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Identification of isoAsp7-Aβ as a major Aβ variant in Alzheimer's disease, dementia with Lewy bodies and vascular dementia

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Abstract

The formation of amyloid- β (A β) aggregates in brain is a neuropathological hallmark of Alzheimer's disease (AD). However, there is mounting evidence that A β also plays a pathogenic role in other types of dementia and that specific post-translational Aβ modifications contribute to its pathogenic profile. The objective of this study was to test the hypothesis that distinct types of dementia are characterized by specific patterns of post-translationally modified A β variants. We conducted a comparative analysis and quantified A β as well as A β with pyroglutamate (pGlu3-A β and pGlu11-A β), N-truncation (A β (4-X)), isoaspartate racemization (isoAsp7-A β and isoAsp27-A β), phosphorylation (pSer8-A β and pSer26-A β) or nitration (3NTyr10-A β) modification in *post mortem* human brain tissue from non-demented control subjects in comparison to tissue classified as pre-symptomatic AD (Pre-AD), AD, dementia with Lewy bodies and vascular dementia. Aβ modification-specific immunohistochemical labelings of brain sections from the posterior superior temporal gyrus were examined by machine learningbased segmentation protocols and immunoassay analyses in brain tissue after sequential Aβ extraction were carried out. Our findings revealed that AD cases displayed the highest concentrations of all A β variants followed by dementia with Lewy bodies, Pre-AD, vascular dementia and non-demented controls. With both analytical methods, we identified the isoAsp7-A β variant as a highly abundant A β form in all clinical conditions, followed by A β (4-X), pGlu3-A β , pGlu11-A β and pSer8-A β . These A β variants were detected in distinct plaque types of compact, coarse-grained, cored and diffuse morphologies and, with varying frequencies, in cerebral blood vessels. The 3NTyr10-AB, pSer26-AB and isoAsp27-AB variants were not found to be present in A β plaques but were detected intraneuronally. There was a strong positive correlation between isoAsp7-A β and That phase and a moderate negative correlation between isoAsp7-A β and performance on the Mini Mental State Examination. Furthermore, the abundance of all A β variants was highest in APOE 3/4 carriers. In aggregation assays, the isoAsp7-A β , pGlu3-A β and pGlu11-A β variants showed instant fibril formation without lag phase, whereas A β (4-X), pSer26-A β and isoAsp27-A β did not form fibrils. We conclude that targeting A β post-translational modifications, and in particular the highly abundant isoAsp7-A β variant, might be considered for diagnostic and therapeutic approaches in different types of dementia. Hence, our findings might have implications for current antibody-based therapies of AD.

Keywords Amyloid- β · Abeta · Post-translational modifications · Alzheimer's disease · Vascular dementia · Dementia with Lewy bodies · Mini Mental State Examination · Automated histology quantification

Abbreviations

Αβ	Amyloid β
AD	Alzheimer's disease
APOE	Apolipoprotein E

Sarah Schrempel and Anna Katharina Kottwitz have contributed equally to this work.

Extended author information available on the last page of the article

ARIA Amyloid-related imaging abnormality CAA Cerebral amyloid angiopathy DAB 3,3'-Diaminobenzidine DLB Dementia with Lewy bodies FA Formic acid GdmCl Guanidinium chloride HFIP 1,1,1,3,3,3-Hexafluoro-2-isopropanol HRP Horseradish peroxidase

LB stage	Unified Staging System for Lewy Body
	Disorders
MMSE	Mini Mental State Examination
NOS2	Inducible nitric oxide synthase
PBS	Phosphate-buffered saline
PBS-T	PBS containing 0.02% Tween 20
PMI	Post mortem Interval
Pre-AD	Pre-symptomatic AD
PTM	Post-translational modifications
ROI	Region of interest
TBS	Tris-buffered saline
TEM	Transmission electron microscopy
ThT	Thioflavin T
TMB	3,3',5,5'-3,3,5,5-Tetramethylbenzidin
VAD	Vascular dementia

Introduction

Alzheimer's disease (AD) and other types of dementia are characterized by the degeneration of defined subsets of neurons and by the deposition of proteins that accumulate as amyloid plaques, neurofibrillary tangles, Lewy bodies and glial cytoplasmic inclusions. Historically, the aggregated proteins were assigned to defined clinical conditions: AB and Tau to AD [19, 37], α -synuclein to the synucleinopathies Parkinson's disease (PD) [86, 110], dementia with Lewy bodies (DLB) [9, 110] and multiple system atrophy (MSA) [5, 116] and Huntingtin to Huntington's disease (HD) [64, 102]. This constricted perspective has changed in the last two decades when proteins formerly considered typical for a specific clinical condition were also observed in other types of pathology [4, 6, 14, 34, 42, 53, 117, 118]. Therefore, we became interested in analyzing the pathology of distinct Aß peptide variants in brains from different types of dementia: AD, DLB and vascular dementia (VAD) and, at an early disease stage, Pre-AD.

It is well established that $A\beta$ per se is a physiological peptide with cellular functions. For example, it was shown to be critical for neuronal survival [82]. Moreover, A β acts as a positive modulator of neurotransmitter release probability in hippocampal synapses [1] and its hippocampal production is enhanced during memory induction in experimental mouse models [84]. In addition, A β concentrations measured by ELISA in brain interstitial fluid have been shown to correlate with neurological status in human subjects [20]. On one hand, in the clinical condition of familial AD, amyloid precursor protein overexpression and mutations or mutations of its processing γ -secretase, respectively, lead to pathological Aβ accumulation [87, 121, 126]. In sporadic AD, on the other hand, the exact mechanisms leading to Aß accumulation are not known, but may include a pathological formation of A β post-translational modifications (PTMs), such



Fig. 1 Schematic representation of the A β peptide variants investigated in the present study in brain tissue of deceased patients categorized in Pre-AD, AD, DLB, VAD and control subjects. Together, eight different A β variants from five groups of post-translational modifications were analyzed side-by-side in a comprehensive manner by immunohistochemical and biochemical methods. A β structure from [25]. *Created with BioRender.com*

as N-terminal truncation [17, 66], pyroglutamylation [63, 97], phosphorylation [55, 57], nitration [61] and isoaspartate formation [107, 115] (for review, see [62, 91]). Such $A\beta$ PTMs might be valuable diagnostic markers and also therapeutic targets for pharmacologic interventions. In case of enzyme-catalyzed generation of PTMs, their formation could be prevented by enzyme inhibition and already formed $A\beta$ PTMs could be targeted by specific monoclonal antibodies. A good example for this strategy is the prevention of pGlu3-A β formation by inhibition of glutaminyl cyclase with Varoglutamstat [100, 119] and clearance of existing pGlu3-A β by the antibody Donanemab [40, 108].

However, to date, there is no comprehensive side-by-side analysis of all the above-mentioned A β variants in different types of dementia, which would allow the identification of a therapeutic target for a specific type of dementia or a combination of dementia diseases. Therefore, we here asked the question whether the different clinical conditions of dementia under investigation are characterized by specific signatures of A β PTMs or display similar patterns of these A β variants (Fig. 1). To address this issue, we used wellpreserved human brain tissue with a short *post mortem* delay and a detailed clinical and pathological characterization that enabled us to correlate clinical with histopathological and biochemical findings.

Materials and methods

The general workflow of the immunohistochemical and biochemical analyses conducted in the present study is summarized in Fig. 2. Human brain tissue from all cases was available to be used for both analytical methods.

Human brain tissue

The human brain tissue was provided by the Arizona Study of Aging and Neurodegenerative Disorders (AZSAND) and Brain and Body Donation Program of Banner Sun Health Research Institute in Sun City, Arizona. Case recruitment and autopsy were performed as approved by institutional review boards overseeing AZSAND's Brain and Body Donation Program [13]. The required consent was obtained for all cases. The diagnosis and staging of AD, DLB and VAD for the cases used in this study were based on standardized clinicopathological criteria as previously described [13]. The diagnosis of AD cases was based on the presence of neurofibrillary tangles and neuritic plaques in the hippocampal formation and neocortical areas and met "intermediate" or "high" levels of AD Neuropathological Change according to the criteria of the National Institute on Aging-Alzheimer's Association (NIA-AA). DLB was defined as dementia occurring either at presentation or within one year of the onset of parkinsonism, with a brain distribution of α -synuclein pathology meeting DLB Consortium criteria for "intermediate" or "high" likelihood [67]. The Unified Staging System and McKeith criteria received good inter-rater reliability scores in a multi-center comprehensive analysis defining consensus criteria for the evaluation of Lewy body pathology in *post mortem* brains [8]. The diagnosis of VAD was based on the National Institute of Neurological Disorders and Stroke (NINDS) and the Association Internationale pour la Recherche et l'Enseignement en Neurosciences (AIREN) criteria [92].

Brain sections of the posterior superior temporal gyrus (Brodmann area 22) from ten AD cases, ten DLB cases, ten VAD cases and thirty age-matched non-demented subjects were evaluated for deposition of different post-translationally modified A β variants by immunohistochemistry. According to their respective Thal phase, the thirty age-matched non-demented subjects were further subdivided into control (Co; Thal 0–1) and pre-symptomatic AD (Pre-AD; Thal 2–5). Biochemical analyses were performed in unfixed brain tissue from the same location (Brodmann area 22) of the other brain hemisphere of the same cases. For detailed characterization of the cases, see Table 1.

Antibodies

For immunohistochemical labelings and immunoassay analyses, primary antibodies specific for the respective A β variants were used (for details, see Table 2). Antibodies raised against A β (3D6; epitope A β amino acids 1 to 5; parent antibody of Bapineuzumab), pGlu3-Aβ (J8), isoAsp7-Aβ (K11), (F2) were generated by Fraunhofer IZI-MWT (Halle, Germany) (see Supplementary Information). The specificities of the antibodies 3D6 against A β [32, 38], J8 against pGlu3-A β [45], K11 against isoAsp7-Aβ [38], 1E4E11 against pSer8-Aβ [56], K13 against pGlu11-Aβ [52] and 5H11C10 against pSer26-A β [59] were recently demonstrated. For the characterization of the binding capacity and specificity of the newly generated monoclonal antibodies 4C3 raised against 3NTyr10-Aβ and F2 against isoAsp27-Aβ, immunoassays based on different N- or C-terminally truncated



Fig. 2 Schematic presentation of the workflow for the immunohistochemical (top) and biochemical (bottom) analyses of A β peptide variants in human brain tissue from Pre-AD, AD, DLB, VAD and control subjects. The immunohistochemically labeled brain slices were digitized with an Axioscan slide scanner and subjected to plaque load quantification by machine learning-based segmentation protocols.

For biochemical analyses, $A\beta$ peptides were extracted from unfixed human brain tissue by sequential centrifugation of Tris-buffered saline (TBS), guanidinium chloride (GdmCl) and formic acid (FA) dilutions followed by quantification by immunoassays using specific monoclonal antibodies. *Created with BioRender.com*

A β species as well as wild type A β were established (see Suppl. Figure 1 and 2). For the detection of A β (4-X), the polyclonal rabbit antibody 58–1 was used for immunohistochemical labeling [10] and the monoclonal antibody 18H6 was employed for immunoassay analyses [22]. In immunoassay analyses, the commercially available horseradish peroxidase (HRP)-conjugated antibody 4G8 (Biolegend, San Diego, CA, USA) against A β (17–24) and HRP-conjugated antibody A β x-40 (clone 11A50B10; Biolegend) against A β (x-40) have been used for the detection of A β and A β 40 species, respectively.

Peptide synthesis

Peptides listed in Suppl. Table 1 were purchased from peptides & elephants GmbH (Hennigsdorf, Germany), Bio-Cat GmbH (Heidelberg, Germany) or Peptide Specialty Laboratories GmbH (Heidelberg, Germany). Generation of full-length and N- or C-terminally truncated pGlu3-A β , 3NTyr10-A β and isoAsp27-A β peptides was performed by Fraunhofer IZI as previously described by Piechotta et al. [80] and Gnoth et al. [38].

Immunohistochemical analysis

Single labeling immunohistochemistry

All immunohistochemical labelings on human brain tissue were performed on 40 µm thick free-floating brain sections at room temperature, unless stated otherwise. After optional pre-treatment (for pre-treatments, see Table 2), brain sections were washed in PBS containing 0.02% Tween 20 (PBS-T) and treated with 1% H₂O₂ in 60% methanol for 60 min. Unspecific staining was blocked by incubating brain sections in blocking solution (PBS-T with 2% (w/v) bovine serum albumin, 0.3% (*w/v*) milk powder, 0.5% (*v/v*) normal donkey serum) for 60 min before incubating brain sections with the primary antibodies in blocking solution at 4 °C for 42 h (for antibody concentrations, see Table 2). Brain sections were incubated in a 1:2 mixture of blocking solution and PBS-T containing the biotinylated donkey anti-mouse or anti-rabbit, respectively, secondary antibody (1:1000; Dianova, BIO-ZOL Diagnostica Vertrieb GmbH, Eching, Germany) for 60 min followed by incubation with ExtrAvidin-conjugated peroxidase (1:2000; Sigma, Merck KGaA, Darmstadt, Germany) in PBS-T for 60 min. Bound peroxidase was visualized in a solution containing 2-4 mg DAB, 40 mg ammonium nickel(II) sulfate and 5 µl H₂O₂ per 10 ml Tris-buffer (0.05 M; pH 8.0) yielding black epitope staining.

Light microscopy

Immunohistochemically stained human brain sections were digitized with an Axio-Scan.Z1 slide scanner connected with a LED light source and a Hitachi HV-F202SCL camera (Carl Zeiss AG, Oberkochen, Germany). Using a $20 \times$ objective lens with 0.8 numerical aperture (Carl Zeiss AG), high-resolution images from areas of interest were taken. Images were analyzed using the Zeiss ZEN 3.8 imaging tool. Photoshop CS2 (Adobe Systems, San José, CA, USA) was used to process the images. Care was taken to apply the same brightness, sharpness, color saturation and contrast adjustments in the processing of the various pictures.

Automated image analysis

To quantify the stained plaque area of immunohistochemically labeled brain sections, Zeiss arivis cloud and Zeiss ZEN 3.8 Intellesis software were used for automated image analysis (for details see Suppl. Figure 3). For each A β variant and for all 60 cases, three representative regions of interest (ROIs) with an area of 4 mm² each were chosen of the grey matter spanning all cortical layers. A separate training was conducted for each A β variant staining using Zeiss arivis cloud. After several iterating training cycles to enhance the capability of the machine learning algorithm to detect immunohistochemically stained plaques, all 60 cases were analyzed by the trained algorithm in a batch analysis using Zeiss ZEN Intellesis software. Obviously false positive detection of unspecific staining was manually removed upon visual inspection.

Staging of plaque morphology and vascular deposition

The staging of compact, coarse-grained, cored and diffuse plaques stained for a specific A β variant was performed on digitized human brain slices with Zeiss ZEN 3.8 according to Boon et al. [15]. The plaque load of each plaque morphology was ranked between no staining (–), some positive staining (+ –), positive staining (+) and prominent positive staining with high plaque load (++). Vascular A β deposition was evaluated similarly and differentiated between 0% (–), 1–40% (+ –), 41–80% (+) and 81–100% (++) of cases with vascular deposition per clinical group.

Case #	Age [years]	[h] [M]	Gender f / m	Brain weight [g]	CAA yes / no	Braak score	Thal phase	APOE genotype	MMSE score	CERAD criteria	NIA-AA criteria	Uni- fied LB stage
Co												
Co1	88	3	f	1030	ou	II	1	3/4	n.a	not AD	not met	0
Co2	82	2.16	ш	1160	yes	III	1	3/3	28	not AD	not met	0
Co3	91	1.5	ш	1440	no	П	0	3/3	27	not AD	not met	0
Co4	53	3.66	ш	1456	no	I	0	3/3	n.a	not AD	not met	0
Co5	61	2.33	ш	1220	yes	I	1	3/3	n.a	not AD	not met	0
Co6	59	3.15	f	1162	ou	I	1	3/3	n.a	not AD	not met	0
Co7	91	2.5	f	1185	yes	III	1	2/2	27	not AD	not met	0
Co8	74	3.25	в	1440	no	Ι	0	2/3	n.a	not AD	not met	0
Co9	82	2.07	f	1249	yes	I	1	2/3	29	not met	not met	0
Co10	06	5.42	в	1239	no	III	0	3/3	22	not met	not met	0
Co11	79	4.3	в	1156	no	П	0	2/3	28	not met	not met	0
Co12	66	3.5	f	975	no	Ш	0	3/3	29	not AD	not met	0
Co13	90	3.41	В	1150	no	I	1	3/3	27	not met	not met	0
Co14	87	3.08	f	1001	no	III	0	3/4	28	not met	not met	0
Co15	87	4	f	1196	yes	Ш	1	3/4	28	not met	not met	0
Co16	72	4.6	ш	1262	yes	I	0	3/3	29	not met	not met	0
Co17	71	3.5	ш	1344	no	Ш	0	3/3	27	not met	not met	0
Co18	100	3.87	ш	1274	no	III	0	n. a	20	not met	not met	0
Co19	76	2.5	В	1340	no	Ι	0	3/3	n.a	not AD	not met	0
Co20	79	ю	В	1100	ou	I	0	3/3	n.a	not AD	not met	0
Mean	80.55	3.24	7/13	1218.95	6/20	II	0		26.85			0
Pre-AD												
Pre-AD1	81	2.75	В	1190	yes	Ш	5	3/3	25	not AD	not met	0
Pre-AD2	90	3	f	1150	ou	III	ς,	3/3	27	not AD	not met	0
Pre-AD3	86	3	ш	1055	no	I	2	3/3	n.a	not AD	not met	0
Pre-AD4	95	2.66	f	1140	no	Ш	2	3/3	n.a	not AD	not met	0
Pre-AD5	73	2.5	ш	1240	no	П	2	2/3	30	not AD	not met	0
Pre-AD6	87	4	f	1145	no	III	2	3/3	n.a	not AD	not met	0
Pre-AD7	53	3.95	f	1015	no	П	2	3/3	n.a	not met	low	0
Pre-AD8	81	c,	f	1275	yes	П	3	4/4	n.a	not AD	not met	0
Pre-AD9	<i>6L</i>	2	ш	1135	no	Ι	2	3/3	n.a	not AD	not met	0
Pre-AD10	70	2	f	1100	yes	I	2	4/4	n.a	not AD	not met	0
Mean	79.50	2.89	6/4	1144.5	3/10	Π	7		27.33			0
AD												

Table 1 (con	ttinued)											
Case #	Age [years]	[h] [M]	Gender f / m	Brain weight [g]	CAA yes / no	Braak score	Thal phase	APOE genotype	MMSE score	CERAD criteria	NIA-AA criteria	Uni- fied LB stage
AD1	76	3.08	f	862	yes	>	n.a	3/3	9	definite AD	high	0
AD2	92	3.73	f	1110	yes	Ν	5	3/4	7	definite AD	high	0
AD3	75	3.83	Е	1070	yes	>	5	3/4	3	definite AD	high	0
AD4	77	3.62	н	1180	yes	Ν	5	3/4	10	definite AD	high	0
AD5	83	4	Е	1184	yes	N	5	3/4	14	definite AD	high	0
AD6	83	3.62	f	1052	yes	>	5	3/3	17	definite AD	high	0
AD7	84	3.33	f	1142	yes	ΙΛ	5	3/3	24	definite AD	high	0
AD8	83	3.58	в	1105	yes	Λ	5	2/4	3	definite AD	high	0
AD9	75	2.97	ш	1251	yes	^	5	3/4	25	definite AD	high	0
AD10	82	4	f	1098	yes	ΙΛ	5	3/3	19	definite AD	high	0
Mean	81.00	3.58	5/5	1105.4	10 / 10	٨	S.		12.80			0
DLB												
DLB1	78	3.33	ш	1300	no	I	3	3/3	n.a	probable AD	intermediate	N
DLB2	80	2.33	f	1080	no	III	4	3/4	0	probable AD	intermediate	N
DLB3	87	3.75	ш	1160	no	Π	0	2/3	22	not AD	not met	III
DLB4	74	2.42	ш	1422	no	Π	3	3/3	6	definite AD	intermediate	N
DLB5	76	3.47	f	1066	no	IV	0	3/3	27	not met	not met	IV
DLB6	68	2.5	ш	1350	no	Ι	ю	3/4	26	definite AD	intermediate	IV
DLB7	96	3.16	f	1195	no	Ш	0	3/3	n.a	not AD	not AD	III
DLB8	92	5	ш	1200	yes	IV	4	3/4	20	probable AD	intermediate	IIb
DLB9	78	3	ш	1225	ou	Ι	5	2/3	n.a	probable AD	intermediate	IV
DLB10	74	3.33	ш	1250	yes	Ш	2	3/3	16	not AD	low	N
Mean	82.40	3.23	3/7	1224.8	2/10	II	2		17.14			N
VAD												
VAD1	86	.0	ш	1120	no	III	0	3/3	20	not AD	not met	0
VAD2	106	2.25	f	896	yes	IV	0	2/3	19	not met	not met	0
VAD3	89	2.83	ш	1158	no	Ш	0	3/3	27	not met	not met	0
VAD4	85	Э	f	1080	ou	Ι	n.a	3/3	20	not AD	not AD	0
VAD5	86	2.97	f	1082	ou	IV	0	3/3	12	not met	not met	0
VAD6	92	3.32	f	1070	no	IV	0	3/3	27	not met	not met	0
VAD7	93	2.92	ш	1015	ou	Π	ŝ	2/3	19	definite AD	low	0
VAD8	92	2.63	f	1075	no	IV	0	3/3	28	not met	not met	0
VAD9	88	2	f	1170	yes	Ш	2	3/4	24	possible AD	low	0
VAD10	82	3.25	ш	1290	no	Ш	0	3/4	2	not AD	not AD	0
Mean	89.90	2.82	6/4	1095.6	2/10	III	1		19.80			0

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Bold represents mean values of the disease conditions

of AD based on criteria of the Consortium to establish a registry for AD; NIA-AA criteria diagnosis of AD based on criteria of the National Institute on Aging–Alzheimer's Association; Unified 113]; APOE genotype apolipoprotein E genotype; MMSE score last Mini Mental State Examination score before death; CERAD criteria diagnosis IIa (brainstem predominant), IIb (limbic predominant), III (brainstem and liminterval; f female; m male; CAA Cerebral Amyloid Angiopathy; Braak score Braak neurofibrillary stage I-VI [18]; Thal phase Thal amyloid staging with phases 0 (A0), LB stage unified staging system for Lewy body disorders with stage 0 (no Lewy bodies), I (olfactory bulb only), and 5 (A3) [112, (A1), 2 (A1), 3 (A2), 4 (A3) bic), IV (neocortical) [12] PMI post mortem

Immunoassays

Human brain tissue preparation

Tissue blocks of human temporal cortex were prepared in the frontal plane according to the atlas of the human brain and stored at - 80 °C until usage. To prepare human brain for A β immunoassay analysis, the protocol of Gnoth et al. [38] was followed with adaptations. In short, a sample of about 500 mg temporal cortex was homogenized in TBS buffer (20 mM Tris/HCl. 150 mM NaCl. 5 mM KCl. pH 7.5) supplemented with Protease Inhibitor Cocktail Tablets (Roche Diagnostics GmbH, Mannheim, Germany) and 1 mM 4-(2-Aminoethyl)-benzolsulfonylfluorid Hydrochlorid (Carl Roth GmbH+Co. KG, Karlsruhe, Germany) at a concentration of 200 mg/ml by using a Precellys homogenizer (VWR International GmbH, Darmstadt, Germany), followed by sonication for 10 s. After centrifugation of the homogenate at 100,000xg for 1 h, supernatant (TBS fraction) was obtained. The resulting pellet was dissolved to 250 mg/ml in 5 M GdmCl, followed by an incubation step in an overhead shaker at room temperature for 3 h and a subsequent centrifugation step at 100,000xg for 1 h. Again, supernatant (GdmCl fraction) was collected and the pellet was resuspended in 500 µl 70% FA, followed by sonication for 20 s and neutralization by 2.9 ml 3.5 M Tris (FA fraction).

Quantification of A β variants using immunoassay analysis

The performance of A β and isoAsp7-A β immunoassay analyses was previously described by Gnoth et al. [38]. In short, coating antibodies were immobilized on polystyrene 96-well microtiter plates at 4 °C overnight. After blocking (4 °C, 2 h), sample dilutions (TBS fraction 1:2-1:50, GdmCl fraction 1:40-1:5000, FA fraction 1:100-1:5000) were selected according to subsequent adsorption signals within the linear range of the standard curve. For the standard curve, synthetic standard peptides were serially diluted and added to the wells in duplicate. This was followed by an incubation period at 4 °C for 2 h. For detection of bound A β peptides, the HRP-conjugated antibody 4G8 was used to detect Aβ, pGlu3-, isoAsp7-, pSer8- and 3NTyr10-Aβ. The HRP-conjugated antibody Aβx-40 was used for pSer26- and isoAsp27-Aß detection. Both detection antibodies were diluted to a final concentration of $1 \mu g/$ ml (see Table 2), added to the samples and incubated at 4 °C for 1 h. A color reaction with commercially available HRP substrate 3,3',5,5'-3,3,5,5-Tetramethylbenzidine (TMB; SureBlue Reserve TMB Microwell Peroxidase Substrate (1-component); KPL, LGC Clinical Diagnostics, Milford, MA, USA) was performed and stopped by the addition of 1.2 N H₂SO₄. Absorption at 450/540 nm was determined Table 2 Primary and secondary antibodies for detection of A^β variants by immunohistochemistry (IHC) and immunoassay analyses

IHC and Immunoassay			IHC		Immunoassay				
Antibody (Clone; Concentration [mg/ml])	Company / Supplier	Host	Pre-treatment	Antibody Dilution	Standard Curve Range or Concentration [ng/ ml]				
Aβ (3D6; 5.00)	Fraunhofer IZI (Halle, Germany)	Mouse	_	1:4000	0.39–0.006				
pGlu3-Aβ (J8; 1.00)	Fraunhofer IZI (Halle, Germany)	Mouse	basic	1:1000	0.39-0.006				
Aβ(4-X) (58–1; 1.00)	University Medical Center Göttin- gen (Göttingen, Germany)	Rabbit	-	1:5000	-				
Aβ(4-X) (18H6; 0.60)	NYU Grossman School of Medi- cine (New York, USA)	Mouse	_	-	30-0.0073				
isoAsp7-Aβ (K11; 3.00)	Fraunhofer IZI (Halle, Germany)	Mouse	_	1:6000	0.5–0.008				
pSer8-Aβ (1E4E11; 1.00)	University Hospital Bonn (Bonn, Germany)	Mouse	acidic	1:1000	3.125-0.049				
pGlu11-Aβ (K13; 2.50)	Fraunhofer IZI (Halle, Germany)	Mouse	_	1:1000	-				
3NTyr10-Aβ (4C3; 1.00)	Fraunhofer IZI (Halle, Germany)	Mouse	_	1:400	6.25-0.01				
pSer26-Aβ (5H11C10; 1.00)	University Hospital Bonn (Bonn, Germany)	Mouse	acidic	1:2000	2.187-0.003				
isoAsp27-Aβ (F2; 1.64)	Fraunhofer IZI (Halle, Germany)	Mouse	_	1:6000	100-1.56				
biotinylated donkey anti-mouse (-; 0.55)	Dianova, BIOZOL Diagnostica Vertrieb GmbH, (Eching, Ger- many)	Donkey	-	1:1000	-				
biotinylated donkey anti-rabbit (-; 0.50)	Dianova, BIOZOL Diagnostica Vertrieb GmbH, (Eching, Ger- many)	Donkey	-	1:1000	-				
HRP-conjugated 4G8 (4G8; 0.50)	Biolegend (San Diego, CA, USA)	Mouse	-	-	1000				
HRP-conjugated Aβx-40 (11A50B10; 0.50)	Biolegend (San Diego, CA, USA)	Mouse	_	-	1000				

Pre-treatments: basic: Tris-buffer (0.05 M; pH 8.0); acidic: citric acid/sodium citrate (0.1 M; pH 6.0) and 88% FA (v/v)

by a Tecan Sunrise plate reader (Tecan Group Ltd., Männedorf, Switzerland). The standard curve was calculated from measured absorption by a 4-Parameter-Logistic-Fit: $y = A2 + (A1 - A2)/(1 + (x/ \times 0)^p)$. Novel immunoassays were adapted and specified in the same way (see Table 2). For specification purposes, the stability of the antibodies in the matrix as well as LOD and LOQ were determined.

Quantification of the A β (4-X) variant was carried out as electrochemiluminescence assay (Meso Scale Discovery, MSD, Rockville, MD, USA). Biotinylated 18H6 ([22]; 0.6 µg/ml), as well as sulfo-tagged 4G8 (1:50; Meso Scale Discovery, MSD) were used as capture and detection antibodies, respectively, on MSD GOLD 96-well Small Spot Streptavidin Plates (Meso Scale Discovery, MSD) following the manufacturer protocol for the human A β antibody set. Standard curves were prepared using synthetic A β (4–40) peptides and data analysis including standard curve calculations was carried out with the Discovery Workbench 4.0.12 software package (Meso Scale Discovery, MSD).

Thioflavin T fibril formation assays

Aβ aggregation studies were based on Piechotta et al. [80]. Lyophilized Aβ peptides were dissolved in 1,1,1,3,3,3-hexafluoro-2-isopropanol (HFIP) to obtain homogeneous preparations without seeds. For peptide preparation, the HFIP was evaporated and immediately prior to analysis, peptide pellets were dissolved in 20 µl of 0.1 M NaOH and incubated for 10 min. After adding 380 µl PBS (138 mM NaCl, 8 mM Na₂HPO₄, 1.5 mM KH₂PO₄, 3 mM KCl, pH 7.1), neutralization with 0.1 M HCl followed. All work steps were carried out on ice. Peptide concentration was determined by Pierce BCA Protein Assay Kit (Thermo Fisher Scientific, Waltham, MA, USA) and adjusted to 10 µM or 15 µM. To monitor the fibril formation of A β variants, monomeric A β peptide was co-incubated with 20 μ M thioflavin T (ThT) in 96-well microtiter plates (PS, half area, black μ Clear; Greiner Bio-One, Kremsmünster, Austria). Measurements were carried out in triplicates. The plate was sealed with adhesive film and incubated at 37 °C and 300 rpm for 20 h. ThT fluorescence intensity was measured using a CLARIOstar (BMG Labtech, Ortenberg, Germany) plate reader (excitation 440 nm, emission 490 nm) and normalized to the initial ThT fluorescence signals. Kinetic parameters of the ThT curves of the fibrilforming A β variants were calculated using logistic sigmoid functions in GraphPad PRISM (version 10, San Diego, CA, USA).

Transmission electron microscopy

Transmission electron microscopy (TEM) of A β aggregates was based on Köppen et al. [53]. In short, fibril samples (10 µl) from an aggregation reaction as described above were directly incubated on an EM Tec Formvar Carbon TEM Support Film on nickel 200 square mesh (Micro to Nano, Haarlem, Netherlands) at room temperature for 10 min and washed three times with distilled water. Staining was obtained with 2% (v/v) uranyl acetate (SERVA Electrophoresis GmbH, Heidelberg, Germany) for 1 min. Fibrils were imaged with a LEO EM 912 Omega TEM (Carl Zeiss AG) at 80 kV, and digital micrographs were obtained with a dual-speed 2 K-on-axis CCD camera-based YAG scintillator and the software ImageSP (version 1.2.13.17 TRS-Tröndle, Moorenweis, Germany).

Statistical analysis of immunohistochemical and immunoassay data

For statistical analysis, the mean plaque area and the mean amount of A β variant was used. Using a Python script for immunohistochemical data, the total plaque area of one ROI of one case and one AB variant was calculated relative to the total area of the ROI (4 mm²). The mean plaque area was then calculated from the total plaque area of three ROIs per case. The mean amount of AB variant per case was calculated using two technical immunoassay replicates. In the case of the A β (4-X) electrochemiluminescence assay, a single measurement was performed. Determination of statistical significance of differences of the stained plaque areas and the amount of a specific A β variant, respectively, between disease conditions was conducted with One-Way ANOVAs followed by Tukey's multiple comparisons test using Graph-Pad PRISM. Differences between groups were considered statistically significant for p values < 0.05. Correlations of plaque area or amount of Aß variants relative to Braak stage, Thal phase and MMSE were calculated as Pearson Correlations using GraphPad PRISM. The Pearson correlation coefficient r and the P value (two-tailed, confidence interval 95%) are indicated in each plot. Pearson correlation was considered to be moderate with correlation coefficients of $0.40 \le r < 0.70$, as strong with $0.70 \le r < 0.90$ and as very strong with $0.90 \le r \le 1.00$. Correlations with $0.00 \le r < 0.40$ were interpreted as weak and negligible [105]. Heat Maps show the mean plaque areas, the mean A β variant amounts and the Pearson correlation coefficient r of the immunohistochemical or biochemical correlations, respectively (values are not normalized).

Results

Immunohistochemical analyses

First, the abundance of post-translationally modified Aß peptides in amyloid plaques was quantified. The Aβ variants 3NTyr10-A β , pSer26-A β and isoAsp27-A β were not detected in amyloid plaque-like formations and, thus, excluded from this analysis. The immunohistochemical appearance of these A β variants is documented in Suppl. Figure 4. All other A β variants were, in varying quantities, detected in amyloid plaques (Fig. 3). In control cases, Braak stage I-III and Thal phase 0-1, a very minor plaque load of < 1% of the grey matter brain area was calculated for all $A\beta$ peptide variants (Fig. 3). In the Pre-AD cases, Braak stage I-III and Thal phase 2-5, the median plaque load varied between 4.8% for Aβ and 0.3% for pGlu11-Aβ. Among the remaining A PTMs, the isoAsp7-A variant was most abundant (2.6% plaque load) followed by A β (4-X) (1.7% plaque load), pGlu3-Aß (1.5% plaque load) and pSer8-Aß (0.6% plaque load) (Fig. 3). Among all clinical conditions, the highest Aβ plaque load was detected for AD, Braak stage V-VI and Thal phase 5 (Fig. 3). For the different A β variants, median plaque load values between 10.7% for A β and 0.5% for pGlu11-Aβ were detected. The abundance of isoAsp7-Aβ (12.3% plaque load) was similar to that of A β and was followed by A β (4-X) (4.2% plaque load), pGlu3-A β (3.4% plaque load) and pSer8-A β (1.0% plaque load). Also in DLB, Braak stage I to IV and Thal phase 0-5, the isoAsp7-A β was most abundant among all A^β variants (3.9% plaque load), followed by pGlu3-A β (2.5% plaque load), A β (4-X) (1.4% plaque load) and pGlu11-AB and pSer8-AB with less than 1% plaque load each (Fig. 3). In the VAD cases, Braak stage I–IV and Thal phase 0-3, all A β variants accounted for less than 1% plaque load, each.

Together, the immunohistochemical analyses of plaque load revealed the highest abundance of all A β variants analyzed in AD, followed by DLB, Pre-AD and VAD, the latter two being close to control cases. Regarding the different A β variants, isoAsp7-A β plaque load was most abundant,





<Fig. 3 Representative examples of immunohistochemical labeling of human cortical brain tissue from control cases and different clinical conditions using antibodies to detect specific post-translational A β modifications as indicated (left). Respective quantifications of A β plaque load as a percentage of brain area covered by plaques is presented (right). Note that the Y-axes differ between the individual A β variants. Differences between clinical groups are statistically significant at *p < 0.05; **p < 0.01; ***p < 0.001. Medians are indicated by horizontal lines

followed by $A\beta(4-X)$ and pGlu3-A β at similar levels and pGlu11-A β and pSer8-A β with the lowest plaque occupancy.

We next asked whether there is an association of the $A\beta$ load with APOE genotype and whether there might be correlations between the abundance of the AB variants investigated and the histopathological Braak and Thal staging on one hand and the MMSE score on the other. The subgroup sizes defined by APOE genotype differed markedly with N=1 for the 2/2, N=8 for the 2/3, N=1 for the 2/4, N=34 for the 3/3, N=13 for the 3/4 and N=2 for the 4/4 allele (see Table 1). The highest abundance for all $A\beta$ variants was detected in APOE 3/4 carriers, followed by APOE 3/3 carriers (Fig. 4a). Within the APOE 3/4 subgroup, the mean plaque load for all AB PTMs was significantly lower than that of A\beta, except for isoAsp7-Aβ, which displayed similar levels. As expected, the A β variants showed a higher correlation with Thal amyloid phases than with Braak Tau stages (Fig. 4b). The strongest correlations with Thal phases were observed for A β , A β (4-X) and isoAsp7-A β (r values $0.70 \le r < 0.90$ indicate a strong correlation) and were highlighted in grey. Moderate correlations (r values $0.40 \le r < 0.70$) were marked with a light grey. Regarding the MMSE, we observed moderate negative correlations for A β , A β (4-X) and isoAsp7-A β (Fig. 4b).

To better visualize the plaque load for the individual $A\beta$ variants in the different types of dementia and the strength of correlations with Braak and Thal staging and MMSE scores, respectively, heat maps were established (Fig. 5). These indicate the particularly high abundance of isoAsp7-A β in Pre-AD, AD and DLB (Fig. 5a), as well as a positive correlation between isoAsp7-A β plaque load and histopathological Braak and Thal staging (Fig. 5b). In addition, a moderate correlation of isoAsp7-A β and A β (4-X) with the decline in MMSE is evident (Fig. 5b).

We also wanted to reveal whether specific A β variants are associated with amyloid plaque types of distinct morphology and with cerebral blood vessels. Therefore, we graded the abundance of each A β variant separately according to its respective detection in the following categories of plaques: compact, coarse-grained, cored and diffuse, as well as in blood vessels. This was done for all clinical conditions and cases (Fig. 6a). The staging was made according to Boon et al. [15] and differentiated between no staining (–), some positive staining (+ –), positive staining (+) and prominent positive staining with high plaque load (++). AD and DLB cases displayed a high abundance of all plaque types labeled for all A β variants. Diffuse plaques were strongly labeled by all A_β antibodies and were also highly abundant in the Pre-AD cases. Regarding the prominent labeling of isoAsp7-Aβ and A β (4-X) variants, isoAsp7-A β was found to be more abundant in plaques of compact and of cored morphology than $A\beta(4-X)$ (Fig. 6a). For the staining of cerebral blood vessels, the grading differentiated between no vascular amyloid deposits in any case of a clinical group (-), in 1 to 40% of the cases (+ -), in 41–80% of the cases (+) and in 81-100% of the cases (++). The pSer8-A β and pGlu11-A β variants were highly abundant in blood vessels in all clinical conditions including Pre-AD and control cases (Fig. 6a). In comparison, the A β (4-X) variant was less abundant in the vasculature of the VAD and Pre-AD group, whereas pGlu3-Aß and isoAsp7-Aß were present in all clinical conditions in a smaller subset of patients.

We next asked whether the A β variants present in amyloid plaques display different aggregation kinetics than their counterparts not detected in plaques. ThT assays were employed to monitor the time-dependent fibril formation of all A β (x-40) variants. Intriguingly, the pSer26-A β (1–40) and isoAsp27-A β (1–40) variants which had not been localized in plaques only showed low ThT fluorescence. On the contrary, $3NTyr10-A\beta(1-40)$ which had not been detected in amyloid plaques, did show ThT fluorescence and, thus, robust time-dependent fibril formation in the aggregation assay. The other A β variants detected in amyloid plaques also showed ThT fluorescence with instant fibril formation for pGlu3-A β (1–40), isoAsp7-A β (1–40) and pGlu11-A β (1–40), and lag phases ranging from 1.32 h for pSer8-A β (1–40) to 3.21 h for 3NTyr10-A β (1–40) (Fig. 6b). The A β (4-X) variant is an exception from this pattern, since it was highly abundant in amyloid plaques, but the A β (4–40) variant did not form fibrils under the conditions of the ThT assay. This might indicate that the A β (4–42) variant dominates in plaques.

To reveal specific characteristics of the fibrils derived from distinct A β variants, TEM was performed (Fig. 6c). In general, the TEM images confirmed and substantiated the data of the fibrillation curves obtained in the ThT assay, although a few A β (4–40) fibrils were detected (Fig. 6c). Remarkably, pGlu3-A β (1–40) and isoAsp7-A β (1–40) fibrils appeared to be shorter and thicker than fibrils of the other A β variants (Fig. 6c).

Biochemical analyses

We used immunoassay analyses to quantify those $A\beta$ variants that were shown above to be present in amyloid plaques. The $A\beta$ variants $3NTyr10-A\beta$, $pSer26-A\beta$ and $isoAsp27-A\beta$ were not detected in amyloid plaque-like formations and, Fig. 4 a Subgroup analyses of the abundance of $A\beta$ variants by APOE genotype. Note the high plaque load for all AB variants in APOE 3/4 carriers and the particularly high abundance of isoAsp7-Aβ compared to the other $A\beta$ PTMs in this subgroup. $\dot{\mathbf{b}}$ Pearson correlation analyses between the immunohistochemically quantified plaque load for individual AB variants and the neuropathological Braak Tau stages (column 1) and Thal amyloid phases (column 2). In addition, correlations between the respective abundance of $A\beta$ variants and the MMSE scores (column 3) are presented. Pearson correlation was considered as moderate with correlation coefficients of $0.40 \le r < 0.70$ (highlighted in light grey) and as strong with $0.70 \le r < 0.90$ (highlighted in grey). Note the moderate and strong correlations of isoAsp7-A β with Braak Tau stages, Thal amyloid phases and MMSE scores. The individual cases relate to the clinical condition as follows: black-Co; grey—Pre-AD; red—AD; blue—DLB; green—VAD







b Correlation coefficient r



Fig. 5 Heat maps for the plaque load of $A\beta$ variants and the correlation coefficient r of the Pearson correlation analyses. **a** Immunohistochemically quantified plaque load of $A\beta$ variants is depicted for the different conditions. **b** For all correlations between the plaque load of

A β variants and Braak and Thal staging and MMSE score, the correlation coefficient r as a measure for the quality of the correlation is presented. Note that the typical correlation pattern is not observed for pSer8-A β

thus, immunoassay analyses of these A β variants are shown in Suppl. Figure 5.

A β was separated based on solubility in TBS, GdmCl and FA fractions. For each A β variant, less than 1% of the sum of all fractions was detected in the TBS fraction, whereas GdmCl and FA fractions accounted for about 50% of each A β variant (Fig. 7). Due to potential dephosphorylation of pSer8-A β in FA [21], pSer8-A β was not measured in this fraction.

Consistent with immunohistochemical data on the quantification of the respective plaque load, immunoassay analyses revealed the highest A β concentrations in AD, followed by DLB, Pre-AD, VAD and controls. Specifically, in control cases, Braak stage I to III and Thal phase 0–1, a low amount of all A β variants was measured (Fig. 7). In the Pre-AD cases, Braak stage I to III and Thal phase 2–5, the median amount of A β was close to control subjects and varied between 341 pg/mg for A β and 9 pg/mg for pSer8-A β in GdmCl and between 161 pg/mg for A β and 17 pg/mg for A β (4-X) in FA.

In AD cases, the highest A β concentration was measured in the GdmCl and the FA fraction for A β with 2,331 pg/mg and 1,141 pg/mg, respectively. The lowest concentration was measured for pSer8-A β with 8 pg/mg in GdmCl. In GdmCl fractions, the amount of A β (4-X) (564 pg/mg) was highest among the A β variants, followed by isoAsp7-A β (386 pg/ mg) and pGlu3-A β (337 pg/mg). In FA, the same pattern as in GdmCl was observed with the highest concentration of A β (4-X) (737 pg/mg), followed by isoAsp7-A β (309 pg/mg) and pGlu3-A β (263 pg/mg).

In DLB cases, Braak stage I to IV and Thal phase 0–5, A β amounts in the GdmCl fraction ranged from 9 pg/mg for pSer8-A β to 951 pg/mg for A β . The variants pGlu3-A β , A β (4-X) and isoAsp7-A β showed similar concentrations with around 69 to 269 pg/mg in both GdmCl and FA. In the VAD cases, Braak stage I to IV and Thal phase 0–3, all A β variants accounted for low concentrations in GdmCl and FA.

In analogy to the correlations of immunohistochemically determined plaque load, we performed a subgroup analysis of the A β load by APOE genotype and correlated the abundance of the A β variants quantified by immunoassay with Braak and Thal staging and with MMSE score (Fig. 8). The highest quantities for all A β variants were detected in APOE 3/4 carriers, followed by APOE 3/3 carriers (Fig. 8a). In these APOE genotypes, A β PTMs displayed significantly lower concentrations than A β . In addition, the A β variants showed a higher correlation with Thal amyloid phases than with Braak Tau stages, with the strongest correlations for A β , A β (4-X) and isoAsp7-A β (r values between 0.70 and 0.90 indicate a strong correlation; highlighted in grey), and a moderate correlation for pGlu3-A β (r value 0.621; marked with a light grey) (Fig. 8b). Regarding the MMSE,

a Plaque morphology

Aβ Variant	Comp	act Pl	aque	C	oarse-G	iraine	d Pla	que		Core	d Pla	que			Diffus	se Pl	aque		Va	scular	· Dep	oosit	ion
	•						2				Y							and the second sec				0 µm	
	Co Pre-AD	AD	DLB VA	DC	D Pre-Al	D AD	DLB	VAD	Co	Pre-AD	AD	DLB	VAD	Co	Pre-AD	AD	DLB	VAD	Co	Pre-AD	AD	DLB	VAD
Αβ	+- ++	++	++ +	+	++	++	++	+		+-	+-	+-	+-	+	++	++	++	+	+-	+-	+	+-	+
pGlu3-Aβ	- ++	+	+ +	+	+	++	++	+-	-	+-	++	++	+-	+	++	++	++	+	+-	-	+-	-	+-
Αβ(4-Χ)	+- +	+	++ +	+	+	++	++	+	-	-	+-	+	-	+	++	++	++	+-	++	+	++	++	+
isoAsp7-Aβ	• ++	++	• •	+	+	++	++	+	•	+-	+	+	+-	+	++	++	++	+	+	+-	+	+	+
pSer8-Aβ	+- ++	+-	++ +	+	++	+	+	+	-	+	++	+	+-	+	++	++	++	+	++	++	++	++	++
pGlu11-Aβ	- +-	+-			++	++	++	+	•	+-	+	+	+-	+-	++	++	++	+	++	++	++	++	**

 Plaque morphology:
 - no staining
 +- some positive staining
 + positive staining
 ++ prominent positive staining with high plaque load

 Vascular deposition:
 - 0 %
 +- 1 - 40 %
 + 41 - 80 %
 ++ 81 - 100 % of cases with vascular deposition





	t _{lag} [h]	t _{1/2} [h]
 Aβ(1-40) 	1.86	7.36
 pGlu3-Aβ(3-40) 	-	1.33
 Aβ(4-40) 	n.d.	n.d.
 isoAsp7-Aβ(1-40) 	0.26	1.47
 pSer8-Aβ(1-40) 	1.32	2.77
 3NTyr10-Aβ(1-40) 	3.21	5.39
 pGlu11-Aβ(11-40) 	-	2.03
 pSer26-Aβ(1-40) 	n.d.	n.d.
 isoAsp27-Aβ(1-40) 	n.d.	n.d.

c TEM



Fig. 6 a Association of A β , pGlu3-A β , A β (4-X), isoAsp7-A β , pSer8-Aß and pGlu11-Aß variants with amyloid plaques of compact, coarse-grained, cored and diffuse morphologies as well as with cerebral vessels. The images on the top show labeling of A β . The quantifications below indicate a high abundance for most Aß variants in all plaque types, in particular in AD and DLB. In blood vessels, pSer8-Aß and pGlu11-Aß variants were particularly abundant in all clinical conditions, whereas pGlu3-Aβ and isoAsp7-Aβ were present in a smaller subset of patients in all clinical conditions. b Aggregation curves of time-dependent fibril formation of the Aß variants. Note the instant fibril formation of pGlu3-A β (1-40). As compared to unmodified A β (1–40), the variants pGlu3-A β (1–40), isoAsp7-Aβ(1-40), pSer8-Aβ(1-40) and 3NTyr10-Aβ(1-40) showed shorter lag phases, indicating more rapid formation of β-sheet containing aggregates. We did not observe significant fibril formation of A β (4–40), pSer26-A β (1–40) and isoAsp27-A β (1–40) variants. t_{lag}lag phase, $t_{1/2}$ —half maximum ThT fluorescence intensity time. c Electron microscopic images of fibrils derived from the respective Aß variants

we observed moderate negative correlations with *r* values between -0.415 and -0.510 for all A β variants except for pSer8-A β (Fig. 8b).

The correlation analyses of the biochemically determined concentrations of the $A\beta$ variants with histopathological and clinical data revealed results almost identical to the immunohistochemical data, indicating strong inter-assay reliability of our analytical methods.

To better visualize the abundance of the individual $A\beta$ variants in different tissue fractions and in distinct types of dementia and to indicate the strength of correlations with Braak and Thal staging and MMSE scores, heat maps were established (Fig. 9). These indicate the particular high abundance of pGlu3-A β , A β (4-X) and isoAsp7-A β in GdmCl and FA fractions in AD and to a lesser extent in DLB (Fig. 9a). For all A β variants except pSer8-A β , a moderate to strong correlation to the Thal amyloid phases was observed (Fig. 9b). Likewise, for all A β variants, except pSer8-A β , moderate negative correlations with MMSE were observed (Fig. 9b).

Discussion

In this study, we present an extensive and, to our knowledge, the most comprehensive analysis of eight distinct $A\beta$ PTMs in human brain tissue affected by different types of dementia. Our key discovery is that the isoAsp7 modification is a highly abundant $A\beta$ PTM in all clinical conditions including Pre-AD, AD, DLB and VAD.

Association of $A\beta$ variants with amyloid plaques in different types of dementia

The two A β variants with the highest abundance identified in this study in all clinical conditions were the isoAsp7-A β and A β (4-X) variants (see Suppl. Figure 6).

The isoAsp7-A β variant appeared to be very prominent among all clinical conditions observed when quantifying immunohistochemical plaque load and concentrations of Aß variants in GdmCl and FA fractions by immunoassay. We observed strong correlations of the abundance of this variant with Thal phase and moderate correlations with the decline in MMSE. In addition, we report robust and instant fibril formation without lag phase in aggregation assays. This $A\beta$ modification was identified over 30 years ago in the brain parenchyma [90]. The isoAsp7-A β variant was observed to induce cerebral amyloidosis when administered peripherally to transgenic mice with amyloid pathology [54] and to be more neurotoxic towards human neural stem cells than $A\beta(1-42)$ [69]. In addition, an $A\beta$ variant with a dual isoAsp modification at residues 1 and 7 was shown to be abundant in parenchymal and vascular A β deposits in AD [75], as well as in the Iowa variant of familial AD [115]. Another study showed the neuroprotective effect of protein-L-isoaspartyl methyltransferase, an ubiquitous enzyme that catalyses the conversion of isoAsp back to Asp, against Aß oligomers by increasing the size and reducing both hydrophobicity and toxicity of A β oligomers [23]. We observed an instant fibril formation for the isoAsp7-Aβ variant in the ThT assay. Intriguingly, targeting isoAsp7-Aβ with the monoclonal antibody K11 used in this study has already been demonstrated to reduce amyloid pathology and to ameliorate behavioural deficits in 5xFAD mice [38, 39]. This makes this A β variant particularly interesting as a therapeutic target for different types of dementia.

 $A\beta(4-X)$ was identified in AD brain and has been shown to account for more than 60% of the FA extractable A β peptides from amyloid plaque cores that were accessible to Edman protein sequencing [66, 68]. However, considering the abundance of pGlu3-AB and pGlu11-AB that escape Edman protein sequencing, its proportion in plaques should be lower but still significant. Subsequently, its presence was demonstrated by mass spectrometric analysis in the AD brain [83], in parenchymal and vascular deposits in the Iowa variant of familial AD [115], as well as in Danish amyloid (ADan) and A β co-deposits in Familial Danish Dementia [114]. Recently, $A\beta(4-X)$ was shown to be produced by ADAMTS4, a secreted metalloprotease that is exclusively expressed in oligodendrocytes [120]. Moreover, N-terminal A β truncation at position 4 was demonstrated to result in the formation of poorly soluble, aggregation-prone peptides with high amyloidogenic propensity and the potential of exacerbated fibrillary deposit formation [22]. A study using



D Springer

<Fig. 7 Quantification of A β variants by immunoassays. Note that the Y-axes differ for the individual A β variants and for TBS, GdmCl and FA fractions. Differences between clinical groups are statistically significant at *p < 0.05; **p < 0.01; ***p < 0.001. Medians are indicated by horizontal lines

a polyclonal A β (4-X) antibody confirmed the presence of $A\beta(4-X)$ in amyloid plaques in two transgenic AD mouse models but also showed that $A\beta(4-X)$ was present in human AD cases in blood vessels and in neuritic plaques but not in diffuse amyloid deposits [123]. These findings were further expanded by a study using another monoclonal antibody (clone 18H6) against A β (4-X). In this study, an association between A β (4-X) and cored plaques, CAA and also diffuse plaques was reported in human AD brain tissue, along with an enhanced brain retention of oligometric A β (4-X) [94]. In our study, the polyclonal A β (4-X) antibody detected the A β variant in all plaque types as well as in vascular amyloid deposits which is in line with the initial characterization of this antibody [10] and a recent study using the monoclonal antibody 18H6 against A β (4-X) [125]. In our ThT assay and transmission electron microscopy, only little fibril formation was detected for A β (4–40), suggesting that the majority of $A\beta(4-X)$ deposited in brain parenchyma and cerebral blood vessels is $A\beta(4-42)$.

There are a number of reports in the literature that point to the presence of individual or multiple $A\beta$ variants in brain tissue of one or more clinical conditions. For example, A β , the pGlu3-A β and pSer8-A β variants were compared side-by-side using immunohistochemical and Western blot analyses in control, Pre-AD and AD [89]. This allowed for a biochemical staging of $A\beta$ deposition, where both A β PTMs were absent in stage 1 as the earliest preclinical stage, only pGlu3-A β was present in stage 2 and both modified A^β variants were present in the last stage of A^β aggregation, stage 3 [89]. The higher abundance of pGlu3-A β compared to pSer8-Aß in Pre-AD (median plaque load of 1.5% for pGlu3-A β and 0.6% for pSer8-A β) and AD cases (median plaque load of 3.4% for pGlu3-A β and 1.0% for pSer8-A β) was also demonstrated in our study and was quantified in detail by automated immunohistochemical analysis and immunoassay methods. Another study analyzed the pGlu3-A\beta and pSer8-A\beta variants in cerebral amyloid angiopathy (CAA) and found a similar pattern of vascular amyloid stages with pGlu3-A preceding pSer8-A deposits in microvessels [36]. It should be noted that the appearance of A_β coincides with widespread Tau pathology at later stages of AD and that there is Tau co-pathology of DLB [70, 106]. Although Tau pathology was not investigated in the present study, we would like to point out that Tau is an important mediator required for pGlu3-Aβ toxicity [77]. Moreover, pGlu3-A load in AD predicted the hyperphosphorylated Tau load and was related to the severity of AD neuropathology and clinical dementia [65]. In addition, pGlu3-A β and isoAsp-A β displayed low immunoreactivity in non-demented controls and were significantly increased in AD [75].

Other studies focused on pSer8-A β and pGlu3-A β , individually, in different clinical conditions and related animal models. For example pSer8-A β was identified in AD brain and reported to form neurotoxic oligomers that act as nuclei for A β fibril formation [55, 56]. In addition, pSer8-A β was found to be present in Down syndrome and transgenic mouse models with A β pathology [60] and in brains of non-human primates and canines [58]. However, the pSer8-A β variant was demonstrated to be exclusively associated with a subset of A β plaques and vascular deposits in familial and sporadic AD and to be absent or only detectable at very small amounts in control, DLB and VAD brains [7]. This accumulation pattern of the phosphorylated A β variant pSer8-A β in Pre-AD, AD, DLB and VAD brain tissue was substantiated in our present study.

The presence of the pGlu3-Aß variant in AD amyloid plaques was initially reported by the Saido and Roher groups [63, 97]. It was subsequently reported to be present in the brains of humans, non-human primates, canines and transgenic mice with amyloid pathology [30, 44, 71, 72, 122]. Because of its instant aggregation [104], high neurotoxicity [2, 77], compromised degradation [95, 96] and seeding capacity [77, 101], pGlu3-Aβ emerged as a pharmacologic target for AD therapy. There are two strategies of interfering with pGlu3-A β , (i) prevention of its generation by inhibition of glutaminyl cyclase [46, 103] and (ii) specific targeting of existing pGlu3-A β by immunotherapy [24, 26, 27, 29, 31]. Recently, intravenous infusions with Donanemab (Kisunla) targeting pGlu3-A β have been approved by the U.S. Food and Drug Administration for early symptomatic AD treatment [51].

As discussed above, in the course of AD, pGlu3-A β is present earlier than the pSer8-A β variant and its abundance is higher. In addition, compared to the other A β variants, pGlu3-A β displayed lower concentrations than isoAsp7-A β and A β (4-X) in brains from Pre-AD, AD, DLB and VAD cases. The related pGlu11-A β variant, on the other hand, has been studied less intensively, but was shown to be co-localized with pGlu3-A β in amyloid plaques in AD brain tissue and to form the central plaque core [79, 111]. In addition, structural characteristics and neurotoxicity of pGlu11-A β were recently reported [98, 99]. We here observed that the pGlu11-A β variant only accounts for a small proportion of the A β plaque load in all clinical conditions investigated.

Another important aspect is that longer A β 42 and A β 43 peptides are more directly linked to AD based on early pathology, familial AD mutations in presenilins, Down syndrome amyloid pathology and A β biomarkers than the shorter A β 40 variants [16, 33, 73, 85]. This also holds true



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∢Fig.8 a Subgroup analyses of the abundance of Aβ variants by APOE genotype in GdmCl fractions. Note the high quantity of Aβ in APOE 3/3 and APOE 3/4 carriers. **b** Pearson correlation analyses between the load for individual Aβ variants quantified by immunoassays in the GdmCl fractions and the neuropathological Braak Tau stages (column 1) and Thal amyloid phases (column 2). In addition, correlations between the abundance of Aβ variants and the MMSE scores (column 3) are presented. Pearson correlation was considered as moderate with correlation coefficients of $0.40 \le r < 0.70$ (highlighted in light grey) and as strong with $0.70 \le r < 0.90$ (highlighted in grey). The individual cases relate to the clinical condition as follows: black—Co; grey—Pre-AD; red—AD; blue—DLB; green – VAD

for higher neurotoxicity and faster aggregation characteristics of A β 42 versus A β 40 [48, 49, 81]. In the present study, pGlu3-A β (3–40), isoAsp7-A β (1–40), pGlu11-A β (11–40) and pSer8-A β (1–40) showed faster aggregation compared to unmodified A β (1–40). Thus, these PTMs could increase the pathogenicity of A β 40 variants towards that of A β 42.

Aβ variants not associated with amyloid plaques

In the present study, the A β variants 3NTyr10-A β , pSer26-A β and isoAsp27-A β were not detected in amyloid plaques and only the 3NTyr10-A β (1–40) variant was shown to form fibrils in aggregation experiments of A β (1–40) variants in vitro. Importantly, in the case of immunohistochemical labeling and immunoassay analyses, pre-analytical treatments of brain tissue and samples may influence the final results and should be considered when comparing different results. However, it seemed striking that the plaque-associated A β variants investigated were post-translationally modified at the N-terminus of the A β peptide. Due to the high β -sheet content of aggregated A β in plaques [3], it might be difficult for monoclonal antibodies against pSer26-A β and isoAsp27-A β to bind to their epitopes, which might be sterically inaccessible in the plaque-like formation.

For 3NTyr10-A β , the NOS2-catalyzed nitration was shown by using enzyme inhibitors and knock-out mice [61]. This A β variant was prone to fibril formation and was detected in the core of amyloid plaques in brains from AD patients and transgenic mice with amyloid pathology [61]. Since these initial data were obtained with a polyclonal antiserum, we developed a specific monoclonal antibody targeting this particular $A\beta$ variant for the present study. While we could reproduce the biochemical characteristics of 3NTyr10-A β in the ThT and TEM assays (see Fig. 6b, c), we did not detect an association of $3NTyr10-A\beta$ with amyloid plaques in any of the clinical conditions (see Suppl. Figure 4). Instead, the monoclonal antibody showed an intraneuronal localization of 3NTyr10-A^β. Discrepancies in the staining pattern of the polyclonal antiserum and the monoclonal antibody might arise from impacts of AB conformation and secondary structure or different detection sensitivities of the antibodies as well as brain tissue storage conditions and pre-analytical sample preparations.

The pSer26-A β has recently been demonstrated to give rise to stable oligomeric assemblies with high neurotoxicity. Furthermore, the pSer26 modification has been shown to impair the fibrillation of the A β peptide [57, 88]. In a transgenic AD mouse model and in human AD brain, pSer26-A β was found to be localized particularly in intraneuronal deposits [57]. The lack of fibril formation of pSer26-A β (see Fig. 6) and its intraneuronal labeling was confirmed here in all types of dementia under investigation, especially in AD and DLB (see Suppl. Figure 4).

In addition to the previously described isoAsp7-Aβ variant (see above), we here also aimed at an analysis on the potential formation of isoAsp27-A β , for which we generated and characterized a monoclonal antibody (see Suppl. Figure 2). There are several reports in the literature showing that Asp isomerization may be a phenomenon associated with aging and neurodegeneration in general and in AD in particular [47, 74, 78, 93]. For this reason, antibodies raised against isoAsp-modified Aß have been proposed as indicators of the plaque age [28]. In general, isoAsp is formed at sites with Asn or Asp amino acid precursors, whereas the rate of isoAsp formation from Asn precursors is typically one-to-two orders of magnitude faster than from Asp [35, 41]. Hence, it is tempting to speculate that $isoAsp27-A\beta$ might be formed in addition to isoAsp7-Aβ. However, the formation and deposition of isoAsp27-Aβ appears negligible and to be restricted to intracellular compartments. A potential rationale for this observation might be the placement of Asn27 at the contact area of A β protofilaments to form fibrils [124]. A modification of this residue to isoAsp might prevent the formation of fibrils. Likewise, isoAsp27-A β (1–40) did not show fibril formation in our analysis.

Taken together, although not abundantly present in different types of dementia, the 3NTyr10-A β , pSer26-A β and isoAsp27-A β variants were detected in the soma of pyramidal neurons. Hence, a potential influence on neuronal and synaptic dysfunction cannot be excluded, as previously suggested for pSer26-A β [57].

It is quite obvious that substantial modifications to the majority of A β peptides are restricted to the N-terminal residues 1–8, as also suggested recently [76]. Cryo-EM and NMR data collectively suggest that these residues are not incorporated into fibrils and are therefore structurally flexible and accessible to enzyme catalysis or spontaneous isoAsp formation.

Association of $A\beta$ variants with cerebral blood vessels

The pGlu3-A β and isoAsp7-A β variants were observed to be less frequently associated with cerebral microvessels than



a Amount of Aβ variants

b Correlation coefficient r 1.0 Aβ (GdmCl) 0.5 pGlu3-Aβ (GdmCl) Aβ variant Aβ(4-X) (GdmCl) 0 isoAsp7-Aβ (GdmCl) -0.5 pSer8-Aβ (GdmCl) -1.0 MMSE Braak Thal r Staging

Fig.9 Heat maps for the amount of $A\beta$ variants quantified in the TBS, GdmCl and FA fractions and the correlation coefficient r of the Pearson correlation analyses. **a** $A\beta$ variants quantified by immunoassays are depicted for the different clinical conditions. **b** For all

the other A β variants investigated in this study. However, both A β variants, as well as the pGlu11-A β , are not completely absent from blood vessels, a finding also reported recently for AD [109]. Vascular A β deposits are believed to contribute to blood–brain-barrier disruption and to account for amyloid-related imaging abnormalities (ARIAs), edema and hemorrhages, during the anti-A β immunotherapies [11, 43, 50]. Thus, targeting A β variants less frequently present in blood vessels should be expected to reduce these adverse events. However, in the TRAILBLAZER trials using the antibody Donanemab to target the pGlu3-A β variant in patients, such events were not less frequent than in other clinical trials [43, 108].

Conclusions

Post-translationally modified A β peptides play an important role in the initiation of aggregation processes that result in the formation of oligomeric/ fibrillary assemblies with high neurotoxicity. We hypothesized that different types of dementia are characterized by specific patterns of modified A β variants. Surprisingly, we did not identify such a disease-specific signature of post-translationally modified A β peptides but a rather consistent pattern of A β variants across different clinical entities. We discovered isoAsp7-A β as the most abundant A β variant, followed by A β (4-X) and pGlu3-A β , in all types of dementia assessed here. It remains

correlations between the A β concentrations and Braak and Thal staging and MMSE score, the correlation coefficient r as a measure for the quality of the correlation is presented. Note that the typical correlation pattern is not observed for pSer8-A β

an open question whether the most abundant A β variant is the best pharmacologic target. However, in preclinical studies, the efficacy of antibodies against isoAsp7-A β [38] and pGlu3-A β [27, 31] has already been demonstrated and the pGlu3-A β antibody Donanemab was recently approved for AD therapy. We believe that it is worth considering Donanemab treatment in other types of dementia and that isoAsp7-A β and A β (4-X) are additional targets for immunotherapy in different clinical conditions with an amyloid component.

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Author contributions Sa.S. designed the study, performed the histological and biochemical tissue processing, immunohistochemical analyses of human brain tissue, imaging via transmission electron microscopy, statistical analyses as well as data visualization of immunohistochemical and biochemical data and was a major contributor in manuscript writing, A.K.K. designed the study, isolated antibodies, validated antibody specificity, performed biochemical tissue processing, biochemical quantifications in human brain tissue, Aß aggregation and imaging via transmission electron microscopy and contributed to manuscript writing. A.P. and K.G. generated antibodies and provided scientific advice. L.B. performed biochemical quantifications in human brain tissue. M.H.R. provided scientific advice and contributed to manuscript writing. M.M. performed imaging via transmission electron microscopy and provided scientific advice. M.S. and M.K. generated antibodies. G.E.S. and T.G.B. helped select human subject cases appropriate for the project, performed clinico-pathological classification of human subjects and contributed to manuscript writing and editing. J.G. and A.R. provided experimental tools, gave scientific advice and reviewed the manuscript. M.H. and J.W. provided experimental tools, gave scientific advice on interpretation of data and contributed to manuscript writing. O.W. provided experimental tools, supervised biochemical analyses, gave scientific advice on interpretation of data and contributed to manuscript writing. St.S. initiated, designed and supervised the study and contributed to manuscript writing. S.R. initiated, designed and supervised the study and was a major contributor in manuscript writing. All authors reviewed the manuscript.

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Declarations

Conflict of interest The authors declare no competing interests.

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