# Relevance of the interplay between amyloid and tau for cognitive impairment in early Alzheimer's disease

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# Journal Requirements

- 1. Regular manuscripts should be **no longer that 10 printed journal pages (30 doubled-spaced pages, including references, figures and tables)** and should include only the most essential figures and tables
- 2. Divide your article into clearly defined and numbered sections. Subsections should be numbered 1.1 (then 1.1.1, 1.1.2, ...), 1.2, etc. use this numbering also for internal cross-referencing
- 3. State the objectives of the work and provide an adequate background, avoiding a detailed literature survey or a summary of the results.
- 4. Avoid extensive citations and discussion of published literature.
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**ABSTRACT** 

Amyloid  $\beta$  (A $\beta$ ) and tau are key hallmark features of Alzheimer's disease (AD) neuropathology.

The interplay of AB and tau for cognitive impairment in early AD was examined with cross-

sectional analysis, measured by cerebrospinal fluid (CSF) biomarkers (Aβ<sub>1-42</sub>, total tau [t-tau] and

phosphorylated tau [p-tau<sub>181P</sub>]), and on cognitive performance by the repeatable battery for

assessment of neuropsychological status (RBANS). Participants (n=246) included cognitively

normal (A $\beta$ -), mild cognitively impaired (A $\beta$ -), preclinical AD (A $\beta$ +), and prodromal AD

 $(A\beta+)$ . Overall, cognitive impairment (RBANS Total Scale score) had a moderate negative

correlation to t-tau (n=246; r=-0.434; p<0.001) and p-tau<sub>181P</sub> (r=-0.389; p<0.001). When

classified by A $\beta$  status, this correlation to t-tau was applicable only in A $\beta$ + participants (n=139;

r=-0.451, p<0.001) but not A $\beta$ - participants (n=107; r=0.137, p=0.16), with identical findings

for p-tau. Both tau (p<0.0001) and interaction of A $\beta_{1-42}$  with tau (p=0.006) affected RBANS, but

not  $A\beta_{1-42}$  alone. Cognitive/memory performance correlated well with CSF tau levels across early

stages of AD, although the correlation is Aβ dependent.

Keywords: Alzheimer's disease, amyloid, tau, interaction, RBANS, cognition

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#### 1. Introduction

The hallmark pathological changes of Alzheimer's disease (AD), amyloid- $\beta$  (A $\beta$ ) plaques and tau neurofibrillary tangles (NFT) are known to start accumulating years before the onset of clinical symptoms (Dubois et al., 2016; Villemagne et al., 2013), with a putative sequence of A $\beta$  events preceding tau. With the molecular definition of the disease (Jack et al., 2018), the progressing clinical disorder of AD is considered a continuum with a seamless sequence of events from healthy elderly individuals with positive biomarker signature and impairments gradually increasing over time to progress from milder stages of cognitive deficits to functional impairment and dementia (Aisen et al., 2017; Jack et al., 2013). This interpetation of AD was enabled by advances in cerebrospinal fluid (CSF) and imaging biomarkers such as A $\beta$  and tau to move from a clinically defined late stage disease to a biological definition of a disease process that starts many years before onset of clinical symptoms. These research avances have resulted in a new set of criteria for AD to define the earliest stages in the AD continuum, *i.e.*, cognitively normal individuals with an abnormal AD biomarker pattern, being preclinical AD (Dubois et al., 2014; Knopman and Caselli, 2012; Vos et al., 2013).

The amyloid cascade hypothesis brings forward the deposition of  $A\beta$  as a result of increased production and/or decreased clearance of  $A\beta$  from the central nervous system, as the initial pathological trigger in the disease continuum, interacting with the microtubule-binding protein tau, forming NFT leading to widespread neuronal degeneration and dysfunction, cognitive decline and dementia (Karran et al., 2011). Genetic evidence from rare familial forms of AD as well as mutations in amyloid- $\beta$  precursor protein (APP) and presentlins 1 and 2 genes support the accumulation of  $A\beta$  as the causative factor for AD (Levy-Lahad et al., 1995; Rogaev et al., 1995; Sherrington et al., 1995).

While genetic evidence points towards  $A\beta$  as the initiator of pathology, the eventual clinical syndrome appears to be more closely related to the progression of the tau pathology. The rate of cognitive impairment in AD and neuronal loss however strongly correlate spatially and temporally with the total tau burden and NFT formation (Nelson et al., 2012; Pontecorvo et al., <u>2017</u>). While A $\beta$  pathology spreads inwards starting from the cortex, progression of tauopathy in AD topographically disconnects and goes in the opposite direction (Braak et al., 2006). In animal models, both A\beta and tau have shown self-propagation capabilities (Jucker and Walker, 2011, 2013), but whether this occurs in humans is not known. Extracellular phospho-tau (p-tau) seeds are known to induce tauopathy in neurons, supporting the hypothesis that tauopathy can spread in a prion-like fashion between neurons, a cascade that could potentially be initiated or accelerated by Aß pathology (Goedert et al., 2017). Despite the prominent correlation of NFTs with cognitive impairment in AD, mutations in tau are not causative for AD but may result in dementia, without an AD clinical or neuropathological phenotype (Hardy and Selkoe, 2002). Increasing preclinical evidence supports a possible interplay between A $\beta$  and tau, whereby A $\beta$ may exert its toxicity through tau by influencing or inducing the hyperphosphorylation, misfolding and secretion of tau (Musiek and Holtzman, 2015; Nisbet et al., 2015) and tau may further enhance Aβ toxicity via a feedback loop. In addition, preclinical data demonstrates that tau is required for learning and memory deficits in the presence of Aβ (Roberson et al., 2007). Aβ is being proposed as the trigger to AD, which may induce the cascade of downstream events including tauopathy, and tau as the bullet, driving the neurodegeneration and cognitive deficits (Bloom, 2014). This potential interplay would further strengthen the interaction between both key hallmark features of the amyloid cascade hypothesis already at the start of the AD continuum, hypothesizing that the relation of CSF tau to cognitive performance is dependent on A $\beta$  pathology as measured by CSF A $\beta$  levels.

The interaction of  $A\beta$  and tau, measured by means of CSF biomarkers ( $A\beta_{1-42}$ , total tau [t-tau] and p-tau<sub>181P</sub>), on cognitive performance as a clinical marker in normal participants and those in the earliest stages of the AD continuum defined as preclinical and prodromal AD (pAD) was investigated.

#### 2. Methods

# 2.1. Study population

Participants screened from two separate clinical trials in early Alzheimer's disease (Study 1: NCT01978548 (<u>Timmers et al., 2018</u>); Study 2: NCT02260674) were included in this study. Both studies were conducted in sites across Europe in Caucasian participants between December 2013 and June 2016.

# 2.2. Study participants

A total of 256 participants (Study 1: n=112; Study 2: n=144), aged 50 to 90 years inclusive, who completed a standardized 4-step screening process for either studies to assess their eligibility according to predefined inclusion and exclusion criteria (Supplemental fig S1), were included in this study (Timmers et al., 2018). Of the 256 participants who completed the screening process, a total of 10 participants were further excluded from final analysis, due to incorrect, incomplete or missing RBANS data recorded (n=7), missing p-tau<sub>181P</sub> data (n=1) or no CDR data (n=2) available, resulting in 246 evaluable participants.

Two paradigms were used to classify participants included (n=246) as depicted in Supplemental Fig. S2.

#### 2.2.1. Biomarker CSF $A\beta_{1-42}$ and cognitive status (CDR).

Participants were identified as "preclinical AD" (CDR=0 and A $\beta$  < 600ng/L [A+] n= 33), prodromal Alzheimer's disease (pAD; CDR  $\geq$  0.5 but no dementia; CDR=0.5 [n=104] and 1.0

[n=2], together with  $A\beta$  < 600ng/L [A+], n=106), "mild cognitive impairment" (MCI; CDR=0.5 and normal  $A\beta$  levels [A-]( $\geq$ 600 ng/L), n=38) or "control" (CDR=0 and normal  $A\beta$  levels [A-], n=69).

# 2.2.2. Biomarker status alone independent of clinical stage; i.e., by $A\beta$ (A-/A+) and tau (T-/T+) status.

Participants were considered T+ if CSF t-tau and/or p-tau<sub>181P</sub> were above the pre-defined cut-off (t-tau >350 ng/L; p-tau<sub>181p</sub>levels >70 ng/L). All participants positive for p-tau<sub>181P</sub> were positive for t-tau given the strong correlation between both markers independent of A $\beta$  status (A-: r= 0.95, p<0.001; A+: r=0.95, p<0.001; Supplemental Fig. S3). Overall, n=70 were A-/T-, n=37 were A-/T+, n=23 were A+/T-, and n=116 were A+/T+.

#### 2.3. Assessments

#### 2.3.1. Mini-Mental State Examination

The MMSE (Folstein et al., 1975) was used during screening as a tool to screen dementia.

## 2.3.2. Clinical Dementia Rating Scale

The Clinical Dementia Rating Scale (CDR), a semi-structured interview of patients and informants, was used to measure a patient's current dementia status (Morris et al., 1997). The CDR demonstrates good reliability (Burke et al., 1988) and has been validated against neuropathological findings (Berg et al., 1993).

2.3.3. Repeatable Battery for the Assessment of Neuropsychological Status (RBANS)

The RBANS (Randolph et al., 1998) was performed twice by participants during screening

(FORM A and FORM B) to explore test-retest reliability. For the purpose of this study, the first completion of the RBANS (either FORM A or B) was included in the analysis.

# 2.3.4. Collection and analysis of CSF $A\beta_{1-42}$ , p-tau<sub>181p</sub> and t-tau Levels

All participants had a baseline CSF sample (12 mL) collected during screening by single lumbar puncture between the L3 and L4 or L4 and L5 intervertebral space. CSF samples were collected in polypropylene tubes to avoid adsorbance of proteins to test tube walls. CSF samples were mixed to avoid possible gradient effects, aliquoted, frozen and stored at -80°C immediately after collection. All samples analyzed in this study had at most 1 freeze-thaw cycle.

CSF samples were analyzed at one central lab (Sahlgrenska University hospital, Sweden) for  $A\beta_{1-42}$ , phosphorylated tau at position threonine 181 (p-tau<sub>181P</sub>) and total tau (t-tau) concentrations using INNOTEST® Phospho-TAU<sub>181</sub>, INNOTEST® hTAU Ag, and INNOTEST®  $\beta$ -AMYLOID<sub>1-42</sub> (Fujirebio, Ghent, Belgium) (Palmqvist et al., 2014; Timmers et al., 2018), following stringent protocols for quality control of analyses. Diagnostic threshold CSF concentrations for AD versus normal control for  $A\beta_{1-42}$  (CSF  $A\beta_{1-42}$  levels below cut-off value of 600 ng/L) were applied to the current sample set to assess the likelihood of having cerebral  $A\beta$  plaque deposition (A+) (Palmqvist et al., 2014; Timmers et al., 2018). In addition, elevated CSF t-tau (>350 ng/L) and/or p-tau181p levels (>70 ng/L) respectively (T+) as set by the analysis lab at the start of Studies 1 and 2 were applied to the current sample set to assess the likelihood of having neuronal injury and/or cerebral tau pathology (T+).

# 2.4. Statistical analyses

Statistical analyses were performed using SAS Software version 9.4 (SAS Institute). For categorical data, group differences were analyzed using Fisher's Exact test. For numerical data when there were no clear departures from a normal distribution (age, height, weight, and BMI), balance across groups was evaluated with an F-test and pairwise group comparisons were analyzed by Student's t-test (equal variances) or Satterthwaite test (unequal variances). Nonparametric statistics (Hollander, 1999) were used for skewed or truncated data (A $\beta_{1-42}$ , t-tau,

p-tau<sub>181P</sub>, RBANS, and MMSE). Balance across groups was evaluated with a Kruskal-Wallis test. For pairwise comparisons, Hodges-Lehmann estimates for median differences between groups were calculated. Confidence intervals (CIs) and nominal p-values for the pairwise differences between groups were based on the Wilcoxon Rank Sum test. The choice of parametric versus nonparametric methods was driven by normality versus non-normality of data distributions. Spearman correlation coefficients were calculated to evaluate the strength of association between RBANS indexes and tau (t-tau and p-tau<sub>181P</sub>). Multiple linear regression models were employed to explore the relationship between cognition (RBANS) and biomarkers ( $A\beta_{1-42}$ , t-tau and the interaction of  $A\beta_{1-42}$  with t-tau). Similar models using p-tau<sub>181P</sub> instead of t-tau were analysed. Both t-tau and p-tau<sub>181P</sub> were not included together in the same model due to colinearity. Rank-transformed values were used for RBANS and biomarker variables in multiple regression models because assumptions for normality in the model residuals were not met.

#### 3. Results

#### 3.1. Demographic, biomarker and cognitive characteristics

Two paradigms were used to classify participants as depicted in Supplemental Fig. S2, i.e., by  $A\beta_{1-42}$  and cognitive status (CDR) and by biomarker status alone indendent of clinical stage. i.e., by  $A\beta_{1-42}$  and tau (T-/T+) status (see 2.2.1 and 2.2.2). For both classifications, the demographic, cognitive (RBANS, CDR, MMSE) and biomarker characteristics ( $A\beta_{1-42}$ , p-tau<sub>181p</sub>, t-tau) are summarized in Table 1A and Table 1B. The distribution of the overall population enrolled based on their  $A\beta$  and tau biomarker profiles is shown in Supplemental Fig. S4 (A:  $A\beta$  and p-tau<sub>181p</sub>; B:  $A\beta$  and t-tau). Gender distribution was equal across groups independent of classification used (by  $A\beta_{1-42}$  and cognitive status or biomarker only). A tendency towards higher representation of male subjects in the A+/T- vs the A+/T+ group was observed which was in line with earlier reports

(<u>Paquet et al., 2018</u>). Participants classified as pAD and A+/T+ had lower weight and BMI and participants classified as A-/T- were younger.

#### 3.2. Prodromal AD, preclinical AD, control and MCI groups

No differences in CSF A $\beta_{1-42}$  levels were observed amongst A+ groups (pAD vs preclinical AD) or amongst A- groups (MCI vs control). Tau levels (t-tau; p-tau<sub>181P</sub>;median [range]) were significantly increased in both pAD (644.5 ng/L [161;1780]; 86.5 ng/L [29;197] vs control and MCI p<0.001) and preclinical AD (404 ng/L [118;1050] vs control p<0.05, vs MCI p<0.01; and 59 ng/L [23;128] vs control p<0.05, vs MCI p<0.01) compared with control (324 ng/L [152;742]; 50 ng/L [25;116]) and MCI groups (300 ng/L [100;661]; 46 ng/L [20;103]). No differences in CSF tau (t-tau or p-tau<sub>181P</sub>) levels were observed between the control and MCI group (Table 1A). The pAD group showed significantly increased tau levels compared with the preclinical AD group (p<0.001 for both t-tau and p-tau<sub>181P</sub>) (Table 1A).

Overall, pAD participants presented with a lower (median [range] global MMSE score (26 [14;30], p<0.001), and a lower RBANS Total Scale score (74 [47;124], p<0.001; Table 1A, Fig. 1A) and significant impairment on all RBANS index scores (Supplemental Table S1; p<0.001) compared with the control and preclinical AD groups, and for most of the RBANS index scores compared with the MCI group. The pAD participants showed highest impairment (lowest score) on the Delayed Memory Index followed by the Attention Index, Immediate Memory Index, Language Index and Visuospatial Index. The preclinical AD group showed a normal cognitive profile on the RBANS Total Scale and Index scores (Table 1A, Fig. 1A, Supplemental Table S1), while the MCI group had a significantly (p<0.001) lower RBANS Total Scale score (Table 1A; Fig. 1A) and index scores (Supplemental Table S1) compared with the preclinical AD and control group. The MCI group showed the highest impairment on the Attention Index, followed by Delayed Memory, Immediate Memory, Language, and Visuospatial Index.

#### 3.3. A-/T-, A-/T+, A+/T-, A+/T+ groups

High tau values (p-tau<sub>181P</sub> and t-tau) were observed only in the A+ participants. Significantly (p<0.001) higher t-tau and p-tau<sub>181P</sub> levels (median [range]) were observed in the A+/T+ group (637 ng/L [354; 1780]; 85 ng/L [49; 197]) compared with the A-/T+ group (458 ng/L [353; 742]; 69 ng/L [46; 116]; Table 1B). Aβ levels did not differ between the A+/T+ (381 ng/L [159;587]) and A+/T- (421 ng/L [259;590]) group. Slightly higher levels of Aβ were observed in the A-/T+ (903 ng/L [607;1580]) vs the A-/T- (790 ng/L [600;1180]) group (p<0.001). Participants characterized as A-/T+ are considered participants with Suspected Non-Aβ Pathology (SNAP) (Supplemental Fig. S4A and S4B).

The A+/T+ participants presented with a significantly lower MMSE (26 [14;30], p<0.001) (Table 1B), RBANS Total Scale score (76 [47;135], p<0.001; Table 1B and Fig. 1B) and RBANS Index scores (except visuospatial index) compared with all the other groups (p<0.001; Supplemental Table S2). The highest impairment (lowest score) was observed on Delayed Memory index, followed by Attention, Immediate Memory, Language Index, and Visuospatial Index (no significant difference with the A+/T- group; Supplemental Table S2.) The A+/T-, A-/T+ or A-/T- group did not differ from each other based on the RBANS Total Scale score or RBANS Index scores.

3.4. Correlation between CSF tau levels (t-tau and p-tau<sub>181P</sub>) and cognitive impairment

In the overall group, the RBANS Total Scale score showed a moderate negative correlation to t-tau (n=246; r=-0.434; p<0.001; Fig. 2A and Table 2) and p-tau<sub>181P</sub> (n=246; r=-0.389; p<0.001

[Table 2]). However, when classified by Aβ status, the RBANS Total Scale score showed a moderate negative correlation to t-tau in the A+ participants (n=139; r=-0.451, p<0.001) and not in the A- participants (n=107; r=0.137, p=0.16) (Fig. 2B). The individual RBANS index scores (Table 2) showed similar observations with only significant negative correlations with t-tau

levels in the A+ group and not in the A− group with the strongest negative correlation observed in the A+ group for the attention index (r=-0.438, p<0.001) and delayed memory index (r=-0.431, p<0.001), followed by the immediate memory index (r=-0.327, p<0.001), language (r=-0.261, p<0.01) and visuospatial index (r=-0.254, p<0.01). Similar findings were observed for p-tau<sub>181P</sub> (Table 2).

Both t-tau (p<0.0001) and the interaction of  $A\beta_{1-42}$  with t-tau (p=0.006) were statistically significant in their effect on RBANS, but not  $A\beta_{1-42}$  (Supplemental Table S3A). When the regression included independent variables for age and sex (0 for female, 1 for male), again both t-tau (p<0.0001) the interaction of  $A\beta_{1-42}$  with t-tau were significant (p=0.0127), but not the other variables (Supplemental Table S3B). APOE  $\epsilon$  4 carrier status was not included in the model since it was only available for the A+ participants. Similar findings were observed for p-tau<sub>181P</sub> (Supplemental Tables S3C and S3D).

#### 4. Discussion

We examined the role of tau and its possible interplay with  $A\beta$ , measured based on CSF biomarkers ( $A\beta_{1-42}$ , t-tau and p-tau<sub>181P</sub>), on cognitive performance (RBANS) as a clinical marker in participants in the early AD continuum ranging from normal to preclinical AD and pAD. Original screening paradigm allowed classification of participants based on their clinical function and evidence of  $A\beta$  pathology along the early stages of the AD continuum. In addition, across the early AD continuum, CSF  $A\beta$  levels were shown to plateau while tau levels steadily increased with increased cognitive impairment. In the current study, the correlation of cognitive/memory impairment with CSF tau was dependent on CSF  $A\beta$  levels indicative for  $A\beta$  pathology, suggesting the importance of an interplay between both hallmark features of the disease. In addition, both tau and the interaction of  $A\beta_{1-42}$  with tau were significant in their effect on RBANS, but not  $A\beta_{1-42}$  alone.

Earlier publications have suggested that tau correlates with, and is predictive of, clinical function and cognitive impairment (Nelson et al., 2009). In the current study, clinical function aligned with cognitive performance (Hobson et al., 2010). The pAD group presented an RBANS Total Scale score and RBANS Index scores in line with AD pathology (Randolph et al., 1998), while the preclinical AD group presented a normal cognitive profile, confirming the usability of the RBANS as a cognitive screening tool in AD (Duff et al., 2008).

CSF tau levels (t-tau and p-tau<sub>181P</sub>) increased across stages in the early AD continuum (preclinical and pAD) with the pAD group showing highest tau levels. No further decline in CSF Aβ levels was observed across early AD stages, suggesting that CSF Aβ levels may plateau early while CSF tau levels further increase and may drive cognitive decline (Andersson et al., 2008; Buchhave et al., 2012). High CSF levels of t-tau and p-tau<sub>181P</sub> have been associated with a distinct cognitive profile with more severe impairment of memory and mental speed, and

executive functions, not explained by disease severity (van der Vlies et al., 2009). In contrast, the amount of  $A\beta$  accumulation or its removal by immunotherapy has been shown not to correlate with cognitive performance (Giannakopoulos et al., 2003; Salloway et al., 2014). In addition, substantial  $A\beta$  deposition may also occur without affecting cognitive performance overall (Perez-Nievas et al., 2013). Based on the above,  $A\beta$  pathology appears to be an early marker of incident disease.

The observation of elevated CSF tau levels already in the preclinical AD stage is in line with previous studies (<u>Hoglund et al., 2017</u>; <u>Mattsson et al., 2017</u>; <u>Toledo et al., 2013</u>; <u>Vos et al., 2013</u>), and similar as to further elevated CSF tau levels along to early AD continuum (<u>Andersson et al., 2008</u>; <u>Stefani et al., 2006</u>; <u>Toledo et al., 2013</u>). However this finding is not observed in some studies (<u>Insel et al., 2018</u>). The CSF tau elevations may indicate that the preclinical and pAD groups are already in a state of neuronal injury/degeneration.

CSF A $\beta$  levels did not significantly differ between A+ populations, despite the fact that pAD participants presumably have more accumulated brain A $\beta$  pathology compared to preclinical AD participants. CSF A $\beta$  and A $\beta$  PET measures do not change precisely in parallel as they measure different aspects of the AD amyloid pathology (Toledo et al., 2015). CSF A $\beta$  significantly decreases early in preclinical AD and then remains rather stable, whereas the A $\beta$  PET signal would continuously increase in the non-demented AD stages, lowering again in the symptomatic stage of the disease (Kadir et al., 2012). In contrast to the pAD group, the MCI group presented with a less severe and slightly distinct cognitive profile from the pAD group. The overall MCI group presented normal CSF tau levels indicating that the cognitive impairment may have another etiology than AD.

When classifying participants solely based on their CSF biomarker profile, it became apparent that CSF A $\beta$  or tau levels alone did not result in significant cognitive changes compared with the participants having normal CSF levels. (Pascoal et al., 2017), This suggests that elevated CSF tau levels are essential for learning and memory deficits in the presence of decreased CSF A $\beta$  levels indicative of amyloid pathology in AD (Desikan et al., 2012; Roberson et al., 2007). While T+ participants were observed in both A+ and A- groups, high CSF tau levels were only observed in A+ participants, suggesting that A $\beta$  pathology may further drive or advance CSF tau levels resulting in synergistic rather than additive effects (Pascoal et al., 2017). Recent in vivo labelling data suggest that neurons may respond to A $\beta$  pathology by increasing their secretion of both total and phosphorylated tau (Sato et al., 2018) Such neurons may be at increased risk of developing tangles and eventually degenerate but this may be downstream of the tau dysmetabolism that the biomarkers reflect.

In the current analysis, cognitive/memory performance correlated with CSF tau levels (t-tau and p-tau<sub>181P</sub>) in the early stages of the AD continuum (Rami et al., 2011; Reijs et al., 2017). However, the correlation observed is A $\beta$  dependent and only present in the A $\beta$  positive group,. This correlation with CSF tau levels (t-tau and p-tau<sub>181P</sub>) was also observed for all relevant RBANS subdomains, Delayed Memory, Attention and Immediate Memory on the background of A $\beta$ . A recent meta-analysis of preclinical data in transgenic mice further supports a primary role for tau in cognitive decline in preclinical AD, while A $\beta$  may rather play an indirect role in the development of NFT (Huber et al., 2018). In contrast, other groups have observed the opposite, showing a correlation of cognitive impairment with A $\beta$  on the backbone of tau in preclinical AD and MCI (Desikan et al., 2012; Haldenwanger et al., 2010).

The data presented in this study are suggestive of an interplay between  $A\beta$  and tau in relation to cognitive impairment/decline in AD. Both are required to induce the impairment while tau

pathology is driving the level of impairment. Supportive hereof is that widespread cortical tau pathology (Braak stage $\geq$ 3) has commonly been observed in patients with A $\beta$  plaques, although not in patients without plaques, supporting the hypothesis that A $\beta$  aggregation is required for the appearance of high cortical tau pathology (Knopman et al., 2003; Petersen et al., 2006; Price and Morris, 1999). In addition, significant interactions between CSF A $\beta$  and CSF p-tau levels have been shown to affect brain structure (MRI) only in preclinical AD patients positive for both CSF A $\beta$  and tau. This interaction is suggestive of a 2-phase phenomon of pathological cortical thickening associated with low A $\beta$ , followed by atrophy and cognitive decline associated with abnormal (high) CSF p-tau levels (Fortea et al., 2014).

Recent preclinical studies have highlighted several possible pathways by which Aβ could directly and indirectly influence the levels of tau, induce hyperphosphorylation of tau and the formation of neurofibrillary tangles (Gotz et al., 2001; Nisbet et al., 2015). In in vitro and in vivo (transgenic mice) models, inhibition of Aβ production by beta and gamma secretases elicits Aβ aggregation and subsequent tau pathology (incl. CSF tau) suggesting tau to be downstream effect of Aβ pathology (Choi et al., 2014; Schelle et al., 2017), hence preventing Aβ aggregation may directly affect tau pathologyOn the other hand, both Aβ and tau have shown self-propagation capabilities in animal models (Jucker and Walker, 2011). Extracellular p-tau seeds can induce tauopathy in neurons, supporting the hypothesis that tauopathy can spread in a prion-like fashion between neurons, a cascade that could be potentially initiated or accelerated by Aβ pathology (Goedert et al., 2017; Nussbaum et al., 2013). In such a case, optimal treatment to prevent cognitive decline/impairment should ideally start prior to tau reaching pathological (high) levels in CSF or before reaching levels associated with cognitive decline, hence in preclinical AD stage 1 or 2 (Sperling et al., 2011), highlighting the importance for the inclusion of biomarkers in the diagnosis of preclinical AD. Current ongoing studies in preclinical AD targeting Aβ (e.g., BACE

inhibitor studies) will provide further insight into this possible interaction between  $A\beta$  and tau and the ability to prevent cognitive decline and increases in tau pathology by  $A\beta$  inhibition.

As all participants were screened in light of an AD clinical trial, the population included may not be representative for the overall continuum as some patients may have presented with subjective memory complaints. However, data presented clearly demonstrated that based on the screening paradigm applied, different population groups could be identified from normal controls to preclinical AD and ppAD and even MCI . Secondly, this study only reports cross-sectional data, and longitudinal data further supporting the hypothesis that tau pathology correlates to cognitive decline on the backbone of  $A\beta$  (CSF) are needed. Furthermore, although similar correlations between CSF tau and cognitive impairment have been reported before, opposite results have also been reported, still describing an interaction between  $A\beta$  and tau in the opposite direction.(Desikan et al., 2012) These differences may be explained by the application of different cognitive measures. Where most studies opted for the classical clinical and cognitive measures such as MMSE, CDR and ADAS-cog, we applied next to the CDR and MMSE, for the RBANS as a sensitive and specific measure for cognitive function, currently applied in several multicentre trials in AD, which may clarify some of the differences observed.

#### 5. Conclusion

Across the early stages of the AD continuum, cognitive/memory performance correlated well with CSF tau levels. However, the observed correlation is A $\beta$ -dependent and only present in the A $\beta$ -positive group indicating the importance of a possible interplay between both neuropathological hallmarks of AD.

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Tables  $\label{eq:tables}$  Table 1A Demographics and biomarker summary by CDR and A $\beta$  (A+ or A-) status

	Control							
	(CDR=0; A-)	Preclinical AD (CDR=0; A+)	MCI (CDR≥0.5; A-)	Prodromal AD (CDR≥0.5; A+)	Total			
	N=69	N=33	N=38	N=106	N=246			
Women, n (%)	37 (53.6)	14 (42.4)	19 (50)	57 (53.8)	127 (51.6)			
Age, mean (SD), years	67.9 (5.62)	69.8 (5.54)	67.0 (6.77)	69.3 (7.08)	68.6 (6.49)			
Age group, n (%)								
< 65	18 (26.1)	7 (21.2)	12 (31.6)	26 (24.5)	63 (25.6)			
≥65 - < 75	44 (63.8)	17 (51.5)	22 (57.9)	52 (49.1)	135 (54.9)			
≥ 75	7 (10.1)	9 (27.3)	4 (10.5)	28 (26.4)	48 (19.5)			
Race, n (%)								
White	68 (98.6)	33 (100)	38 (100)	103 (97.2)	242 (98.4)			
Black or African	1 (1.4)	0	0	0	1 (0.4)			
American	, ,				, ,			
Other	0	0	0	3 (2.8)	3 (1.2)			
Weight, mean (SD), kg	74.1 (14.28)	72.9 (11.86)	75.3 (16.69)	68.7 (12.26) <sup>a</sup>	71.8 (13.88)*			
Height, mean (SD), cm	168.5 (9.87)	168.5 (9.43)	166.1 (11.24)	166.2 (10.16)	167.1 (10.08)			
BMI, mean (SD), kg/m <sup>2</sup>	26.0 (3.74)	25.6 (3.25)	27.2 (4.83)	$24.8 (3.32)^{b}$	25.6 (3.84)**			
MMSE total score	29.2 (1.40)	20.2 (1.90)	27.2 (2.76)	25 2 (2 24)	26.9 (2.05)			
Mean (SD)	28.3 (1.49)	28.3 (1.80)	27.3 (2.76)	25.3 (3.24)	26.8 (2.95)			
Median (range)	29 (24; 30)	29 (21; 30)	28 (19; 30)	26 (14; 30) <sup>c</sup>	28 (14; 30)***			
APOE ε4 carrier, n (%) Yes	0	17 (51.5)	0	58 (54.7)	75 (30.5)			
Missing/unknown	69 (100)	5 (15.2)	38 (100)	10 (9.4)	122 (49.6)			
CDR global score, n	09 (100)	3 (13.2)	36 (100)	10 (3.4)	122 (49.0)			
(%)								
0	69 (100)	33 (100)	0	0	102 (41.5)			
≥0.5	0	0	38 (100)	106 (100)	144 (58.5)			
RBANS total score	· ·	O .	30 (100)	100 (100)	111 (30.3)			
Mean (SD)	102.0 (12.70)	97.5 (17.37)	84.4 (15.19)	73.5 (16.13)	86.4 (19.69)			
Median (range)	102 (67; 136)	101 (57; 135)	87 (51; 114) <sup>d</sup>	74 (47; 124) <sup>c</sup>	87 (47; 136)***			
Biomarkers	, , ,	, , ,	, , ,	, , ,	, , ,			
Aβ 1-42 (innotest), ng/L	,							
Mean (SD)		416.5 (103.89)	840.4 (152.33)	390.8 (94.61)	595.6 (263.79)			
Madian (ranga)	960 (600, 1590)	205 (244, 500)°	815 (616; 1320)	384.5 (159;	526.5 (159;			
Median (range)	800 (000; 1380)	393 (244; 390)	813 (010; 1320)	586) <sup>e</sup>	1580)***			
<600 ng/L, n (%)	0	33 (100)	0	106 (100)	139 (56.5)			
T-tau, ng/L								
Mean (SD)	345.2 (135.65)	427.9 (192.57)	321.2 (137.53)	730.9 (333.26)	518.8 (309.06)			
Median (range)	324 (152; 742)	404 (118; 1050) <sup>f</sup>	300 (100; 661)	644.5 (161; 1780) <sup>g</sup>	448 (100; 1780)***			
>350 ng/L, n (%)	26 (37.7)	20 (60.6)	11 (28.9)	96 (90.6)	153 (62.2)			
P-tau, ng/L	_= (2,)	_= (00.0)	(-0.5)	- ( , 0.0)	(0)			
Mean (SD)	53.4 (18.73)	62.2 (21.89)	48.9 (8.43)	90.0 (33.51)	69.7 (31.94)			
Median (range)	50 (25; 116)	59 (23; 128) <sup>f</sup>	46 (20; 103)	, ,	64 (20; 197)***			
>70 ng/L, n (%)	13 (18.8)	12 (36.4)	4 (10.5)	76 (71.7)	105 (42.7)			
Aß amyloid-ß: AD Alzheimer's disease: APOF gene encoding for anolinoprotein F: BMI hody mass								

Aβ, amyloid-β; AD, Alzheimer's disease; APOE, gene encoding for apolipoprotein E; BMI, body mass index; CDR, clinical dementia rating; MCI, mild cognitive impairment; MMSE, mini-mental state

examination; P-tau, phospho tau; RBANS, repeatable battery for the assessment of neuropsychological status; SD, standard deviation; T-tau, total tau.

- \*\*\*, \*\*, \* indicates nominal p-value <0.001, <0.01, or <0.05, respectively, across groups.
- <sup>a</sup> nominal p-value <0.01 vs Control; <0.05 vs MCI.
- b nominal p-value <0.05 vs Control; <0.01 vs MCI.
- <sup>c</sup> nominal p-value <0.001 vs Control, vs Preclinical AD and vs MCI.
- <sup>d</sup> nominal p-value <0.001 vs Control, vs Preclinical AD and vs Prodromal AD.
- <sup>e</sup> nominal p-value <0.001 vs Control and vs MCI.
- f nominal p-value <0.05 vs Control; <0.01 vs MCI and <0.001 vs Prodromal AD.
- g nominal p-value < 0.001 vs Control, vs Preclinical AD and vs MCI.

Table 1B Demographics and biomarker summary by A $\beta$  (A+ or A-)/tau (T+ or T-) status

Age, mean (SD), years Age group, n (%)  < 65  ≥65 - < 75  ≥ 75  Race, n (%)  White Black or African American Other  Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m²  MMSE total score Mean (SD) Median (range)  APOE ε4 carrier, n (%)  Yes No	N=70 38 (54.3)	N=37	N=23		N=246
Women, n (%)       3         Age, mean (SD), years       66         Age group, n (%)       2         ≥65 - < 75	38 (54.3)	10 (40 4)		N=116	
Age group, n (%)  < 65  ≥65 - < 75  ≥ 75  Race, n (%)  White  Black or African  American  Other  Weight, mean (SD), kg  Height, mean (SD), cm  BMI, mean (SD), kg/m²  MMSE total score  Mean (SD)  Median (range)  APOE ε4 carrier, n (%)  Yes  No		18 (48.6)	8 (34.8)	63 (54.3)	127 (51.6)
<pre>     &lt; 65     ≥65 - &lt; 75     ≥75  Race, n (%)     White     Black or African     American     Other  Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m² MMSE total score     Mean (SD)     Median (range)     APOE ε4 carrier, n (%)     Yes     No </pre>	5.5 (5.90) <sup>a</sup>	69.6 (5.85)	68.6 (5.12)	69.6 (7.01)	68.6 (6.49)*
≥65 - < 75 ≥ 75  Race, n (%) White Black or African American Other Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m² MMSE total score Mean (SD) Median (range) APOE ε4 carrier, n (%) Yes No	23 (32.9)	7 (18.9)	7 (30.4)	26 (22.4)	63 (25.6)
≥ 75 Race, n (%) White Black or African American Other Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m² MMSE total score Mean (SD) Median (range) APOE ε4 carrier, n (%) Yes No	42 (60)	24 (64.9)	13 (56.5)	56 (48.3)	135 (54.9)
White Black or African American Other Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m² MMSE total score Mean (SD) Median (range) APOE &4 carrier, n (%) Yes No	5 (7.1)	6 (16.2)	3 (13.0)	34 (29.3)	48 (19.5)
Black or African American Other Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m² MMSE total score Mean (SD) Median (range) APOE \$\varepsilon 4\$ carrier, n (%) Yes No	,	, ,	, ,	, ,	, ,
American Other  Weight, mean (SD), kg 75 Height, mean (SD), cm 166 BMI, mean (SD), kg/m² 26 MMSE total score Mean (SD) 28 Median (range) 28 APOE & carrier, n (%) Yes No	69 (98.6)	37 (100)	22 (95.7)	114 (98.3)	242 (98.4)
Other Weight, mean (SD), kg 75 Height, mean (SD), cm 166 BMI, mean (SD), kg/m² 26 MMSE total score Mean (SD) 28 Median (range) 28 APOE \$\varepsilon 4\$ carrier, n (%) Yes No		, ,	, ,	, ,	, ,
Weight, mean (SD), kg Height, mean (SD), cm BMI, mean (SD), kg/m² MMSE total score Mean (SD) Median (range)  APOE ε4 carrier, n (%) Yes No	1 (1.4)	0	0	0	1 (0.4)
Height, mean (SD), cm BMI, mean (SD), kg/m <sup>2</sup> MMSE total score Mean (SD) Median (range) APOE & carrier, n (%) Yes No	0	0	1 (4.3)	2 (1.7)	3 (1.2)
Height, mean (SD), cm BMI, mean (SD), kg/m <sup>2</sup> 26 MMSE total score Mean (SD) Median (range) 27 APOE ε4 carrier, n (%) Yes No	.0 (16.45)	73.5 (12.37)	74.7 (10.34)	68.7 (12.40) <sup>b</sup>	71.8 (13.88)**
MMSE total score  Mean (SD) 28  Median (range) 28  APOE ε4 carrier, n (%)  Yes  No	5.7 (10.56)	169.4 (9.99)	171.2 (9.91)	165.8 (9.82)	167.1 (10.08)
MMSE total score  Mean (SD) 28  Median (range) 28  APOE ε4 carrier, n (%)  Yes  No	5.8 (4.28)	25.7 (3.92)	25.4 (2.28)	$24.9 (3.48)^{\circ}$	25.6 (3.84)*
Median (range) 28 APOE ε4 carrier, n (%) Yes No					
APOE ε4 carrier, n (%) Yes No	8.0 (1.93)	27.8 (2.37)	28.0 (2.18)	25.6 (3.27)	26.8 (2.95)
Yes No	3 (19; 30)	28 (21; 30)	28 (21; 30)	26 (14; 30) <sup>d</sup>	28 (14; 30)***
No					
	0	0	11 (47.8)	64 (55.2)	75 (30.5)
Missing/unknown	0	0	10 (43.5)	39 (33.6)	49 (19.9)
Wilsonig/ unknown	70 (100)	37 (100)	2 (8.7)	13 (11.2)	122 (49.6)
CDR global score, n (%)					
	13 (61.4)	26 (70.3)	13 (56.5)	20 (17.2)	102 (41.5)
≥0.5	27 (38.6)	11 (29.7)	10 (43.5)	96 (82.8) <sup>d</sup>	144 (58.5)***
RBANS total score					
Mean (SD) 94	.8 (14.61)	97.5 (18.41)	93.6 (18.78)	76.4 (18.17)	86.4 (19.69)
Median (range) 94.	5 (55; 125)	99 (51; 136)	100 (59; 124)	76 (47; 135) <sup>d</sup>	87 (47; 136)***
Biomarkers					
Aβ 1-42 (innotest), ng/L					
Mean (SD) 806	5.2 (137.15)	943.3 (195.94)	432.3 (110.96)	389.9 (93.07)	595.6 (263.79)
Median (range) 7	(90 (600; 1180) <sup>e</sup>	903 (607; 1580) <sup>f</sup>	421 (259; 590)	381 (159; 587)	526.5 (159; 1580)***
<600 ng/L, n (%)	0	0	23 (100)	116 (100)	139 (56.5)
T-tau, ng/L					
Mean (SD) 25	7.1 (62.11)	487.3 (107.44)	257.5 (69.10)	738.6 (303.76)	518.8 (309.06)
Median (range) 264	(100; 350)	458 (353; 742) <sup>f</sup>	262 (118; 349)	637 (354; 1780) <sup>g</sup>	448 (100; 1780)***
>350 ng/L, n (%)	0	37 (100)	0	116 (100)	153 (62.2)
P-tau, ng/L		, ,		, ,	, ,
_	1.5 (9.68)	71.5 (15.53)	41.8 (9.29)	91.7 (29.93)	69.7 (31.94)
Madian (ranga)	.5 (20; 62)	69 (46; 116) <sup>f</sup>	41 (23; 56)	85 (49; 197) <sup>g</sup>	64 (20;
>70 ng/L, n (%)		` ' '		*	197)***

Aβ, amyloid-β; *APOE*, gene encoding for E; BMI, body mass index; CDR, clinical dementia rating; MMSE, mini-mental state examination; P-tau, phospho tau; RBANS, repeatable battery for the assessment of neuropsychological status; SD, standard deviation; T-tau, total tau.

\*\*\*, \*\*, \* indicates nominal p-value <0.001, <0.01, or <0.05, respectively, across groups.

<sup>&</sup>lt;sup>a</sup> nominal p-value <0.05 vs A-/T+; <0.01 vs A+/T+.

b nominal p-value <0.05 vs A-/T+ and vs A+/T-; <0.01 vs A-/T-.

<sup>&</sup>lt;sup>c</sup> nominal p-value <0.01 vs A-/T-.

d nominal p-value <0.001 vs A-/T-, vs A-/T+ and vs A+/T-.

e nominal p-value <0.001 vs A-/T+, vs A+/T- and vs A+/T+.

f nominal p-value <0.001 vs A-/T-, vs A+/T- and vs A+/T+.

g nominal p-value <0.001 vs A-/T-, A-/T+ and vs A+/T-.

Table 2. Correlation of RBANS indexes vs T-tau and P-tau

	All N=246		Αβ- N=107		Αβ+ N=139	
	T-tau	P-tau	T-tau	P-tau	T-tau	P-tau
RBANS total score	-0.434***	-0.389***	0.137	0.173	-0.451***	-0.421***
RBANS sum of	-0.433***	-0.391***	0.139	0.176	-0.449***	-0.422***
index						
RBANS delayed	-0.433***	-0.377***	0.236*	0.310**	-0.431***	-0.406***
memory						
RBANS immediate	-0.368***	-0.329***	0.086	0.109	-0.327***	-0.307***
memory						
RBANS attention	-0.339***	-0.315***	0.119	0.120	-0.438***	-0.410***
Index						
RBANS language	-0.321***	-0.311***	-0.052	-0.049	-0.261**	-0.275**
index						
RBANS visuospatial	-0.186**	-0.165**	0.063	0.069	-0.254**	-0.223**
index						

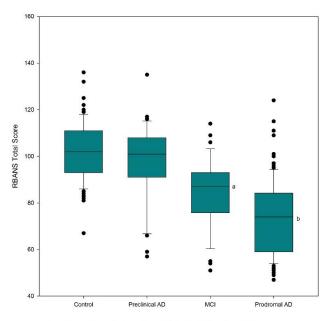
RBANS, repeatable battery for the assessment of neuropsychological status.

<sup>\*\*\*, \*\*, \*</sup> indicates nominal p-value <0.001, <0.01, or <0.05, respectively.

# **Figures**

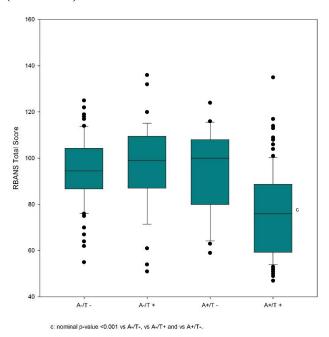
# Figure 1. RBANS Total Scale Score

# A: CDR/Aβ Status



a: nominal p-value <0.001 vs Control, vs Preclinical AD and vs Prodromal AD. b: nominal p-value <0.001 vs Control, vs Preclinical AD and vs MCI.

# B. Aβ (A+ or A-)/Tau (T+ or T-) Status

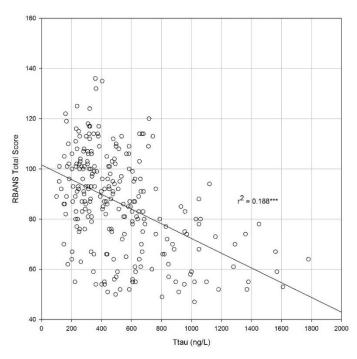


RBANS: repeatable battery for the assessment of neuropsychological status

Participants are considered A $\beta$ + (A+) if CSF A $\beta_{1-42}$  < 600 ng/L and Tau+ (T+) if CSF T-Tau >350 ng/L or CSF p-Tau<sub>181P</sub> > 70 ng/L. (A) Control (CDR 0; A $\beta$ -; n=69), pre-clinical AD (CDR 0; A $\beta$ +; n=33), MCI (CDR  $\ge$ 0.5; A $\beta$ -; n=38), prodromal AD (CDR  $\ge$ 0.5; A $\beta$ +; n=106); (B) A-/T- (n=70), A-/T+ (n=37), A+/T- (n=23), A+/T+ (n=116).

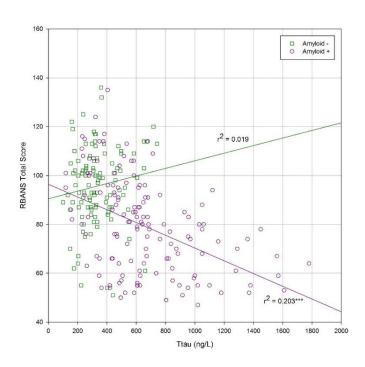
Figure 2. Correlation RBANS Total Scale Score and T-tau

# **A:** Overall Population



<sup>\*\*\*</sup> indicates nominal p-value <0.001

# B. Aß Status



<sup>\*\*\*</sup> indicates nominal p-value <0.001

RBANS: repeatable battery for the assessment of neuropsychological status; Ttau: total tau A Spearman correlation coefficient was calculated to evaluate the possible correlation between RBANS Total Scale Score and t-tau for the overall population (A) and by A $\beta$  (amyloid) status (B). (A) R<sup>2</sup>=0.188, rho = -0.434; p<0.001. n = 246; B. Amyloid -: R<sup>2</sup>=0.019, rho = 0.137, p>0.05, n=107; Amyloid +: R<sup>2</sup>=0.203, rho = -0.451, p<0.001, n=139).