., 2011; Bayer and Wirths, 2014; Cabrera et al., 2018). After this original discovery (Masters et al., 1985), other research groups have verified the presence of  $A\beta_{4-42}$ by other methodologies. Miller et al. (1993) employed matrixassisted, laser-desorption-time-of-flight mass spectrometry of AB peptides isolated from plaque cores or the cerebrovasculature obtained from patients with AD. The authors demonstrated that the C-terminus of A $\beta$  peptides within plaques ended with A $\beta_{42}$ , whereas cerebrovascular A $\beta$  ended at A $\beta_{40}$ . They also proved that N-truncated  $A\beta_{4-X}$  represents a dominant fraction within plaque cores. Lewis et al. (2006) employed surface-enhanced laser desorption/ionization mass spectrometry for identifying the composition of AB peptides in AD and vascular dementia patients. In the brain of AD patients,  $A\beta$  started mostly with Phe-4, but other N-termini starting with Asp-1, Ala-2, pE-3, and Arg-5 were also detected in extractions from plaque cores. Using specific A $\beta$  antibodies, Lemere et al. (1996) showed that patients with Down syndrome harbor Aß starting at Asp-1 or pE-3 in a subset of plaques. Zampar et al. (2020) used capillary isoelectric focusing for showing Aβ antibody specificity of the N-terminus. Furthermore, the authors used immunohistochemistry with the verified N-terminal specific AB antibodies to stain post-mortem brain tissue from sporadic AD patients. They concluded that the staining signal for  $A\beta_{1-X}$  was much weaker in plaques as compared to cerebrovascular amyloid. In contrast, the signal for  $A\beta_{4-X}$  was much more evident in amyloid plaques.

Sergeant et al. (2003) verified that N-truncated A $\beta$  represented the majority of all variants in both AD and pre-symptomatic AD with a substantial amount being A $\beta_{4-42}$ . The authors explored brain specimen from non-demented individuals with low amyloid load and tangle formation by Western blotting and mass spectrometry of the formic acid soluble fraction of amyloid plaques. Rosen et al. (2016) compared the composition of amyloid peptides isolated from AD neocortex and aged squirrel monkeys using immunochemical staining with an A $\beta_{4-X}$  specific antibody and by mass spectrometry. The authors confirmed the high prevalence of N-truncated A $\beta$  peptides including A $\beta_{4-42}$  in the AD brain, while the prevalence in the non-human primate brain was low.

Although it has been shown that  $A\beta_{4-X}$  is generated and secreted *in vitro*, it was a matter of concern whether N-truncation and post-translational modifications of  $A\beta$ represent a post-mortem artifact due to long-term storage or tissue handling. Such an assumption can now be neglected with the therapeutic effect of an  $A\beta_{pE3}$ -specific antibody in patients with AD after passive immunization (Mintun et al., 2021). Wildburger et al. (2017) employed high-resolution mass spectrometry to explore the question of whether N-truncation and other post-translational modifications of  $A\beta$  are found in AD brains due to post-mortem artifacts. The authors studied different  $A\beta$  pools depending on their solubility and concluded that the N-truncated variants did not correlate with post-mortem interval.

# $A\beta_{4-X}$ CAN BE GENERATED BY ENZYMATIC CLEAVAGE

The generation of N-terminal  $A\beta_{4-X}$  by known enzymatic cleavages has been reviewed before in detail (Baver and Wirths, 2014). It can be generated by a two-step process starting with  $\beta$ site APP cleaving enzyme 1 (BACE-1) cutting APP between Met at postion-1 and Asp at position +1 liberating the N-terminus of full-length  $A\beta_{1-X}$  (Vassar et al., 1999). ADAMTS4 (a disintegrin and metalloproteinase with thrombospondin motifs 4) and neprilysin (NEP) further cuts between Glu at position +3 and Phe at position +4 liberating the N-terminus of  $A\beta_{4-X}$  (Bayer and Wirths, 2014; Walter et al., 2019; Figure 1A). NEP cleaves AB at multiple sites thereby detoxifying amyloid-β peptides (Bayer and Wirths, 2014). Walter et al. (2019) identified a recognition site for the secreted form of metalloprotease ADAMTS4 within the full-length Aβ sequence. The induction of ADAMTS4 expression in cell culture led to increased secretion of  $A\beta_{4-40}$  the levels of  $A\beta_{1-x}$  were not altered. Furthermore, the authors identified adult oligodendrocytes as the only source of ADAM4TS triggered AB4-X generation in the murine brain. The main function of NEP on A $\beta$  is degradation and catabolism of the peptide (Iwata et al., 2001; Leissring et al., 2003). The loss of NEP activity leads to enhanced levels of brain and plasma levels of full-length AB, the elevated half-life of soluble AB, and increased amyloid plaque pathology in the J9 mouse model of AD (Farris et al., 2007). Hama et al. (2004) showed that different intracellular compartments are involved in the degradation of amyloid peptides by NEP. Thus NEP is mainly responsible for the intraand extracellular neuronal clearance of AB peptides and more importantly also at the presynaptic site (Iwata et al., 2004). Elevated NEP impaired hippocampal synaptic plasticity and cognitive function in the APP23 mouse model for AD (Huang et al., 2006). Besides general clearance of full-length A $\beta$  peptides, NEP is also involved in the generation of N-truncated Aβ peptides  $A\beta_{4-X}$  (Bayer and Wirths, 2014), and further clearing of  $A\beta_{4-42}$  *in vivo* and *in vitro* (Hornung et al., 2019).  $A\beta_{4-9}$ , a main degradation product of NEP, is a major Cu<sup>2+</sup> binding and has been suggested as a possible Cu<sup>2+</sup> carrier in the brain (Bossak-Ahmad et al., 2019) and NEP modulation (Mital et al., 2018). Modulating Cu metabolism is discussed as a relevant therapeutic target (Lei et al., 2020). In APP23 mice, Cu supplementation lowered amyloid plaque load and stabilized Cu-dependent superoxide dismutase-1 activity (Bayer et al., 2003). In mild to moderate AD patients cognitive decline correlated with low plasma concentrations of Cu (Pajonk et al., 2005). However, treatment for 12 months with supplemental Cu had no effect on cognition in patients with mild AD in a phase 2 clinical trial (Kessler et al., 2008). Alternatively, N-truncated  $A\beta_{4-X}$  may be generated directly by cutting APP between Glu at position 3 and Phe at position 4 by unknown enzymatic activity. Of note, N-truncated  $A\beta_{4-X}$  is secreted together with  $A\beta_{1-42}$  in APP overexpressing cells in vitro, which may indicate unknown enzymatic activity in neurons (Haass et al., 1993). For example,



**FIGURE 1** | Schematic presentation of the potential generation of N-terminal A $\beta_{4-x}$  with known enzymatic cleavages of the amyloid precursor protein (APP) and A $\beta_{1-x}$ . (A)  $\beta$ -site APP cleaving enzyme 1 (BACE-1) or meprin- $\beta$  cuts APP between amino acid position -1 and +1 liberating the N-terminus of full-length A $\beta_{1-x}$ . While ADAMTS4 further cuts between position +3 and +4 liberating A $\beta_{4-40/42}$ , the normal function of neprilysin is to detoxify A $\beta$  as it has several N-terminal activities within A $\beta$  with N-truncation of A $\beta_{4-x}$  being only one alternative. (B) Hypothetical pathway(s) by which N-truncated A $\beta_{4-x}$  is generated by sequential cleavage by meprin- $\beta$ , aminopeptidase A (APA), and/or dipeptidyl peptidase 4 (DPP 4).

N-truncated A $\beta_{5-X}$  is mainly produced from the caspase-cleaved form of APP and not from full-length A $\beta$  (Murayama et al., 2007). Another potential alternative may be the generation the A $\beta$ 4-x peptide, a sequential cleavage of full-length A $\beta$  by aminopeptidase A, meprin- $\beta$  or dipeptidyl-peptidases (Sevalle et al., 2009; Antonyan et al., 2018; Schlenzig et al., 2018; Valverde et al., 2021). At least theoretically, A $\beta_{4-x}$  could be derived from A $\beta_{2-x}$  or A $\beta_{3-x}$  peptide (**Figure 1B**).

## ACUTE EFFECT OF N-TRUNCATED $A\beta_{4-42}$

Exposure of soluble oligomers of  $A\beta_{4-42}$  preparations induced neuron degeneration after 7 days of cultured rat primary cortical neurons (Antonios et al., 2013, 2015). Injecting  $A\beta_{4-42}$  into the lateral ventricles of wildtype mice induced working memory deficits after 4 days (Antonios et al., 2013, 2015). In both assays, the neurodegenerative effect of A $\beta_{4-42}$  was similar to the exposure of  $A\beta_{1-42}$  and  $A\beta_{pE3-42}$ . In 1995, Pike et al. (1995) claimed that  $A\beta$  peptides with N-terminal truncations including  $A\beta_{4-X}$  exhibited enhanced peptide aggregation relative to full-length AB species. Furthermore, they concluded from obtained CD (circular dichroism) spectra that  $\beta$ -sheets were the predominantly formed conformations by all AB variants. They exhibited fibrillary morphology viewed by transmission electron microscopy, and induced degeneration of cultured rat hippocampus neurons. Bouter et al. (2013) demonstrated that soluble oligometric aggregates derived from  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$ have specific structural features distinct from full-length  $A\beta_{1-42}$ , although fibril formation as reported previously (Pike et al., 1995) was comparable. Spectral alterations using ultraviolet circular dichroism spectroscopy showed that  $A\beta_{4-42}$  forms a folded conformation upon heating (Bouter et al., 2013). Using

the liquid state nuclear magnetic resonance technique  $A\beta_{4-42}$ and  $A\beta_{pE3-42}$  exhibited soluble and stable aggregates, which was less pronounced for  $A\beta_{1-42}$ . However, the size of  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$  aggregates were different from those formed by  $A\beta_{1-42}$ (Bouter et al., 2013). Even though  $A\beta_{1-42}$ ,  $A\beta_{pE3-42}$  and  $A\beta_{4-42}$ are unstructured in the monomeric state. After heating all  $A\beta$ peptides formed of folded structures. Interestingly, monomeric  $A\beta_{4-42}$  and  $A\beta_{pE3-42}$  rapidly converted into soluble oligomeric forms in contrast to full-length  $A\beta_{1-42}$ , which stay in equilibrium for a longer time between monomers and oligomers (Bouter et al., 2013).

Parodi-Rullan et al. (2020) studied the effect of full-length and different N-terminal truncated A $\beta$  variants on bloodbrain barrier permeability, cerebral microvascular endothelial cell viability, and angiogenesis. The authors demonstrated that A $\beta_{4-42}$  followed by A $\beta_{4-40}$  was the most potent inhibitor of angiogenesis, and they were also much stronger than A $\beta_{1-42}$  and A $\beta_{1-40}$ .

## CHRONIC EFFECT OF N-TRUNCATED $A\beta_{4-42}$

The development of the APP/PSEN-1 double-transgenic mouse model, APP/PS1KI, for AD revealed, besides massive neuron loss in the hippocampus, many N-truncated  $A\beta_{X-42}$  variants including  $A\beta_{4-42}$  elucidated by two-dimensional Western blotting, which were subsequently verified by mass spectrometry (Casas et al., 2004). The 5XFAD mouse model is more widely used in the scientific community expressing mutant APP and PSEN-1 transgenes (Oakley et al., 2006). Mass spectrometric analysis with N-terminal specific A $\beta$  antibodies of 5XFAD mouse brain elucidated that the vast majority of A $\beta$  peptides were full-length A $\beta_{1-42}$  (Wittnam et al., 2012).

Using a highly specific antiserum against the N-terminus of  $A\beta_{4-X}$  abundant plaque staining was observed in APP/PS1KI and 5XFAD transgenic mouse brain (Wirths et al., 2017). In the human AD brain, this antiserum demonstrated staining of plaque cores of senile plaques, but none in diffuse amyloid deposits. The peptide content of plaques from AD brain and an AD mouse model (presenilin-2/APP transgenic mice, PS2APP) was analyzed using laser dissection microscopy combined with mass spectrometry (Rufenacht et al., 2005). The authors described various N-terminal truncated A $\beta$  peptides in PS2APP and AD amyloid plaques however with significantly elevated levels in AD brain.

Kawarabayashi et al. (2001) studied another AD mouse model with mutant APP (Tg2576) and elucidated that only a minor fraction of A $\beta$  was N-terminally truncated in contrast to AD brain (Kawarabayashi et al., 2001). This was verified, by another study by Kalback et al. (2002). The authors employed size-exclusion and reverse-phase chromatography, amino acid sequencing, and mass spectrometry of amyloid plaques of Tg2576 mice. The authors found that the amyloid plaques differed in their physical and chemical properties from those isolated from the AD brain. In Tg2576 mice, most peptides were full-length A $\beta_{1-42}$ , whereas, in AD brains, most peptides were N-truncated. The brain tissue of the transgenic mouse models mentioned above were freshly prepared without any post-mortem delay, therefore changes in pH, long-storage artifacts, state of agony, and medication, etc. can be ruled out as an explanation of the appearance of N-truncated A $\beta$  variants in transgenic mouse brain.

The Tg4-42 mouse model for AD expresses only  $A\beta_{4-42}$ , which represents a unique model system for studying the effect of chronic exposure of  $A\beta_{4-42}$  in the mouse brain (Bouter et al., 2013; Figure 2). The long-lasting exposure of  $A\beta_{4\text{-}42}$ induced an age-dependent neuron loss in the hippocampus, which correlated with hippocampus-dependent spatial reference memory deficits. Tg4-42 mice demonstrated synaptic hyperexcitability, changes in short-term synaptic plasticity but no effects on short- and long-term potentiation in the hippocampus (Dietrich et al., 2018). Busche et al. (2012) have demonstrated that full-length  $A\beta$  induced neuronal hyperactivity in brain slice cultures of wild-type mice. Synaptic hyperactivity of hippocampal pyramidal neurons is therefore an early event in AD pathology. <sup>18</sup>F-Fluorodeoxyglucose (<sup>18</sup>F-FDG)-PET in combination with magnetic resonance imaging (MRI) in Tg4-42 mice was used for analyzing cerebral brain glucose metabolism in vivo (Hinteregger et al., 2021). Tg4-42 mice demonstrated lower glucose uptake correlating with neuron loss and memory deficits in an age-dependent manner (Bouter et al., 2018). The reduction of glucose metabolism was detected already in young Tg4-42 prior to neuron death and neurological deficits. In clinical settings, the quantification of brain glucose uptake using <sup>18</sup>F-FDG-PET is an established diagnostic tool widely used for differential diagnosis of patients with dementia, including AD (Chetelat et al., 2020). The expert panel suggested a diagnostic algorithm with appropriate time-points using amyloid-PET and <sup>18</sup>F-FDG-PET for better clinical management of patients with AD. <sup>18</sup>F-FDG-PET signal is a powerful tool to monitor cerebral glucose consumption in vivo, a measure for synaptic activity (Sokoloff, 2008). It is also a valuable tool in preclinical research (Bouter and Bouter, 2019). A comparison between 5XFAD and Tg4-42 showed, that only young Tg4-42 developed robust neurological deficits, whereas in aged mice both models elicited similar memory deficits and fear conditioning tasks (Bouter et al., 2014). Reduced acoustic startle response, prepulse inhibition, and motor coordination were reported in Tg4-42 (Sichler et al., 2019; Wagner et al., 2019). Differentially expressed mRNAs in the brain of Tg4-42 and 5XFAD were identified by deep sequencing. A significant number of mRNAs were associated with memory deficits and neuron loss, which points to common disease pathways in both AD models (Bouter et al., 2014). Moreover, small RNA sequencing of the microRNAome in the hippocampus of Tg4-42 mice revealed microRNAs involved in learning, memory function, and synaptic signaling (Bouter et al., 2020). Metabolic changes of the glutamate/4-aminobutyrate-glutamine axis correlated with neurological deficits, neurodegeneration, and elevated CSF levels of neurofilament light chain in aged Tg4-42 mice (Hinteregger et al., 2021). Tg4-42 mice harbor more than 20 copies of the transgene in exon 2 of the retinoic acid receptor  $\beta$  (RARB) leading to decreased expression of RARB, which could have an (at least partial) effect on the phenotype



consequence of chronic exposure of A<sub> $\beta_{4-42}$ </sub> (4), massive reduction in glucose metabolism is detected by <sup>18</sup>F-PET/magnetic resonance imaging (MRI), loss of degenerating CA1 pyramidal neurons, and loss of spatial reference memory analyzed by the Morris water maze test (starting at 4–6 months of age). The figure shows methodologies in addition to pathological events using appropriate symbols. Created with BioRender.com.

(Hinteregger et al., 2020). Co-expression of A $\beta_{4-42}$  with A $\beta_{pE3-42}$  within the same neuron accelerated neurodegeneration in the hippocampus, enhanced loss of anxiety and motor deficits as well as sensori-motor deficits (Lopez-Noguerola et al., 2018).

## N-TRUNCATED $A\beta_{4-X}$ AS THE TARGET FOR IMMUNOTHERAPY

The discovery by Schenk et al. (1999) represents an important development in the AD field because it was the first time that disease-modulation was shown to be possible. The authors

showed that pre-aggregated synthetic  $A\beta_{1-42}$  drastically reduced amyloid plaques together with associated astrogliosis in the PDAPP transgenic mouse model for AD. This observation was supported by a subsequent report by Morgan et al. (2000) using another transgenic mouse model for AD (a cross of the Tg2576 and the PS1M146L transgenic line (Holcomb et al., 1998), demonstrating that memory decline assessed by the radial-arm water-maze test was stabilized by active immunization with full-length A\beta.

Janus et al. (2000) observed similar treatment effects of vaccination with  $A\beta_{1-42}$  in the TgCRND8 model. This was

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followed by exploring the effect of active immunization with AN1792 (pre-aggregated full-length  $A\beta_{1-42}$ ) in patients with AD. In phase I and IIa clinical trials a subset of the patients developed aseptic meningoencephalopathy and was therefore discontinued despite some hints that it might stabilize cognitive decline in a small subset of patients (Hock et al., 2002; reviewed in Morgan, 2011). Boche et al. (2008) demonstrated that amyloid plaques are solubilized by anti-A $\beta$  antibodies after active immunization with pre-aggregated full-length  $A\beta_{1-42}$ . Consequently, the solubilized amyloid- $\beta$  is drained *via* the perivascular pathway and was found to be increased in the brain.

Passive immunization with antibodies against AB had beneficial treatment effects in AD mouse models and is, therefore, another promising approach to modulating amyloid pathology in vivo. Demattos et al. (2001) used the monoclonal antibody m266 directed against the central domain of AB for passive immunization of PDAPP mice leading to reduced amyloid load. Stabilizing memory deficits using an object recognition task and a holeboard learning and memory task without a treatment effect on amyloid load in PDAPP mice was also reported (Dodart et al., 2002). However, Solanezumab the humanized version of m266 did not reach clinical endpoints on cognitive or functional abilities in phase III clinical trials with AD patients (Doody et al., 2014). In clinical phase III trials, bapineuzumab, the humanized monoclonal antibody derived from murine 3D6 was directed against Aβ<sub>1-5</sub> (Johnson-Wood et al., 1997) and did not improve clinical outcomes in patients with AD (Salloway et al., 2014). A comprehensive review of other antibody candidates for the treatment of AD has entered clinical trials and novel drug targets have recently been discussed in detail by Cummings et al. (2020).

Mclaurin et al. (2002) reported that active immunization with the full-length  $A\beta_{1-42}$  peptide of TgCRND8 mice reduced amyloid plaques and rescued memory deficits. Interestingly, the therapeutically active antibodies in this experimental setup recognized residues 4–10 of  $A\beta_{1-42}$ , an indication that N-truncated  $A\beta_{4-X}$  may be a relevant drug target. The therapeutic active anti- $A\beta_{4-10}$  antibodies also influenced

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aggregation propensity and toxicity *in vitro*. We have generated novel murine monoclonal A $\beta$  antibodies against N-truncated A $\beta_{4-X}$  (and A $\beta_{pE3-X}$ ) using preparations of freshly prepared A $\beta_{4-40}$  and used the antibody NT4X for further studies (Antonios et al., 2013, 2015). NT4X rescued the acute toxic effects of A $\beta_{4-42}$  using rat primary neurons and *in vivo* by cerebroventricular injection into wildtype mice (Antonios et al., 2013). NT4X also rescued the chronic effects of A $\beta_{4-42}$  on pyramidal neuron loss in the hippocampus and spatial reference memory deficits after passive immunization (Antonios et al., 2015). Therefore, besides A $\beta_{pE3-42}$ , A $\beta_{4-42}$  might be another relevant drug target in AD.

The question of whether N-truncations of AB within plaques represent a post-mortem artifact or might even precede the symptomatology of AD was addressed by Russo et al. (1997), demonstrating that both  $A\beta_{1-X}$  and  $A\beta_{pE-X}$  can form stable water-soluble aggregates not related to aggregates amyloid within plaques. Moreover, Rijal Upadhaya et al. (2014) studied post-mortem brain tissue from AD cases with symptomatic and preclinical AD by Western blot and demonstrated that  $A\beta_{pE3-X}$ is an informative biomarker for biochemical amyloid- $\beta$  staging.  $A\beta_{pE-X}$  and a phosphorylated A $\beta$  variant were not only detectable in plaques but also in soluble aggregates. The authors concluded that the different  $A\beta$  variants occur in a hierarchical sequence that allows the distinction of three stages, and may therefore be relevant for therapeutic intervention (Rijal Upadhaya et al., 2014). Finally, Mintun et al. (2021) conducted a phase 2 clinical trial of Donanemab in patients with early symptomatic AD, an antibody that specifically detects  $A\beta_{pE3-X}$  in plaques. The patients were selected based on the amount of tau and amyloid deposition on positron-emission tomography. The outcome of the study showed that a group of patients had a significantly better cognitive score than the placebo group as well as lower amyloid and tau load.

### **AUTHOR CONTRIBUTIONS**

TB wrote the article and designed the figures.

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