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## Luminex-based quantification of Alzheimer's Disease neuropathologic change in formalin-fixed post-mortem human brain tissue

C. Dirk Keene<sup>a,\*</sup>, Angela M. Wilson<sup>a,\*</sup>, Mitchell D. Kilgore<sup>a</sup>, Lauren T. Bruner<sup>a</sup>, Nadia O. Postupna<sup>a</sup>, and Martin Darvas<sup>a</sup>

<sup>a</sup>Department of Pathology, University of Washington, Seattle WA 98104

### Abstract

The vast majority of archived research and clinical pathological specimens are stored in the form of formalin fixed, paraffin-embedded (FFPE) tissues, but, unlike fresh frozen tissue samples, highly quantitative measures in FFPE tissues are limited to immunohistochemical and immunofluorescence thresholding image analysis studies, cell counting, and ordinal ranking systems. This poses a significant obstacle for clinical investigations that aim to correlate diagnostic markers of neurodegenerative diseases like Alzheimer's disease (AD) with parameters like age, gender, drug exposures, genotype, disease stage, co-morbidities or environmental factors. To overcome this limitation, we have developed Luminex-based techniques and protocols for the quantification of amyloid  $\beta$  and hyperphosphorylated Tau in FFPE brain sections. We validated the Luminex assay in FFPE sections from prefrontal cortex, hippocampus and neostriatum from 30 cases that underwent prior neuropathological diagnostic assessment of AD following the current NIA-AA recommendations for AD: 10 cases diagnosed as not or low, 10 cases as intermediate, and 10 cases as high AD neuropathologic change. Consistent with the neuropathologic assessment, Luminex assay detected high amounts of amyloid beta in the frontal cortex and striatum, and high amounts of hyperphosphorylated Tau in the frontal cortex and hippocampus, of cases with high AD neuropathologic change. This assay can be expanded to detect diverse antigenic targets of interest, as we show here with IBA1 and GFAP. This novel approach supports multiplexed highly quantitative, molecularly specific neuropathology measures to further explore mechanisms of neurodegeneration in AD.

### Keywords

Luminex; Amyloid  $\beta$ 42; phospho-Tau; Alzheimer's disease; cortex; neostriatum; hippocampus; formalin-fixed; paraffin-embedded

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Correspondence: Martin Darvas (206-897-5581), [mdarvas@uw.edu](mailto:mdarvas@uw.edu), Department of Pathology, Box 359645, University of Washington, Seattle 98104.

\*These authors contributed equally.

### DISCLOSURE/CONFLICT OF INTEREST

The authors declare no conflict of interest.

The pathognomonic lesions of Alzheimer's disease (AD) are neuritic plaques and neurofibrillary tangles (NFT) [1]. Amyloid  $\beta$  peptide 42 (A $\beta$ 42) is a key component of neuritic plaques, and antibodies against A $\beta$ 42 are used in immunohistochemistry (IHC) procedures required for the neuropathologic diagnosis of AD. NFTs are primarily composed of hyperphosphorylated Tau (phospho-Tau), and antibodies against phospho-Tau (pTau) epitopes are used in routine IHC procedures for neuropathologic assessment of AD. In fact, the current National Institute on Aging – Alzheimer's Association (NIA-AA) consensus criteria for the neuropathologic diagnosis of AD are based on histologic characterization of the anatomic distribution of amyloid plaques and NFTs [1, 2]. These criteria are based on previously established criteria for A $\beta$  plaque [3] and NFT distribution [4, 5], and cortical density of neuritic plaques [6]. These diagnostic criteria are not quantitative measures but rather ordered rankings, which severely limits their utility in research studies that aim to correlate pathological features of AD with parameters like age, gender, drug exposures, genotype, co-morbid processes, or environmental factors.

Because the vast majority of archived pathological tissues in research tissue banks and in clinical archives are maintained in the form of formalin fixed paraffin-embedded (FFPE) tissue, approaches to quantify AD neuropathologic change in brain tissue are mostly restricted to immunohistochemistry (IHC) or histochemical methods. These approaches are inherently limited by issues relating to a variability of image thresholds, obtaining uniform optics across tissue, quenching endogenous signals, and field selection. Although unbiased-stereology based protocols allow counting of antibody-stained cells and other discrete structures [7], they do not usually provide a measure for quantifying antibody-binding events and are labor intensive.

Recently, automated image-processing protocols and tissue microarray techniques, similar to those employed in cancer studies, have been applied to neurodegenerative diseases to overcome some of these limitations [8, 9]. For example, using computer-assisted image analyses, Boyle et al have been able to associate IHC measures of amyloid and NFT pathology with ante-mortem cognitive parameters [10]. However, sensitivity to detect low abundance targets and optical correction factors including, but not limited to, background signal subtraction, can reduce sensitivity. A different approach to generate quantitative measures of AD neuropathologic change is a combination of histologic evaluation and ELISA performed on a standard glass slide ("Histelide"), which has been used on over 300 brain autopsies so far [11–14]. The Histelide method is molecularly specific, but is limited by high cost and inability to effectively multiplex the approach. Here, we report a fundamental transformation of quantitative FFPE-tissue based techniques, by adding a Luminex-based approach to the analysis of FFPE tissue in neurodegenerative diseases that is cost-effective, sensitive, and can be multiplexed to limit tissue usage, while allowing cytoarchitectural correlation in immunohistochemically stained adjacent sections. Luminex technology combines magnetic-fluorescent bead-based immunoassays with advanced fluidics, optics and signal processing to permit quantification of multiple antigens within a single sample [15]. This technology has been widely used in frozen-tissue extracts, as well as in serum and cerebrospinal fluid samples from patients with a wide range of diseases and disorders [16], including AD [17–20].

Our laboratory has developed FFPE-compatible protocols that allow extraction of A $\beta$ 42, pTau, glial fibrillary acidic protein (GFAP) and ionized calcium-binding adapter molecule 1 (IBA1). In this manuscript we demonstrate the use of Luminex technology to quantify expression of these proteins in human FFPE sections from three brain regions included in the National Institute on Aging and the Alzheimer's Association (NIA-AA) recommendations for neuropathologic assessment of AD [2]. Using this method, we verified increased amounts of A $\beta$ 42 and pTau in brains with higher levels of AD neuropathologic change. The highly quantitative and multiplexed approach allows comparisons between levels of pathologic peptides and other markers, in the same tissue section, thus allowing for more precise assessment of correlations with risk factors and other parameters of interest. Using Luminex to quantify A $\beta$ 42 and pTau in FFPE tissue offers several advantages that complement existing IHC-based analyses of neurodegenerative change in AD, and provides new opportunities for molecular analyses of AD and other neurodegenerative diseases. Because of its inherent quantitative nature, high degree of sensitivity and specificity, and the ability to simultaneously analyze multiple (up to 48 if duplicate runs are used on a 96-well plate) samples on one Luminex plate, this assay is well suited for high-throughput analysis of A $\beta$ 42 and pTau levels in FFPE autopsy brain tissue.

## MATERIALS AND METHODS

### Human Subjects

This study was approved by the University of Washington (UW) Institutional Review Board. All cases were from the UW Alzheimer's Disease Research Center (ADRC) Neuropathology and Targeted Molecular Testing (NPTMT) Core. All study subjects were participants in the Adult Changes in Thought study. Diagnostic evaluation of each case was performed by a board-certified neuropathologist in the NPTMT Core at the time of autopsy in accordance with the NIA-AA guidelines [1], which require assignment of an "ABC" score designating each subject as having "not", "low", "intermediate", or "high" AD neuropathologic change [2]. In order to test a range of AD pathologic change, we selected 9–10 cases from not/low, intermediate, and high categories that were matched by age, sex, post-mortem interval, and year of death (see Table 1).

### Preparation of antibody conjugated Luminex™ beads

Coupling Luminex MagPlex-COOH beads (Bio-Rad; Hercules, CA) to monoclonal antibodies was based on the procedures outlined in the X-MAP Cookbook [15]. All primary antibodies that contained amine containing additives or preservatives were cleaned using Micro Bio-Spin 6 Tris chromatography columns (Bio-Rad; Hercules, CA) according to the manufacturer's instructions. Table 2 lists the antibodies that were conjugated to MagPlex beads to detect A $\beta$ 42, pTau, GFAP and IBA1.

Conjugation of antibodies to Luminex™ MagPlex-COOH beads was performed using the Bio-Rad bead making kit (Bio-Rad; Hercules, CA), and conjugated beads were quantified using a TC20 cell counter (Bio-Rad; Hercules, CA). For confirmation of antibody coupling to beads, serial dilutions of phycoerythrin (PE) labeled anti-species IgG detection antibodies (Jackson ImmunoResearch; West Grove, PA) were used to detect primary bead-coupled

antibodies from mouse or rabbit hosts (Table 2). The beads were then read using the BioPlex® 200 Luminex instrument (Bio-Rad; Hercules, CA).

### Preparation of Biotinylated antibodies

The Luminex assay described in this manuscript uses a sandwich-assay format that requires two antibodies for each antigen. MagPlex bead-conjugated antibodies described in the previous section and in Table 2 served as antigen-capture antibodies and biotinylated antibodies (see Table 3 for list of detection antibodies) as detection antibodies. GFAP and IBA1 antibodies were obtained as biotin conjugates from their respective manufacturers (both Abcam; Cambridge, MA). Antibodies for detection of A $\beta$ 42 (Biolegend; San Diego, CA) and Tau (Thermo Fisher Scientific; Waltham, MA) were biotinylated in our laboratory using a One-step Antibody Biotinylation kit (Miltenyi; Auburn, CA).

### Generation of recombinant A $\beta$ 42 peptide

Coding sequences for green-fluorescent protein in frame with 2A peptide from porcine teschovirus-1 polyprotein and with a fusion construct encoding the first 243 amino acids of BRI followed by 42 amino acids encoding A $\beta$ 42 were cloned into an expression plasmid under the control of a human polypeptide elongation-factor 1 $\alpha$  promoter and containing a woodchuck hepatitis-virus post-transcriptional regulatory element, and a human growth hormone polyadenylation signal. A similar strategy has been successfully used for expression of secreted A $\beta$  peptides in cell lines and rats [21, 22]. We then transfected HEK cells with the A $\beta$ 42 expression plasmid using polyethylenimine (PEI, Polysciences; Warrington, PA) transfection. Cells were incubated with the DNA/PEI mix for 12 hours, after which we removed the medium containing the DNA/PEI mix and replaced it with growth medium (DMEM, 10% fetal bovine serum, penicillin/streptomycin - all from Life Technologies; Carlsbad, CA). Seventy-two hours after the transfection we collected the cell supernatant, removed cell contaminants by centrifugation (1,200 rpm), sterile-filtered the supernatant, added sodium azide (0.09%) for preservation, and stored aliquots at  $-20^{\circ}\text{C}$ .

### Protein extraction of formalin-fixed paraffin-embedded brain tissue

Autopsy brain tissue was immersion fixated in 10% neutral buffered formalin for 2–3 weeks. Brain regions containing Brodmann area 9 of middle frontal gyrus (MFG), hippocampus at the level of the corpus geniculatum laterale, and neostriatum at the level of the anterior commissure, were then sampled, embedded in paraffin, microtome sliced in 15- $\mu\text{m}$  sections, and placed on charged microscope slides [13]. Three adjacent 15- $\mu\text{m}$  sections per case were deparaffinized by immersion for 3 minutes in each of the following solutions: 3 $\times$  in 100% Xylene, 100% isopropanol, 96% isopropanol, 70% isopropanol, and 50% isopropanol. Sections were then washed in PBS containing 0.05% Tween 2 $\times$  for 4 minutes [14]. For antigen retrieval, sections were boiled for 20 minutes in 0.1 M citrate buffer at a pH of 6.1. Slides were then immersed in PBS containing 0.05% Tween. Tissue from three sections was then scraped off the slides with a razor blade and placed into a 2-ml tube.

*For extraction of A $\beta$ 42 and pTau*, 150  $\mu\text{l}$  of chilled RIPA buffer (1% NP-40, 0.5% sodium deoxycholate, 150 mM sodium chloride, 50 mM Tris hydrochloride, 0.5 mM magnesium sulfate – all from Sigma-Aldrich; St. Louis, MO) with Complete Mini protease inhibitor

(Sigma-Aldrich; St. Louis, MO) was immediately added to the tube. Samples were then sonicated on ice (10 pulses, 3×), and centrifuged for 45 minutes at 21,000 g and 4°C. Supernatants containing RIPA-soluble proteins were pipetted off into new 2 ml tubes. The remaining pellet was washed with an additional 50 µl RIPA buffer with Complete Mini, and centrifuged a second time. The RIPA-buffer containing supernatant was then combined with the first RIPA-buffer containing supernatant. The combined RIPA-buffer supernatant contains soluble proteins and pTau, but not Aβ42. For Aβ42 extraction, 150 µl of chilled 5M guanidine-hydrochloride (Gu-HCl) buffer containing Complete Mini was then added to the remaining pellet in tube, vortexed, then sonicated on ice for 10 pulses. Centrifugation at 13,000 g for 30 minutes at 4°C produced a Gu-HCl soluble supernatant containing the Aβ42 fraction and other RIPA insoluble proteins. All extracts were stored at –80 °C.

*IBA1 and GFAP* were extracted from deparaffinized sections using an FFPE extraction kit (Qiagen; Germantown, MD) by following the 2D-PAGE/MS protocol of the kit instructions. The resulting protein extract was dissolved in 100 µl RIPA with Complete Mini. All extracts were stored at –80 °C.

*Total protein content* was determined in all samples using a BCA kit (Pierce; Rockford, IL) together with colorimetric detection (absorbance at 562 nm) in a plate reader.

### **Luminex assay using customized antibodies and MagPlex beads**

Protein samples and standards were always diluted in PBS containing 1% bovine serum albumin (BSA, Sigma-Aldrich; St. Louis, MO) and conjugated MagPlex beads were always diluted in PBS with 1% BSA and Complete Mini. Prior to adding conjugated MagPlex beads to samples, beads were thoroughly resuspended by vortexing and sonication for 20 s each. Diluted samples and standards were then loaded onto a black flat-bottom 96-well plate (Greiner; Monroe, NC), mixed with resuspended MagPlex beads at a concentration of 2,500 beads per well, and incubated overnight. This and all following incubation steps were performed shaking in the dark and at room temperature. On the next day, beads were washed twice and then incubated for 3–4 hours with 100 ng of biotinylated antibody (in PBS with 1% BSA) per well. After that, beads were washed twice and then incubated for 1 hour with 1 µg of PE per well (Life Technologies; Carlsbad, CA). After that beads were washed twice, resuspended in PBS, incubated for 5 minutes, and then read using the Bio-Plex® 200 Luminex instrument.

*For analysis of Aβ42*, we first used different amounts of Gu-HCl soluble protein extracts to determine the optimal amount of sample (based on total protein content) per well, and then used that amount for the analysis of all experimental samples.

*For analysis of pTau*, we first used different amounts of RIPA-soluble protein extracts to determine the optimal amount of sample (based on total protein content) per well, and then used that amount for the analysis of all experimental samples.

*For analysis of IBA1 and GFAP*, we first used different amounts of RIPA-soluble protein extracts (Qiagen-kit) to determine the optimal amount of sample (based on total protein content) per well, and then used that amount for the analysis of all samples.

### Luminex assay using the ALZ kit

Recombinant A $\beta$ 42 was also measured using the INNO-BIA AlzBio3 kit (Fujirebio through INNOGENETICS; Gent, Belgium) according to the manufacturer's instructions.

### Commercial standards used for GFAP and IBA1 assays

For GFAP and IBA1 Luminex assays we identified suitable standards that we used for validation purposes and which will allow quantification of GFAP and IBA1 in future experiments. For GFAP we used porcine full-length protein purified from spinal cord (Millipore; Burlington, MA) and for IBA1 we used full-length recombinant human IBA1 (Abcam; Cambridge, MA). We prepared the following concentrations for GFAP: 320000 pg/ml, 160000 pg/ml, 80000 pg/ml, 40000 pg/ml, 20000 pg/ml, 10000 pg/ml, 5000 pg/ml, 2500 pg/ml, 1250 pg/ml and 625 pg/ml. For IBA1 we prepared the following concentrations: 40000 pg/ml, 20000 pg/ml, 10000 pg/ml, 5000 pg/ml, 2500 pg/ml, 1250 pg/ml, 625 pg/ml and 312.5 pg/ml.

### Measuring grey-matter area using immunohistochemistry and image analysis

Because most of the proteins measured in this study are known to be differentially distributed between grey and white matter, we needed to control for variability in the grey/white matter composition between samples. To accomplish that, 5  $\mu$ m sections immediately anterior to sections utilized for protein extraction were treated with Epitope Retrieval Solution 2 (Leica Biosystems, Buffalo Grove, IL) for 20 min, immunostained for synaptophysin with rabbit monoclonal MRQ-40 antibody (1:100, Cell Marque, Rocklin, CA), and counterstained with hematoxylin. The sections were then scanned at 40 $\times$  magnification with an Aperio AT2 Digital Slide Scanner (Leica Biosystems, Buffalo Grove, IL) and subjected to histomorphologic analysis using Aperio GENIE Histology Pattern Recognition software (Leica Biosystems, Buffalo Grove, IL) to differentiate between synaptophysin immunoreactive regions and calculate the grey matter area of each section. We used this procedure to control for potential differences of grey matter area in this analysis cohort.

### Statistical Analysis

All experimental and basic patient data are shown as mean  $\pm$  standard error of the mean (SEM). Data were analyzed using one- or two-way analysis of variance (ANOVA), followed by Tukey's post-test, or a two-tailed t-test. Correlations were evaluated by calculating Pearson's correlation coefficient, and linearity was assessed by linear regression. Alpha was always set at 0.05.

## RESULTS

### Generation of recombinant standards for quantification of A $\beta$ 42

To assess whether we could generate recombinant A $\beta$ 42 standards for the quantification of A $\beta$ 42, we transfected HEK cells with pGFP-BRI-A $\beta$ 42 and used the commercial Innogenetics Luminex kit to detect and quantify A $\beta$ 42 in the supernatant of transfected cells (72-hours post transfection). We confirmed the presence of A $\beta$ 42 in the supernatant of



transfected cells and measured an A $\beta$ 42 concentration of 934 pg/ml in the supernatant (Figure 1a). Next, we tested serial dilutions of recombinant A $\beta$ 42 and A $\beta$ 42 standards from the Innogenetics kit (each at 29 pg/ml, 58 pg/ml, 113 pg/ml, 233 pg/ml, 467 pg/ml and 934 pg/ml) using the Innogenetics kit for detection and quantification (Figure 1b). Analysis of fluorescent intensities of diluted recombinant A $\beta$ 42 and A $\beta$ 42 standards from the Innogenetics kit by 2-way ANOVA revealed a significant effect of A $\beta$ 42 concentration ( $F_{5,36} = 321.7$ ,  $p < 0.01$ ), with significant differences between different A $\beta$ 42 dilutions for both recombinant A $\beta$ 42 and A $\beta$ 42 standards ( $p < 0.05$ ), but with no significant differences between recombinant and commercial A $\beta$ 42 standards at equal dilutions ( $p > 0.05$ ). We then tested our recombinant A $\beta$ 42 standards with the newly developed Luminex A $\beta$ 42 assay (capture with H31L21 clone, detection with 6E10 clone). Using the same concentrations of recombinant A $\beta$ 42 that we analyzed with the Innogenetics Luminex kit, we were able to generate a standard curve (Figure 1c) similar to the one generated by commercial A $\beta$ 42 standards. We designated our recombinant standards as S1 (934 pg/ml), S2 (467 pg/ml), S3 (233 pg/ml), S4 (113 pg/ml), S5 (58 pg/ml) and S6 (29 pg/ml). We further determined whether our recombinant A $\beta$ 42 standards could be stored and used over an extended period of time (Figure 1d). 2-way ANOVA of fluorescent intensities of S1–S6 standards measured immediately after, 2 months and 6 months after preparation of recombinant A $\beta$ 42 revealed significant main effects of standard concentration ( $F_{5,30} = 197$ ,  $p < 0.01$ ), but no significant effects of time passed between preparation of standards and measurement of fluorescent intensities ( $p > 0.05$ ).

### Detection of A $\beta$ 42 in human FFPE tissue

To determine whether we could use the same antibodies that are routinely used in neuropathology immunohistochemistry assessments for Luminex-based detection of A $\beta$ 42, we extracted Gu-HCl soluble fractions from the MFG from 3–5 additional AD cases that were not part of the experimental cohort, and tested increasing amounts of each fraction for the presence of A $\beta$ 42. Increasing amounts of Gu-HCl soluble protein extracts resulted in increasing fluorescent signal intensities in the A $\beta$ 42 assay (capture with H31L21 clone, detection with 6E10 clone), and these signal intensities were approximately linear for protein amounts between 15.625 ng and 250 ng, approximately spanning a 35-fold dynamic range (Figure 2a). Based on the observed input curve, we determined that ~150 ng of Gu-HCl soluble proteins allows for a strong A $\beta$ 42 signal that is well within the linear range of the assay.

### Quantification of A $\beta$ 42 in FFPE tissue from AD patients and controls

Next we measured A $\beta$ 42 levels in Gu-HCl extracts from FFPE MFG, hippocampal and neostriatal tissue from patients with not/low, intermediate, and high levels of AD pathology (assessed per NIA-AA criteria), and then compared A $\beta$ 42 levels between samples for each brain region. Comparisons of A $\beta$ 42 levels in the MFG (Figure 2b) by ANOVA revealed significant effects of NIA-AA AD severity ( $F_{2,26} = 5.67$ ,  $p < 0.01$ ). *Post-hoc* comparison with Tukey's multiple comparison test confirmed significantly increased A $\beta$ 42 levels in samples with high AD severity when compared to samples with not/low AD ( $p < 0.01$ ) and to samples with intermediate AD severity ( $p < 0.05$ ). Comparison of A $\beta$ 42 levels in the hippocampus (Figure 2c) by ANOVA did not reveal significant effects of NIA-AA AD

severity ( $p > 0.05$ ). Although we saw visible trends for increased A $\beta$ 42 levels in the neostriatum (Figure 2d) of samples with intermediate and high NIA-AA AD severity, ANOVA of A $\beta$ 42 levels in the neostriatum did not reveal statistically significant effects of NIA-AA AD severity ( $p > 0.05$ ).

### Detection of pTau in human FFPE tissue

To determine whether we could use the same antibodies that are routinely used in neuropathology for Luminex-based detection pTau, we extracted RIPA soluble fractions from the MFG from 3–5 additional AD cases that were not part of the 30 cases/controls cohort, and tested increasing amounts of each fraction for the presence of pTau. Increasing amounts of RIPA-soluble protein extracts resulted in increasing fluorescent signal intensities in our pTau assay (capture with AT8 clone, detection with HT7 clone), and these signal intensities were approximately linear for protein amounts between 187.5 ng and 3,000 ng, approximately spanning a 29-fold dynamic range (Figure 3a). Based on the observed input curve, we determined that ~2,000 ng of RIPA soluble proteins allows for a strong pTau signal that is well within the linear range of the assay.

### Quantification of pTau in FFPE tissue from AD patients and controls

We measured levels of pTau in RIPA extracts from the same FFPE MFG, hippocampal and neostriatal tissue that we used for measuring A $\beta$ 42, and then compared pTau levels between samples that were grouped based on NIA-AAA AD severity. ANOVA of MFG pTau levels (Figure 3b) revealed significant effects of NIA-AA AD severity ( $F_{2,26} = 4.96$ ,  $p < 0.01$ ), and *post-hoc* pair-wise comparisons with Tukey's multiple comparison test further showed that pTau levels in the group with high NIA-AA AD severity were significantly higher than in the groups with not/low and intermediate NIA-AA AD severity (each  $p < 0.05$ ). Similarly, comparison of pTau levels in the hippocampus (Figure 3c) by ANOVA revealed significant effects of NIA-AA AD severity ( $F_{2,25} = 3.82$ ,  $p < 0.05$ ). *Post-hoc* pair-wise comparisons with Tukey's multiple comparison test further show that pTau levels in the group with high NIA-AA AD severity were significantly higher than in the group with not/low NIA-AA AD severity ( $p < 0.05$ ). In contrast, comparison of pTau levels in the neostriatum (Figure 3d) by ANOVA did not reveal any significant effects of NIA-AA AD severity (each  $p > 0.05$ ). We noticed that pTau signal in most samples from the neostriatum was at or close to the level of PBS reagent blanks.

### Correlation analysis of A $\beta$ 42 and pTau levels in FFPE sections from AD patients and controls

We performed correlation analyses of A $\beta$ 42 and pTau levels in all analyzed brain regions. A $\beta$ 42 levels in the MFG were correlated positively with A $\beta$ 42 levels in hippocampus (Figure 4a) and neostriatum (Figure 4b), and A $\beta$ 42 levels in the hippocampus were correlated positively with A $\beta$ 42 levels in the neostriatum (Figure 4c). A $\beta$ 42 levels in the neostriatum were correlated positively with pTau levels in the MFG (Figure 4d). pTau levels in the MFG were correlated positively with pTau levels in hippocampus (Figure 4f). Although pTau levels in the neostriatum correlated significantly with A $\beta$ 42 and pTau levels in some other brain regions, these correlations cannot be interpreted because pTau levels in the neostriatum were essentially as low as PBS blanks. All correlations, including not



statistically significant correlations and correlations with pTau in the neostriatum are listed in Table 4.

### **Grey-matter area measurements in all FFPE sections from AD patients and controls**

In order to potentially correct for variability in white matter content in tissue sections (with the vast majority of A $\beta$ 42 and pTau expected in cortex), we measured the area of grey matter in sections that were immediately adjacent to all FFPE sections that we used for protein extractions for the Luminex assays (Supplemental Figure 1). Comparisons of grey matter areas in MFG, hippocampus and striatum by ANOVA did not reveal any significant effects of NIA-AA AD severity ( $F_{2,79} = 2.7$ ,  $p > 0.05$ ).

### **Association of quantitative measurements of A $\beta$ 42 and pTau with genetic risk for AD**

*APOE- $\epsilon$ 4* is the strongest genetic risk factor for sporadic AD with increased risk with one or two copies of the *APOE- $\epsilon$ 4* allele [23]. Our validation cohort included 19 cases without the *APOE- $\epsilon$ 4* and 11 cases with at least one *APOE- $\epsilon$ 4* allele. Comparisons of A $\beta$ 42 levels in carriers and non-carriers of the *APOE- $\epsilon$ 4* allele across all sampled brain regions (Figure 5a) by ANOVA revealed a significant effect of *APOE* genotype on A $\beta$ 42 levels ( $F_{1,82} = 6.13$ ,  $p < 0.05$ ). However, *post-hoc* pair-wise comparisons failed to show significant ( $p > 0.05$ ) mean differences between specific groups. Comparisons of pTau levels in carriers and non-carriers of the *APOE- $\epsilon$ 4* allele across all sampled brain regions (Figure 5b) by ANOVA did not reveal a significant effect of *APOE* genotype on pTau levels ( $F_{1,82} = 0.76$ ,  $p > 0.05$ ).

### **Association of quantitative measurements of A $\beta$ 42 and pTau with ante-mortem cognitive status**

Quantitative measures of AD pathology can be used to probe for associations between A $\beta$ 42 and pTau and cognitive status (No dementia vs. Dementia). We compared A $\beta$ 42 and pTau levels in cases with and without dementia, and found that cases with dementia had significantly higher A $\beta$ 42 and pTau levels in MFG and significantly higher A $\beta$ 42 levels in the neostriatum than non-demented cases ( $p < 0.05$ , Supplemental Figure 2). Hippocampal levels of A $\beta$ 42 and pTau were not significantly higher in demented vs. non-demented cases ( $p > 0.05$ , Supplemental Figure 2); although there was a trend for increased hippocampal pTau in demented subjects. pTau levels in the neostriatum were too low to be included in this analysis.

### **Detection of GFAP and IBA1 in human FFPE tissue**

To determine whether we could use Luminex-based technology to probe for pathologic pathways in FFPE MFG, we first validated GFAP and IBA1 Luminex reagents with commercially available GFAP (Figure 6a) and IBA1 (Figure 6b) peptides. Increasing concentrations of peptides resulted in increasing fluorescent signal intensities in both GFAP (capture with EPR1034Y clone, detection with polyclonal antibody) and IBA1 (capture with EPR16589 clone, detection with GA5 clone) assays, and these signal intensities were approximately linear for peptide concentrations between 5,000 and 16,000 pg/ml for GFAP and between 300 and 40,000 pg/ml for IBA1. Next we extracted RIPA and Gu-HCl soluble fractions from MFG from 3–5 additional AD cases that were not part of our 30 cases/

controls cohort, and tested increasing amounts of each fraction for the presence of GFAP and IBA1. No signals could be detected for GFAP and IBA1 in these fractions (data not shown). However, when we used MFG protein lysates extracted with a commercial FFPE extraction kit (Qiagen), we were able to detect GFAP and IBA1. Increasing amounts of Qiagen protein extracts resulted in increasing fluorescent signal intensities in the GFAP assay, and these signal intensities were approximately linear for protein amounts between 100 ng and 3,200 ng (Figure 6c). Similarly, increasing amounts of Qiagen protein extracts resulted in increasing fluorescent signal intensities in the IBA1 assay, and these signal intensities were approximately linear for protein amounts between 125 ng and 4,000 ng (Figure 6d).

## DISCUSSION

Here we describe the development of a Luminex-based method to generate highly quantitative, molecularly specific measurements of pathologic peptides and reactive and inflammatory proteins in FFPE brain tissue.

A major obstacle to the analysis of FFPE protein lysates is the formation of methylene bridges between side chains of proteins, cross-linking and preserving the peptides of cellular proteins. To establish an FFPE-compatible Luminex assay, we first developed procedures to solubilize A $\beta$ 42 and pTau in protein extracts from FFPE tissue. We based these procedures on immunohistology protocols that are routinely used for antigen retrieval and subsequent immunostaining for A $\beta$ 42 and pTau [13]. We found that deparaffinization followed by citric-acid boil, homogenization in RIPA buffer and sequential extraction with RIPA buffer (solubilizes pTau) and then Gu-HCl (solubilizes A $\beta$ 42) produced protein extracts that could be quantified using standard protein quantification kits and that resulted in robust Luminex detection of A $\beta$ 42 and pTau. As a result, this method allows to standardize downstream analyses through normalization of measured signal intensities to total amounts of protein per analyzed fraction (e.g. RIPA-soluble or Gu-HCl soluble fractions). By varying the amount of total protein analyzed, we obtained signal intensities that were almost perfectly linear over a 35-fold dynamic range for A $\beta$ 42 and almost perfectly linear over a 29-fold dynamic range for pTau, which are in good agreement with several published studies [13, 14, 24–27]. To align with clinical and research neuropathologic assessments in AD, we chose A $\beta$ 42 and pTau antibodies that are currently recommended for standard of care.

We generated a cell-culture based source of A $\beta$ 42 standards that performed as well as commercial A $\beta$ 42 standards when used with a widely used commercial A $\beta$ 42 Luminex assay (INNO-BIA AlzBio3 kit), which then allowed the generation of standard curves for the quantification of A $\beta$ 42 with our own Luminex assay. The relative ease of preparation and the verified shelf life of at least 6 months makes our cell-culture based A $\beta$ 42 standard a useful and cost-effective alternative to commercially available A $\beta$ 42 standards.

We then generated protein preparations from FFPE brain samples obtained from 30 autopsies of research subjects with AD neuropathologic change ranging from not/low to intermediate and high AD pathologic change (NIA-AA criteria). We used samples collected according to NIA-AA recommendations for AD and included MFG, hippocampus and

neostriatum in our analysis [2]. Consistent with the NIA-AA based neuropathologic diagnosis of AD, the Luminex assay detected low levels of A $\beta$ 42 and pTau in MFG samples with low levels of AD neuropathologic change, and higher levels of A $\beta$ 42 and pTau in MFG samples with high levels of AD neuropathologic change. Also in strong agreement with the NIA-AA based neuropathologic diagnosis of AD, Luminex measurements confirmed significantly higher hippocampal pTau levels in cases with high NIA-AA AD neuropathologic change. While the Luminex assay detected upwards trending levels of A $\beta$ 42 in the neostriatum of cases with high NIA-AA AD neuropathologic change, we could not detect significant levels of neostriatal pTau. Taken together, our quantitative Luminex data are in solid agreement with A $\beta$ 42 and pTau neuropathologic changes typical of NIA-AA AD severity. Although we detected strong differences between the not/low and high NIA-AA AD neuropathologic change groups, we only observed trends for increased A $\beta$ 42 and pTau levels in cases with intermediate NIA-AA AD neuropathologic change. Increasing sample sizes in future studies will most likely resolve this issue.

We observed, that the levels of A $\beta$ 42 in MFG were roughly twice as high as in either the hippocampus or the neostriatum, and pTau levels in the hippocampus were almost 10 times higher than in the MFG. Hence, this method not only allows detection of inter-individual disease-associated levels of A $\beta$ 42 and pTau across the same brain regions, but also intra-individual differences between various brain regions and absolute quantitation of target protein/peptide when standards are available.

We performed a simple and preliminary analysis of the association between quantitative assessments of A $\beta$ 42 and pTau with ante-mortem clinical diagnosis of dementia, and found elevated levels of A $\beta$ 42 and pTau in samples from demented individuals. We are aware that our validation cohort has not been specifically selected for such an association study, but are encouraged by this current finding to further explore this issue in a larger future investigation. Similarly, our observation that carriers of the *APOE*- $\epsilon$ 4 allele had elevated A $\beta$ 42 levels is interesting, but limited by the small sample size and the nature of the validation cohort.

Our correlation analyses showed that A $\beta$ 42 levels in all brain regions correlated positively with A $\beta$ 42 levels in all other brain regions, and that pTau levels in MFG and hippocampus were also positively correlated. Only A $\beta$ 42 levels in the neostriatum were weakly correlated with pTau levels in the MFG, presumably reflecting advanced progression of overall AD neuropathologic change associated with neostriatal A $\beta$ 42 pathology [1, 2]. These preliminary findings warrant further investigations with larger study cohorts.

Hence, this highly quantitative, molecularly specific Luminex-based assay will be useful for future studies correlating pathology with genetic and environmental risk factors, clinical and biomarker profiles, and therapeutic interventions. Future applications may specifically include investigation of pathologic changes related to cognitive resilience/reserve to the effects of elevated A $\beta$ 42 and pTau levels or to the pathological and phenotypic heterogeneity of AD. Applying this method to specific hippocampal regions, including CA1 and subiculum, and entorhinal cortex can also help differentiate regional pathology in

association with phenotypic outcomes, genetic and environmental risk, and therapeutic interventions.

We envision combining IHC assessments of histopathology in adjacent sections to correlate quantitative findings when needed, and expect this method to complement histologic data sets with standardized and highly quantitative measures of specific markers of neurodegeneration. The combination of a high throughput, low cost approach that uses very little tissue to quantify, in a molecularly specific manner, antigenic targets in banked FFPE tissues will enable diverse studies in AD, but with applications in other diseases of the central nervous system and other organ systems that have large FFPE local or national resources and are not readily accessible through other means.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## Abbreviations:

<b>AD</b>	Alzheimer's disease
<b>A<math>\beta</math>42</b>	amyloid $\beta$ 42
<b>pTau</b>	phospho-Tau
<b>IHC</b>	immunohistochemistry
<b>SA</b>	streptavidin
<b>PA</b>	phycoerythrin
<b>FFPE</b>	formalin-fixed, paraffin-embedded

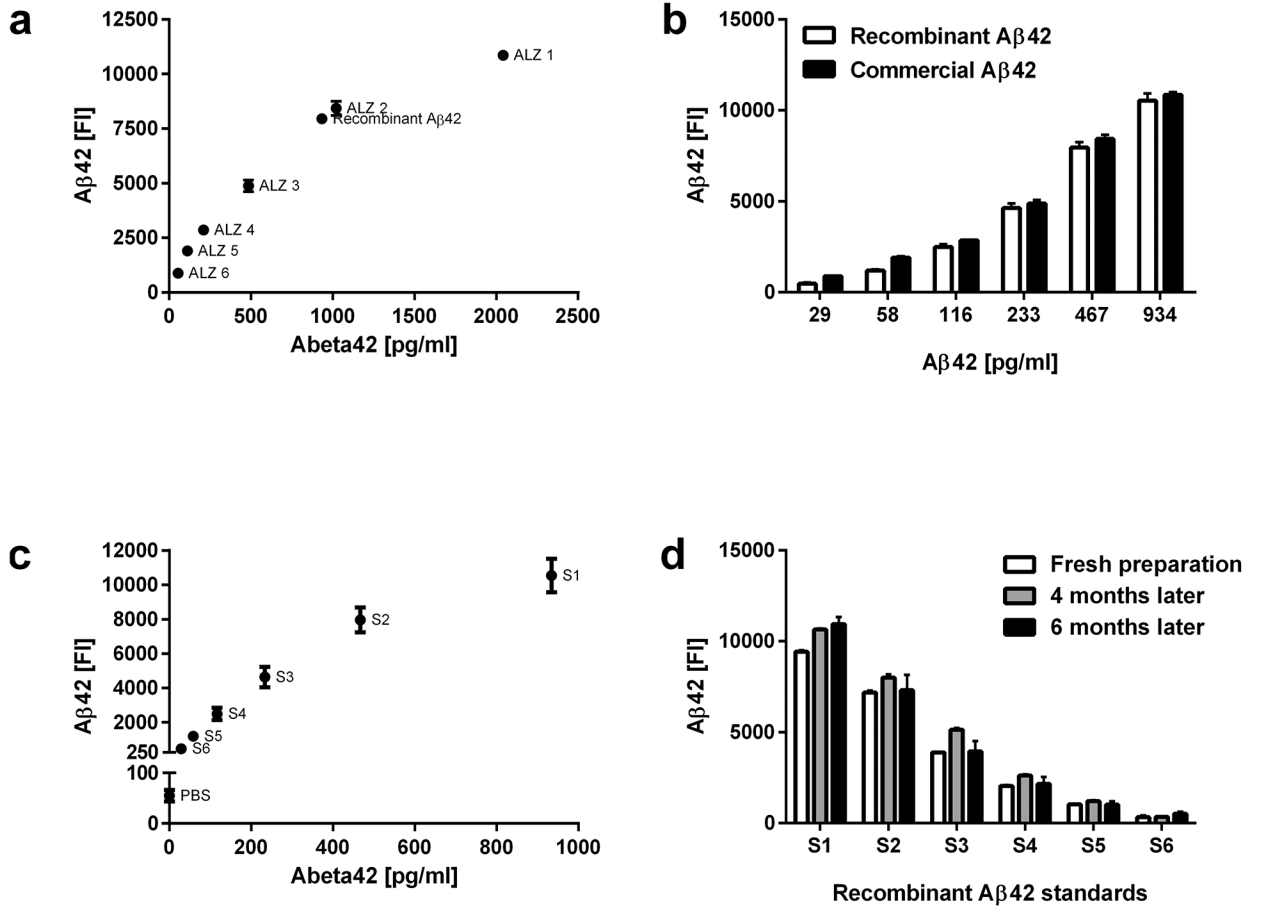
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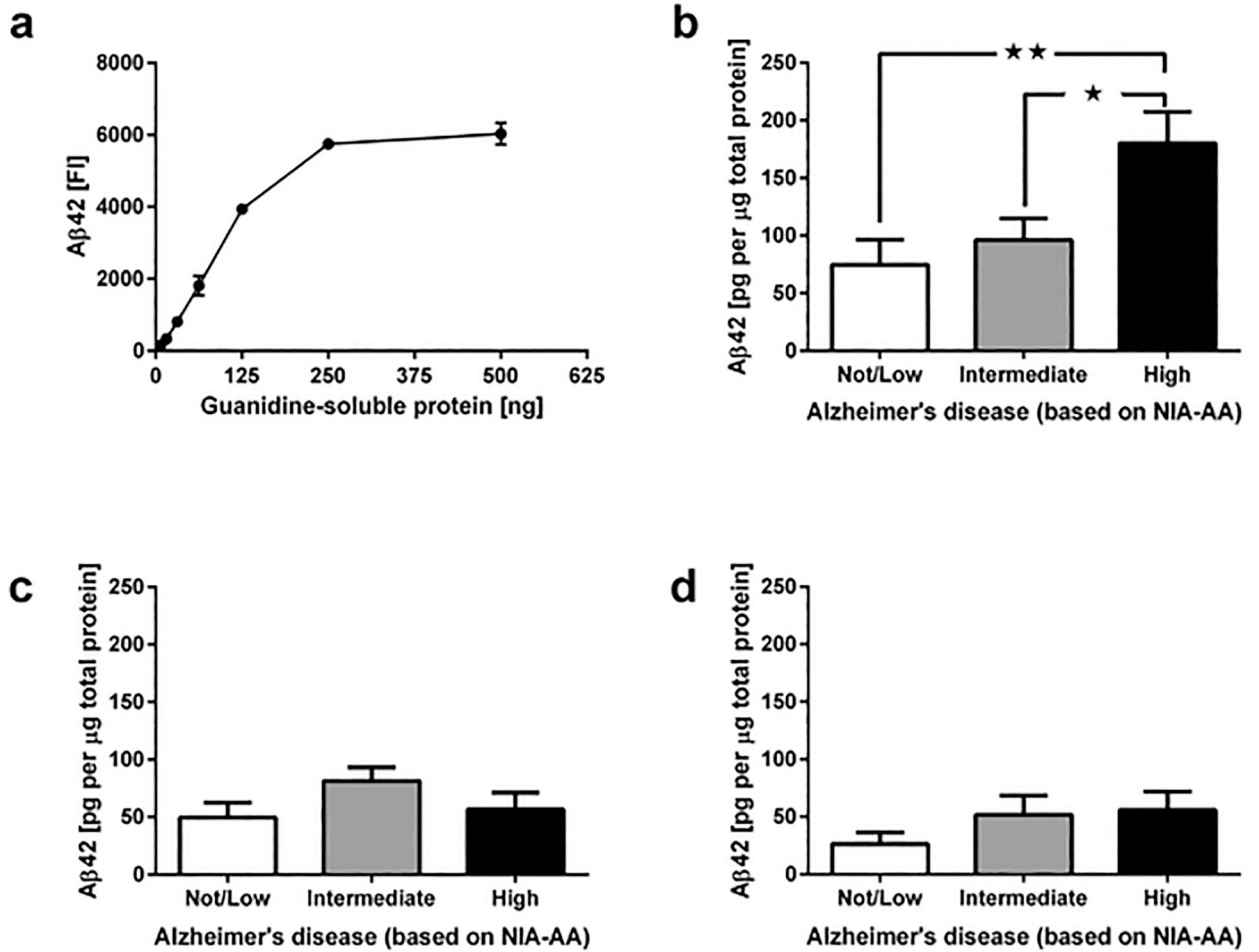
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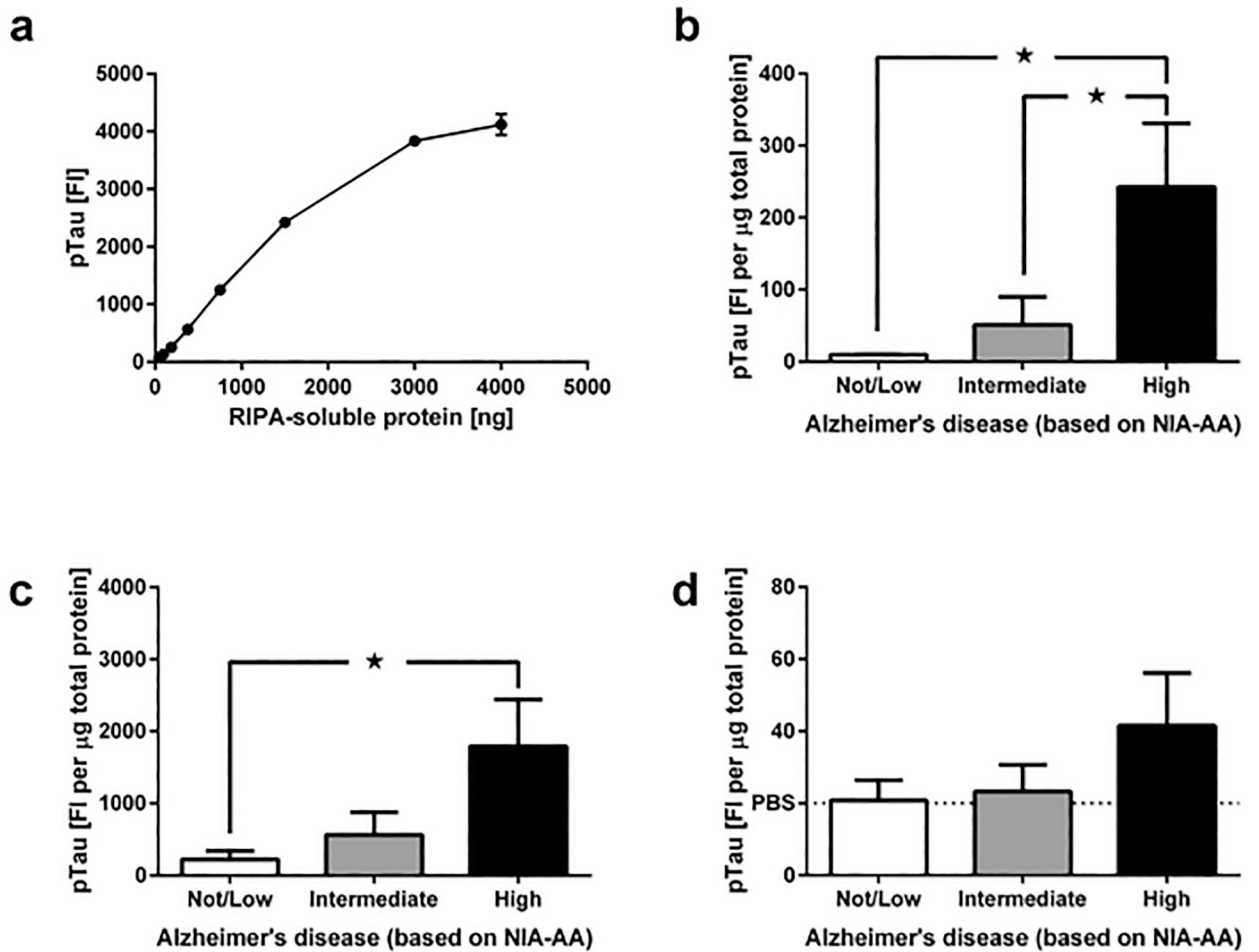


**Figure 1.** Validation of recombinant Aβ42 standards. **(a)** Quantification of recombinant Aβ42 using commercial Innogenetics-kit standards and the Innogenetics kit. **(b)** Comparison of serial dilutions of Innogenetics-kit and recombinant Aβ42 standards measured with the Innogenetics kit. **(c)** Generation of a standard curve with recombinant Aβ42 standards measured with our newly developed Luminex assay. **(d)** Test-retest of serially diluted recombinant Aβ42 standards measured with our newly developed Luminex assay. All data are presented as mean ± SEM.



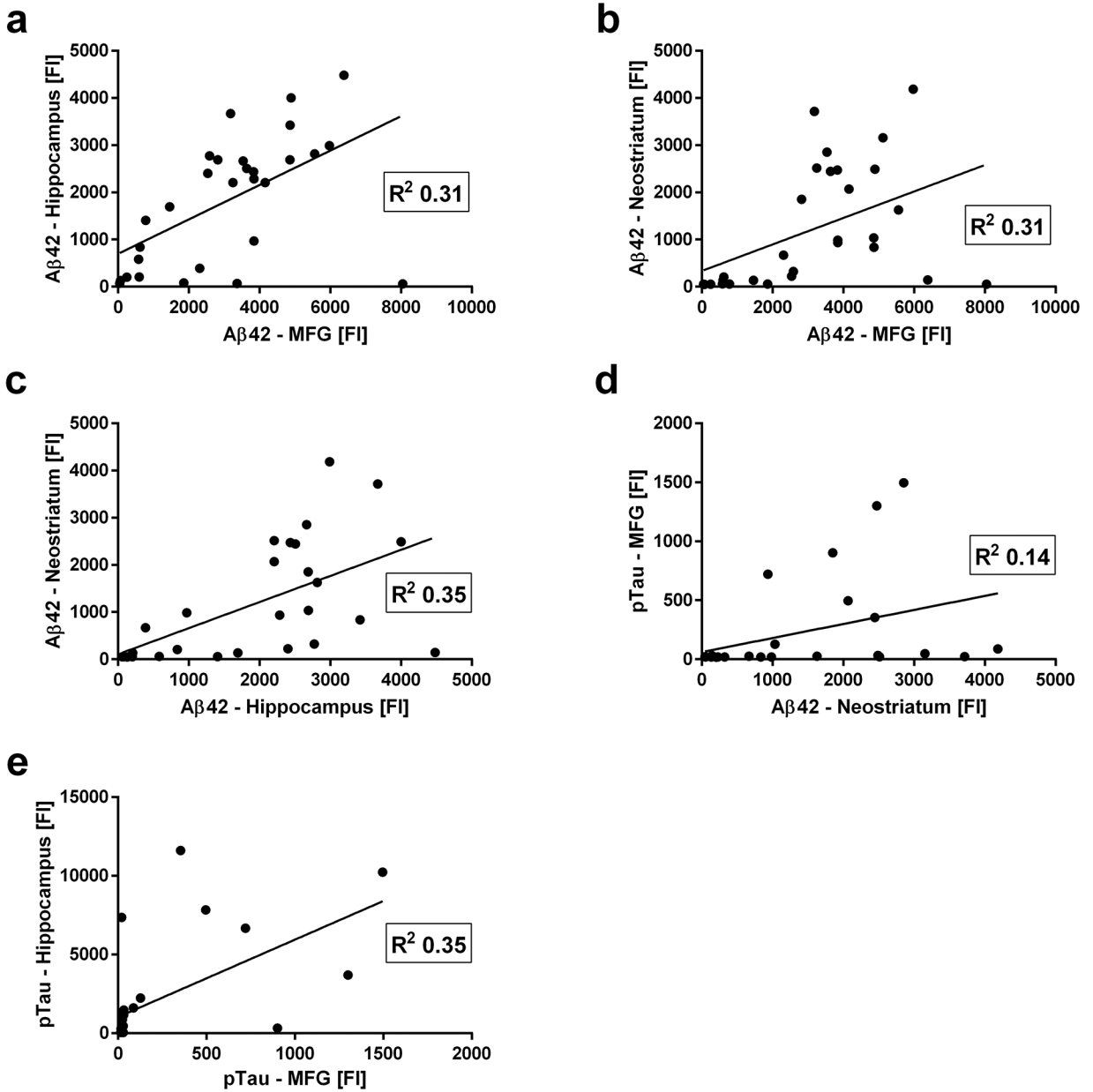
**Figure 2.**

Determination of optimal FFPE sample amounts for Luminex analysis of Aβ42, and quantification of Aβ42 in FFPE tissue from cases with Not/Low, Intermediate and High NIA-AA AD severity. **(a)** Fluorescent-intensity signals for Aβ42 as function of increasing amounts of guanidine-soluble extracts from middle frontal gyrus. **(b)** Aβ42 levels in MFG samples from cases with varying NIA-AA AD severity (Not/Low: n = 10, Intermediate: n = 10, High: n = 9). **(c)** Aβ42 levels in hippocampal samples from cases with varying NIA-AA AD severity (Not/Low: n = 10, Intermediate: n = 10, High: n = 9). **(d)** Aβ42 levels in neostriatal samples from cases with varying NIA-AA AD severity (Not/Low: n = 10, Intermediate: n = 10, High: n = 9). Aβ42 levels in panels **(b-d)** are normalized to the amount of guanidine-soluble protein (pg per μg). All data are presented as mean ± SEM. ★ p < 0.05 and ★★ p < 0.01.

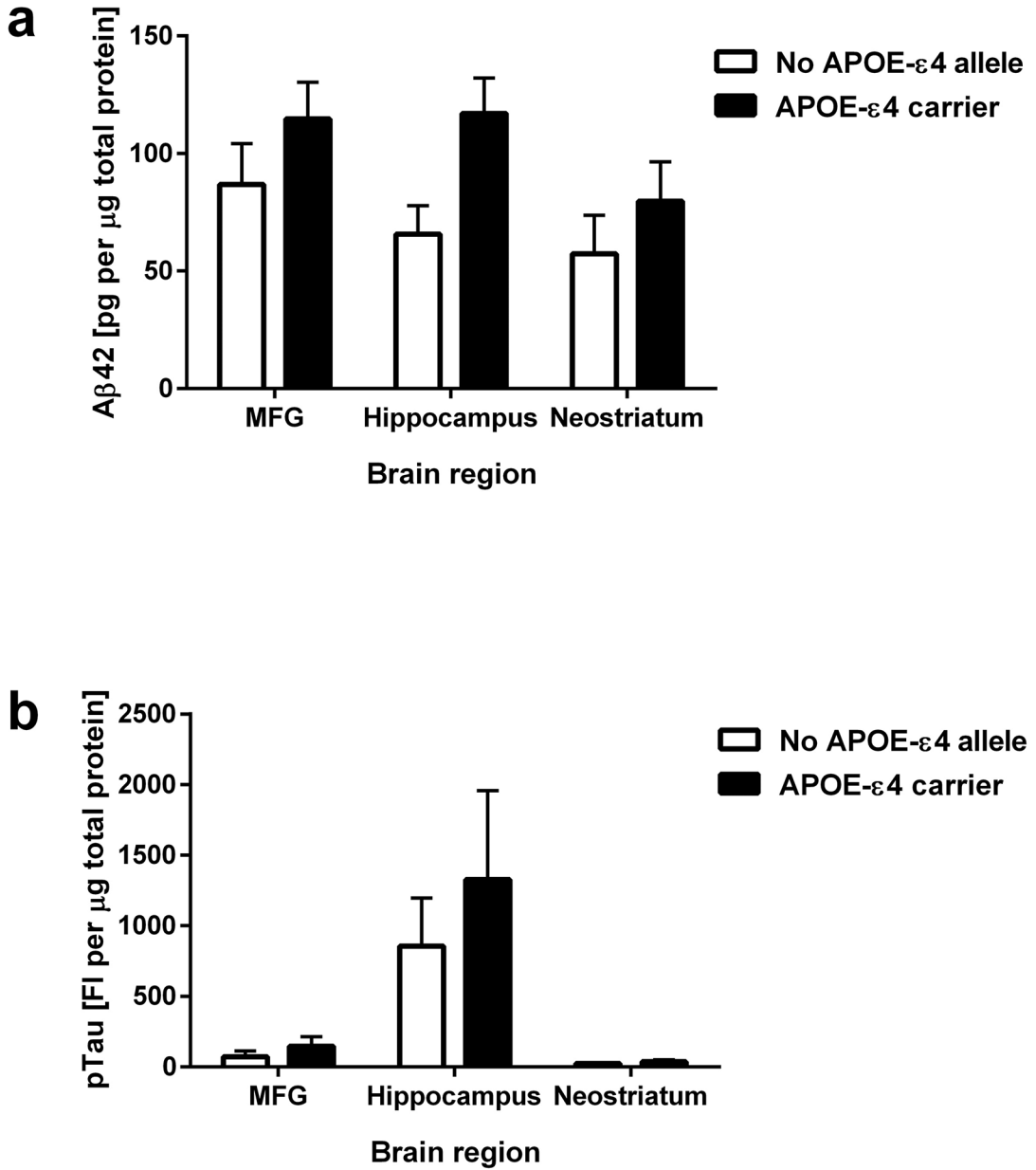


**Figure 3.**

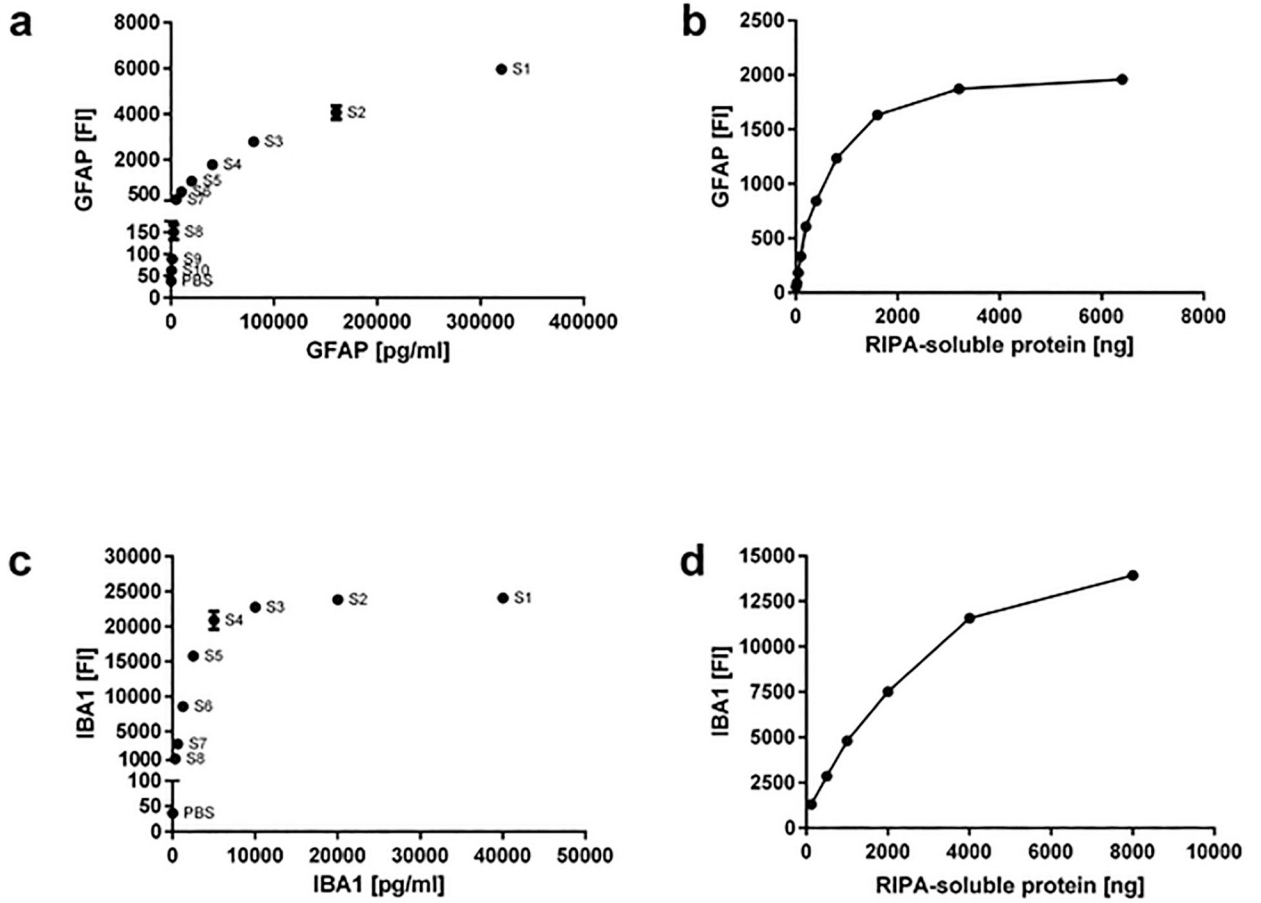
Determination of optimal FFPE sample amounts for Luminex analysis of pTau, and quantification of pTau in FFPE tissue from cases with Not/Low, Intermediate and High NIA-AA AD severity. **(a)** Fluorescent-intensity signals for pTau as function of increasing amounts of RIPA-soluble extracts from middle frontal gyrus. **(b)** pTau levels in MFG samples from cases with varying NIA-AA AD severity (Not/Low: n = 10, Intermediate: n = 10, High: n = 9). **(c)** pTau levels in hippocampal samples from cases with varying NIA-AA AD severity (Not/Low: n = 10, Intermediate: n = 10, High: n = 9). **(d)** pTau levels in neostriatal samples from cases with varying NIA-AA AD severity (Not/Low: n = 10, Intermediate: n = 10, High: n = 9). pTau levels in panels **(b-d)** are normalized to the amount of RIPA-soluble protein (fluorescent intensity per µg). All data are presented as mean  $\pm$  SEM. ★  $p < 0.05$  and ★★  $p < 0.01$ .



**Figure 4.** Scatter plots of Luminex fluorescent signal intensities for Aβ42 and pTau in MFG, hippocampus and neostriatum. We performed correlation analyses of Aβ42 and pTau signals across all analyzed brain regions and confirmed the following statistically significant positive correlations. **(a)** Aβ42 levels in MFG vs. Aβ42 levels in hippocampus. **(b)** Aβ42 levels in MFG vs. Aβ42 levels in neostriatum. **(c)** Aβ42 levels in hippocampus vs. Aβ42 levels in neostriatum. **(d)** Aβ42 levels in neostriatum vs. pTau levels in MFG. **(e)** pTau levels in MFG vs. pTau levels in hippocampus. Statistically significant  $R^2$  values are shown in each scatter plot.



**Figure 5.** Association of Aβ42 and pTau levels with genetic risk for AD. **(a)** Aβ42 and **(b)** pTau levels in MFG, hippocampus and neostriatum of samples from cases with no *APOE-ε4* allele (n = 18) and with at least one *APOE-ε4* allele (n = 11). All data are presented as mean ± SEM.





**Table 1.**

Neuropathologic, clinical and demographic characteristics of subjects.

	<b>Group</b>	<b>Number of Subjects</b>	<b>Age in years (Mean <math>\pm</math> SD)</b>	<b>PMI in hours (Mean <math>\pm</math> SD)</b>	<b>Sex (%female)</b>
Genetic Clinical Neuropathology	Not/low AD	10	83.5 $\pm$ 10.51	5.03 $\pm$ 1.59	70
	Intermediate AD	10	89 $\pm$ 5.21	4.48 $\pm$ 1.73	60
	High AD	9	86.4 $\pm$ 5.36	5.26 $\pm$ 1.53	80
	No dementia	12	85.78 $\pm$ 8.50	4.90 $\pm$ 1.59	83.3
	Dementia	17	87.08 $\pm$ 6.08	4.96 $\pm$ 1.60	58.8
	No <i>APOE-ε4</i> allele	18	88.95 $\pm$ 6.22	4.80 $\pm$ 1.63	61.1
	1 <i>APOE-ε4</i> allele	11	81.73 $\pm$ 7.67	5.13 $\pm$ 1.51	81.8

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**Table 2.**

Antibodies that were conjugated to magnetic beads to capture antigens.

Antigen	Product code	Manufacturer	Host	Clone
A $\beta$ 42	700254	Invitrogen	Rabbit	H31L21
pTau	MN1020	Invitrogen	Mouse	AT8
GFAP	ab218309	Abcam	Rabbit	EPR1034Y
IBA1	ab221790	Abcam	Rabbit	EPR16589

Abbreviations: Amyloid beta 42 (A $\beta$ 42), phospho-tau (pTau), glial fibrillary acidic protein (GFAP), ionized calcium binding adapter molecule 1 (IBA1).

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**Table 3.**

Antibodies that were biotinylated to detect captured antigens.

Antigen	Product code	Manufacturer	Host	Clone
A $\beta$ 42	803001	Biologend	Mouse	6E10
Tau	MN1000	Invitrogen	Mouse	HT7
GFAP	LS-B2645-50	LS BioSciences	Goat	NA (polyclonal)
IBA1	MAB3402B	EMD Millipore	Mouse	GA5

Abbreviations: Amyloid beta 42 (A $\beta$ 42), glial fibrillary acidic protein (GFAP) and ionized calcium binding adapter molecule 1 (IBA1).

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**Table 4.**

Correlations ( $R^2$ ) between A $\beta$ 42 and pTau levels in MFG, hippocampus and neostriatum in all analyzed samples. Statistically significant correlation coefficients are marked: ★  $p < 0.05$  and ★★  $p < 0.01$ .

	A $\beta$ 42			pTau			
	MFG	HPC	NEO	MFG	HPC	NEO	
A $\beta$ 42	MFG	-	0.31★★	0.21★	0.01	0.01	0.09
	HPC	-	-	0.35★★	0.06	0.02	0.01
	NEO	-	-	--	0.14★	0.11	0.30★★
pTau	MFG			-	0.35★★	0.23★★	
	HPC				-	0.23★★	
	NEO					-	

Abbreviations: Amyloid beta 42 (A $\beta$ 42), phospho-tau (pTau), middle frontal gyrus (MFG), hippocampus (HPC) and neostriatum (NEO).