

Head-to-head comparison of 10 plasma phospho-tau assays in prodromal Alzheimer's disease

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8 **Running title:** Comparison of plasma p-tau assays

9

10 **Keywords:** blood p-tau; Alzheimer's disease; amyloid- β ; dementia

11

12 **Abbreviations:** A β = amyloid- β ; AUC = area under the curve; CSF = cerebrospinal fluid; MCI
13 = mild cognitive impairment; MMSE = mini-mental state examination; MS = mass spectrometry;
14 ROC = receiver operating characteristic; PET = positron emission tomography; p-tau =
15 phosphorylated tau

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1 Abstract

2 Plasma phospho-tau (p-tau) species have emerged as the most promising blood-based biomarkers
3 of Alzheimer's disease. Here, we performed a head-to-head comparison of p-tau181, p-tau217
4 and p-tau231 measured using 10 assays to detect abnormal brain amyloid- β status and predict
5 future progression to Alzheimer's dementia. The study included 135 patients with baseline
6 diagnosis of mild cognitive impairment (mean age 72.4 years; 60.7% women) who were
7 followed for an average of 4.9 years. Seventy-one participants had abnormal A β -status (i.e.,
8 abnormal CSF A β 42/40) at baseline; and 45 of these A β -positive participants progressed to
9 Alzheimer's dementia during follow-up. P-tau concentrations were determined in baseline
10 plasma and CSF. P-tau217 and p-tau181 were both measured using immunoassays developed by
11 Lilly Research Laboratories (Lilly) and mass spectrometry assays developed at Washington
12 University (WashU). P-tau217 was also analysed using Simoa immunoassay developed by
13 Janssen Research and Development (Janss). P-tau181 was measured using Simoa immunoassay
14 from ADxNeurosciences (ADx), Lumipulse immunoassay from Fujirebio (Fuji) and Splex
15 immunoassay from Mesoscale Discovery (Splex). Both p-tau181 and p-tau231 were quantified
16 using Simoa immunoassay developed at the University of Gothenburg (UGOT). We found that
17 the mass spectrometry-based p-tau217 (p-tau217^{WashU}) exhibited significantly better performance
18 than all other plasma p-tau biomarkers when detecting abnormal A β status (AUC=0.947;
19 $p_{\text{diff}} < 0.015$) or progression to Alzheimer's dementia (AUC=0.932; $p_{\text{diff}} < 0.027$). Among
20 immunoassays, p-tau217^{Lilly} had the highest AUCs (0.886-0.889), which was not significantly
21 different from the AUCs of p-tau217^{Janss}, p-tau181^{ADx} and p-tau181^{WashU} (AUC_{range}, 0.835-0.872;
22 $p_{\text{diff}} > 0.09$), but higher compared with AUC of p-tau231^{UGOT}, p-tau181^{Lilly}, p-tau181^{UGOT}, p-
23 tau181^{Fuji}, and p-tau181^{Splex} (AUC_{range}, 0.642-0.813; $p_{\text{diff}} \leq 0.029$). Correlations between plasma

1 and CSF values were strongest for p-tau217^{WashU} (R=0.891) followed by p-tau217^{Lilly} (R=0.755;
2 $p_{\text{diff}}=0.003$ vs p-tau217^{WashU}) and weak to moderate for the rest of the p-tau biomarkers (R_{range},
3 0.320-0.669). In conclusion, the findings suggest that among all tested plasma p-tau assays, mass
4 spectrometry-based measures of p-tau217 perform best when identifying mild cognitive
5 impairment patients with abnormal brain A β or those who will subsequently progress to
6 Alzheimer's dementia. Several other assays (p-tau217^{Lilly}, p-tau217^{Janss}, p-tau181^{ADx}, and p-
7 tau181^{WashU}) showed relatively high and consistent accuracy across both outcomes. The results
8 further indicate that the highest performing assays have performance metrics that rival the gold
9 standards of A β -PET and CSF. If further validated, our findings will have significant impacts in
10 diagnosis, screening and treatment for Alzheimer's dementia in the future.

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1 Introduction

2 Alzheimer's disease neuropathologic changes in the brain, i.e., accumulation amyloid- β ($A\beta$)
3 plaques and neurofibrillary tangles containing hyperphosphorylated tau (p-tau), can be detected
4 in living people using positron emission tomography (PET) scanning or quantification of $A\beta$
5 and p-tau proteins levels in CSF.¹ Although $A\beta$ - and tau-PET as well as CSF $A\beta_{42/40}$ and p-tau
6 are highly accurate and validated diagnostic and prognostic biomarkers of Alzheimer's disease²⁻⁴
7 that have been widely used in research settings, blood-based tests are needed for implementation
8 in clinical practice globally and to facilitate patient screening and selection in clinical trials.^{3,5}
9
10 In CSF, soluble p-tau species change in different stages and progression of Alzheimer's disease.⁶
11 A growing number of studies have demonstrated that three variants of p-tau, p-tau181, p-tau217
12 and p-tau231, measured in blood plasma hold great promise as biomarkers of Alzheimer's
13 disease related $A\beta$ and tau pathologies.⁷⁻¹¹ At the same time, there are reported differences in the
14 performance of different plasma p-tau species and assays. For example p-tau217 (measured
15 using either mass spectrometry [MS] or immunoassays) has consistently shown higher accuracy
16 for detecting abnormal CSF and PET biomarker status and differentiating Alzheimer's disease
17 from other neurodegenerative disorders (in both clinical and neuropathological cohorts) and
18 controls than p-tau181, even though the effect sizes were in many cases relatively small.^{7,10,12,13}
19 Some data also suggest that while plasma p-tau231 and p-tau181 perform equally well as
20 diagnostic biomarkers in later dementia phase of Alzheimer's disease, p-tau231 starts to increase
21 earlier than p-tau181 and is more strongly associated with $A\beta$ and tau PET measures in
22 preclinical disease stages.¹⁴⁻¹⁶ However, it is at present unclear how much varying performance
23 of the plasma p-tau biomarkers is attributable to analytical measurement methods. Several
24 immunoassays¹⁷ and an MS-based method⁷ have been developed for determination of different
25 p-tau species in plasma and used across different studies making their interpretation challenging.
26 MS is considered as "the gold standard" for protein identification and analysis and although
27 published work shows that MS-based plasma $A\beta$ measures might more accurately reflect brain
28 $A\beta$ pathology in Alzheimer's disease than immunoassays,¹⁸ a direct comparison of these
29 methods for blood p-tau quantification is currently lacking. Some studies, on the other hand,
30 compared several of the available plasma p-tau immunoassays. P-tau217 measured with two

1 different immunoassays developed by Lilly Research Laboratories and Janssen Research and
2 Development have both been shown to accurately predict abnormal CSF A β status and future
3 conversion to Alzheimer's disease dementia in patients with mild cognitive impairment (MCI).¹⁹
4 In contrast, a certain degree of variability has been found in performance of different p-tau181
5 immunoassays^{12,20}. Interestingly, differences in the performance between plasma p-tau217 and p-
6 tau181 appears much smaller when both biomarkers are measured with Lilly immunoassays that
7 only differ in phospho-specific capture antibodies compared to the differences between Lilly p-
8 tau217 and other p-tau181 immunoassays.^{10,12,13} Collectively, these findings suggest that
9 immunoassay components (e.g., antibodies, other reagents, detection systems) may affect the
10 performance of p-tau biomarkers and illustrate the importance of conducting head-to-head
11 comparisons of different plasma p-tau immunoassays. On the other hand, mass spectrometry
12 measurement of tau peptides generated by trypsinization or other enzymatic digestions may be
13 confounded by the presence of various endogenously produced tau truncated species.²¹
14 Expanding on previous preliminary studies, with the additional aim to compare MS-based
15 methods and immunoassays, we analyzed p-tau181, p-tau217 and p-tau231 using 10 assays in
16 plasma samples from a cohort of MCI patients who were followed for up to 9.5 years to monitor
17 progression of clinical symptoms. We tested the ability of p-tau biomarkers to identify
18 participants with abnormal CSF A β status and to predict future progression from MCI to
19 Alzheimer's disease dementia.

20

21 **Materials and methods**

22 **Participants**

23 The study was approved by the Ethics Committee at the University of Lund and the patients
24 and/or their relatives gave their informed consent (for research). We included 135 individuals
25 with clinical diagnosis of MCI at baseline who were recruited at the Memory Clinic at Skåne
26 University Hospital in Malmö, Sweden.^{19,22,23} All participants underwent a thorough physical,
27 neurological, and psychiatric examination, as well as a clinical interview focusing on cognitive
28 symptoms and activities of daily living function by physicians with an expertise in cognitive

1 disorders. Patients with MCI at baseline had to fulfill the criteria by Petersen,²⁴ including (1)
2 memory complaint, preferably corroborated by an informant; (2) objective memory impairment
3 adjusted for age and education, as judged by the physician; (3) preservation of general cognitive
4 functioning, as determined by the clinician's judgment based on a structured interview with the
5 patient and a Mini Mental Status Examination (MMSE) score greater than or equal to 24; (4)
6 zero or minimal impairment of daily life activities; and (5) not fulfilling the DSM-III-R criteria
7 for dementia. The exclusion criteria were (1) significant unstable systemic illness or organ
8 failure; (2) current significant alcohol or substance misuse; and (3) cognitive impairment that
9 could be explained by other specific non-neurodegenerative disorders such as brain tumor or
10 subdural hematoma. Study participants were followed for an average of 4.9 (SD=2.1) years. The
11 MCI-ADD group included participants who progressed to Alzheimer's disease dementia during
12 follow-up. Patients who received a diagnosis of Alzheimer's disease were required to meet the
13 DSM-III-R criteria for dementia and the criteria of probable Alzheimer's disease defined by
14 NINCDS-ADRDA²⁵ and have abnormal CSF A β 42/40 ratio.¹⁹ The criteria for non-AD dementia
15 diagnosis in this MCI cohort have been previously described.^{22,23} Stable MCI (sMCI) patients
16 and MCI who progressed to non-Alzheimer's disease dementia were classified as non-
17 progressors and further stratified into A β -negative (A-) and A β -positive (A+) groups based on
18 the CSF A β 42/40 ratio status. The characteristics of the study participants are given in Table 1.

19 **CSF and plasma sampling and analysis**

20 CSF and blood sample were drawn in the morning while participants were not necessarily non-
21 fasting. Blood was collected in six K2-EDTA-plasma tubes and centrifuged at 2000g, +4°C for
22 10 minutes. Following centrifugation plasma was aliquoted into 1.5-ml polypropylene tubes (1
23 ml per tube) and stored at -80°C. CSF was obtained by lumbar puncture and stored at -80°C in
24 polypropylene tubes following the Alzheimer's Association flow chart for lumbar puncture and
25 CSF sample processing.²⁶ All samples went through one freeze-thaw cycle before the analysis
26 when 0.2-0.5ml were further aliquoted into LoBind tubes. P-tau217 was measured as
27 phosphorylation occupancy at Thr217 using MS assay developed at Washington University (p-
28 tau217^{WashU}),⁷ Meso Scale Discovery (MSD) immunoassay developed by Lilly Research
29 Laboratories (p-tau217^{Lilly})^{10,27} and Single molecule arrays (Simoa) immunoassay developed by
30 Janssen Research and Development (p-tau217^{Janss}).^{19,28,29} P-tau181 was measured as

1 phosphorylation occupancy at Thr181 using MS-WashU assays (p-tau181^{WashU}),⁷ MSD
2 immunoassay developed by Lilly Research Laboratories (p-tau181^{Lilly}),^{8,30} Simoa immunoassay
3 developed at the University of Gothenburg (p-tau181^{UGOT}),⁹ Simoa immunoassay developed by
4 ADx Neurosciences (p-tau181^{ADx}),^{20,31} Lumipulse immunoassay developed by Fujirebio (p-
5 tau181^{Fuji}) and Splex immunoassay from MSD (p-tau181^{Splex}). P-tau231 was measured using in-
6 house Simoa immunoassay developed at the University of Gothenburg (p-tau231^{UGOT}).¹⁴ We
7 also tested p-tau231^{Splex} assay from MSD. However, this assay failed to detect any measurable p-
8 tau231 in a pilot study of eight plasma samples (four from A β -negative and the other four from
9 A β -positive individuals) analyzed across 2 runs and therefore was not included in the present
10 study. P-tau217^{Lilly} and p-tau217^{Janss} data in overlapping sample have been reported previously.¹⁹
11 CSF samples (N=78) were analyzed using p-tau217^{WashU}, p-tau217^{Lilly}, p-tau217^{Janss}, p-
12 tau181^{WashU}, p-tau181^{ADx}, p-tau181^{UGOT}, p-tau181^{Fuji} and p-tau231^{UGOT} assays. CSF A β 40 and
13 A β 42 levels were assessed using commercially available MSD immunoassays. Amyloid
14 positivity was defined based on CSF A β 42/40 and a previously described threshold of 0.07.^{22,23}
15 All samples were analyzed by staff blinded to the clinical data. Further details of the p-tau
16 analyses are described in the Supplementary Methods and data on assay performance are shown
17 in Table 2 and Supplementary Figure 1.

18 **Statistical analysis**

19 SPSS (version 28, IBM, Armonk, NY, US) and R (version 4.1.2) in RStudio³² were used for
20 statistical analysis. Demographic and clinical data were compared with Mann-Whitney U,
21 Kruskal-Wallis and chi-square (sex and *APOE* ϵ 4 positivity) tests. Group differences in the
22 log₁₀-transformed biomarker levels were assessed with univariate general linear models
23 adjusting for age and sex and additionally for duration of follow-up when comparing MCI
24 participants who progressed to Alzheimer's disease dementia with those who did not. In figures,
25 fold changes relative to the mean of the A- sMCI group are presented to aid interpretation of
26 biomarker levels across comparisons. Correlations between CSF and plasma were examined
27 using Spearman test and we used bootstrapping (n=2000 iterations) to test differences in the
28 correlation coefficients. Diagnostic accuracies of CSF biomarkers were assessed using receiver
29 operating characteristic (ROC) curve analysis. The Youden index with bootstrapping (n=2000
30 iterations) was used to determine sensitivity, specificity and accuracy with 95% confidence

1 interval (CI) at optimal thresholds. Area under the curve (AUC) of two ROC curves were
2 compared with DeLong test with adjustment for multiple comparisons using Benjamini-
3 Hochberg false discovery rate method.³³ For p-tau181^{UGOT} and p-tau181^{Splex} assays, plasma
4 samples from 124 and 101 participants, respectively, were analyzed and included in the main
5 analysis. However, we performed a sensitivity analysis in subsamples where all plasma p-tau
6 measures were available. Two-sided $p < 0.05$ was considered statistically significant.

7 **Data availability**

8 Anonymized data will be shared by request from a qualified academic investigator for the sole
9 purpose of replicating procedures and results presented in the article and as long as data transfer
10 is in agreement with EU legislation on the general data protection regulation and decisions by the
11 Ethical Review Board of Sweden and Region Skåne, which should be regulated in a material
12 transfer agreement.

13 **Results**

14 **Participants**

15 The study included 45 MCI patients who progressed to AD dementia (MCI-ADD), 64 non-
16 progressors with normal A β -status (A-) and 26 A+ non-progressors (Table 1). There were
17 differences in age ($H(2)=19.0$, $p < 0.001$), sex ($\chi^2(2)=8.1$, $p=0.018$), MMSE ($H(2)=30.1$, $p < 0.001$),
18 *APOE* $\epsilon 4$ carriership ($\chi^2(2)=33.0$, $p < 0.001$) and follow-up duration ($H(2)=23.3$, $p < 0.001$)
19 between the groups. The MCI-ADD group was on average older, had lower MMSE and shorter
20 follow-up time than both non-progressor groups ($p < 0.001$). There were more women among
21 MCI-ADD compared with A+ non-progressors ($p=0.005$) and A- non-progressors ($p=0.056$),
22 whereas *APOE* $\epsilon 4$ positivity rate was lower in A- non-progressors than both A+ non-progressors
23 and MCI-ADD ($p < 0.001$)

1 **Associations with A β pathology**

2 We first assessed how well plasma p-tau species measured with different assays identified
 3 individuals with abnormal baseline A β status among all study participants with baseline
 4 diagnosis of MCI (Figure 1A, Table 3). In ROC curve analysis, the mass spectrometry-based p-
 5 tau217 assay (p-tau217^{WashU}) performed significantly better than all other p-tau biomarkers with
 6 AUC of 0.947 (95% CI, 0.907-0.987; $p_{diff}<0.015$). Among immunoassays, p-tau217^{Lilly} had the
 7 highest AUC (AUC=0.886; CI, 0.827-0.944), which was not significantly different from the
 8 AUCs of p-tau217^{Janss} (AUC=0.858; 95% CI, 0.795-0.920; $p_{diff}=0.38$), p-tau181^{ADx} (AUC=0.841;
 9 95% CI, 0.768-0.913; $p_{diff}=0.24$) and p-tau181^{WashU} (AUC=0.835; 95% CI, 0.765-0.906;
 10 $p_{diff}=0.20$), but higher compared with AUC of p-tau231^{UGOT}, p-tau181^{Lilly}, p-tau181^{UGOT}, p-
 11 tau181^{Fuji}, and p-tau181^{Splex} (AUC_{range}, 0.642-0.784; $p_{diff}\leq 0.029$). For comparison, the AUCs of
 12 the best performing CSF p-tau assays in a subsample of 78 participants with CSF measures
 13 available ranged between 0.948 and 0.975 (p-tau217^{WashU}, AUC=0.975; p-tau181^{ADx},
 14 AUC=0.961; p-tau181^{WashU}, AUC=0.954; p-tau217^{Lilly}, AUC=0.952; p-tau217^{Janss}, AUC=0.948).
 15 CSF p-tau showed significantly higher AUCs than corresponding plasma p-tau for most assays
 16 (Supplementary Table 1).

17
 18 When testing differences in plasma p-tau levels between A+ and A- groups, we found that all 10
 19 p-tau biomarkers were significantly higher in A+ MCI than A- MCI (Figure 2). However, the
 20 fold increase in the A+ group compared with the A- group was largest for the p-tau217^{WashU}
 21 (mean=3.6, SD=1.9), followed by p-tau217^{Janss} (mean=2.7, SD=1.8), P-tau217^{Lilly} (mean=2.0,
 22 SD=1.0), and p-tau181^{ADx} (mean=1.8, SD=0.8) and ranging between 1.2 and 1.4 for the rest of
 23 the biomarkers.

24 **Prediction of future progression to Alzheimer's disease dementia**

25 We next studied the performance of the plasma p-tau biomarkers to predict future clinical
 26 progression to Alzheimer's disease dementia (Figure 1B, Table 4). When distinguishing MCI
 27 patients who progressed to Alzheimer's disease dementia during follow-up from those who did
 28 not, p-tau217^{WashU} again showed significantly higher AUC than all other p-tau biomarkers
 29 (AUC=0.932; 95% CI, 0.891-0.974; $p_{diff}<0.027$) followed by p-tau217^{Lilly} (AUC=0.889; 95% CI,

1 0.833-0.946). P-tau217^{Janss} (AUC=0.872; 95% CI, 0.814-0.931; $p_{diff}=0.53$), p-tau181^{ADx}
 2 (AUC=0.846; 95% CI, 0.777-0.916; $p_{diff}=0.16$) and p-tau181^{WashU} (AUC=0.835; 95% CI, 0.764-
 3 0.906; $p_{diff}=0.09$) were non-inferior to p-tau217^{Lilly}, whereas p-tau231^{UGOT}, p-tau181^{Lilly}, p-
 4 tau181^{UGOT}, p-tau181^{Fuji}, and p-tau181^{Splex} all had significantly lower AUCs (AUC_{range}, 0.688-
 5 0.813; $p_{diff} \leq 0.013$). For comparison, the AUCs of the best performing CSF p-tau assays in a
 6 subsample of 78 participants with CSF measures available ranged between 0.907 and 0.943 (p-
 7 tau217^{WashU}, AUC=0.943; p-tau217^{Janss}, AUC=0.928; p-tau217^{Lilly}, AUC=0.926; p-tau181^{ADx},
 8 AUC=0.924; p-tau181^{Fuji}, AUC=0.907). The differences in AUCs between CSF and
 9 corresponding plasma p-tau assays were not significant (Supplementary Table 1).

10
 11 We also found differences in plasma concentrations of all p-tau biomarkers except p-tau181^{Fuji}
 12 between the A- non-progressor, A+ non-progressor and MCI-ADD groups (Figure 3). Post-hoc
 13 analysis revealed that plasma levels of p-tau217 (when measured with three different assays), but
 14 not p-tau181 or ptau231, were higher in MCI-ADD than A+ non-progressors ($p < 0.002$). At the
 15 same time, the three p-tau217 biomarkers as well as the best performing p-tau181 biomarkers (p-
 16 tau181^{WashU} and p-tau181^{ADx}) were increased in both A+ non-progressors and MCI-ADD
 17 compared with A- non-progressors ($p \leq 0.001$). P-tau217^{WashU} showed the largest fold increase in
 18 both MCI-ADD (mean=4.3, SD=1.7) and A+ non-progressors (mean=2.5, SD=1.4) compared
 19 with A- non-progressors. Fold increase was also larger in MCI-ADD (mean_{range}, 2.0-3.2) than in
 20 A+ non-progressors (mean_{range}, 1.4-1.9) for p-tau217^{Lilly}, p-tau217^{Janss} and p-tau181^{ADx}.

21 **Correlations between plasma and CSF p-tau**

22 Finally, we examined associations between plasma and CSF p-tau biomarkers (Figure 4). CSF p-
 23 tau concentrations are presented in Supplementary Table 2. In line with other results of this
 24 study, the strongest correlations between CSF and plasma were seen for p-tau217^{WashU} ($R=0.891$;
 25 95% CI, 0.832-0.930), followed by p-tau217^{Lilly} ($R=0.755$; 95% CI, 0.635-0.839) with significant
 26 difference in correlation coefficients between the two biomarkers ($p=0.003$). The correlations
 27 were weak to moderate for the rest of the biomarkers (R_{range} , 0.320-0.669).

28

1 Plasma p-tau217^{WashU} correlated strongly with plasma p-tau217^{Lilly}, p-tau217^{Janss}, p-tau181^{ADx}
 2 and p-tau181^{WashU} (R_{range} , 0.712-0.862; Supplementary Figure 2) while correlations with other
 3 plasma p-tau biomarkers were weak to moderate (R_{range} , 0.376-0.619; Supplementary Figure 2).

4 Sensitivity analysis

5 The results were similar when statistical analysis was performed in smaller sub-samples where p-
 6 tau181^{UGOT} and p-tau181^{Splex} data were available (Supplementary Tables 3-6). Briefly, plasma p-
 7 tau217^{WashU} showed the best performance when detecting both abnormal A β status and
 8 progression to Alzheimer's disease dementia (AUC_{range} , 0.927-0.955), followed by p-tau217^{Lilly}
 9 (AUC_{range} , 0.878-0.900), p-tau217^{Janss} (AUC_{range} , 0.860-0.870), p-tau181^{ADx} (AUC_{range} , 0.832-
 10 0.860) and p-tau181^{WashU} (AUC_{range} , 0.809-0.827). None of the AUCs of P-tau231^{UGOT}, p-
 11 tau181^{Lilly}, p-tau181^{UGOT}, p-tau181^{Fuji}, p-tau181^{Splex} were consistently above 0.800.

12 Discussion

13 Recently developed blood tests for A β and p-tau are anticipated to transform Alzheimer's disease
 14 research and care. Here we sought to directly compare currently available methods for
 15 determinations of p-tau in blood in order to establish which of these methods are accurate
 16 enough to be useful for implementation in clinical practice or drug trials. In this study including
 17 patients with MCI, plasma p-tau217 quantified using MS-based assay showed very high
 18 accuracy when both identifying participants with abnormal A β status and those who progress to
 19 Alzheimer's disease dementia during follow-up with $AUCs > 0.93$ which was higher than for the
 20 other p-tau biomarkers. Further, this assay exhibited significantly higher correlations with p-tau
 21 levels in CSF than the other p-tau assays. However, p-tau217^{Lilly}, p-tau217^{Janss}, p-tau181^{ADx} and
 22 p-tau181^{WashU} all displayed relatively high and consistent accuracy across both outcomes
 23 (AUC_{range} , 0.835-0.889), whereas the performance of other biomarkers (p-tau231^{UGOT}, p-
 24 tau181^{Lilly}, p-tau181^{UGOT}, p-tau181^{Fuji}, p-tau181^{Splex}) was significantly inferior (AUC_{range} , 0.642-
 25 0.813). Of note, there was no added value of combining different plasma p-tau species (p-
 26 tau217^{WashU}, ptau181^{ADx} and p-tau231^{UGOT}) when either distinguishing normal from abnormal
 27 A β status or predicting future progression to Alzheimer's disease dementia (data not shown).

28

1 MS-based measure of plasma p-tau217 has previously shown very good accuracy to detect A β
2 pathology in 2 mixed cohorts of cognitively healthy controls, MCI participants and patients at
3 different stages of Alzheimer's disease.⁷ Using an improved version of the same MS assay (now
4 requiring lower volume of plasma) we demonstrate that p-tau217^{WashU} accurately predicted
5 abnormal A β status as well as future progression to Alzheimer's disease dementia in a sample of
6 MCI patients. One novel finding of the present study is that MS p-tau217^{WashU} performed
7 significantly better than p-tau217 quantified with immunoassays. A possible explanation for this
8 may be that MS-based detection methods are highly accurate and potentially more so than
9 immunoassays, and therefore could more reliably quantify low abundance proteins in protein-
10 rich matrices such as blood as was seen for plasma A β .¹⁸

11
12 We also found that p-tau217^{WashU} performed better than p-tau181^{WashU} corroborating the results
13 of an earlier MS-based study.⁷ The higher performance of p-tau217 over p-tau181 has been
14 shown for immunoassays-based p-tau measures^{10,12,13} as well as for CSF p-tau217 and p-
15 tau181^{34,35} and could be due to the specificity of p-tau217 for Alzheimer's disease (this
16 biomarker is found at considerably lower levels in people without Alzheimer's disease compared
17 to p-tau181) and to a greater dynamic range of p-tau217, i.e. larger fold increase in relation to
18 developing A β and tau pathologies. Among eight immunoassays tested in the present study, p-
19 tau217^{Lilly} displayed numerically highest AUCs which were significantly different from the
20 AUCs of several p-tau181 biomarkers. However, p-tau217^{Lilly}, p-tau217^{Janss} and p-tau181^{ADx} all
21 exhibited comparable accuracies for both abnormal A β status and progression to Alzheimer's
22 disease dementia indicating substantial variability in the performance of p-tau181 that is most
23 likely caused by the differences in antibodies and analytical procedures used across the assays.

24
25 Our study has several limitations. The overall sample size was moderate with a relatively small
26 number of A+ non-progressor and participant with CSF data which might have affected the
27 analysis. The cohort was restricted to MCI participants and it is possible that the performance of
28 the plasma p-tau assays varies across disease stages warranting future investigations in
29 individuals with preclinical Alzheimer's disease. Nevertheless, our findings in MCI patients are
30 very relevant given that this patient group represent the most likely target population to receive
31 disease-modifying therapies in the clinical settings in the coming years. Replication in more

1 heterogeneous and ethnically diverse population-based cohorts is also needed. Finally, future
2 larger studies should establish if combining individual plasma p-tau biomarkers with other
3 accessible demographic and clinical measures could further improve their diagnostic and
4 prognostic accuracy as has previously been shown for plasma p-tau217.³⁶

5
6
7 In conclusion, we show that there are significant and meaningful differences in the performance
8 of plasma p-tau assays that have to be taken into account when interpreting results from
9 published work. Our data support superior performance of MS p-tau217 to detect abnormal A β
10 status and progression to Alzheimer's disease dementia in MCI patients. In addition, we report
11 relatively high and consistent accuracy for several p-tau immunoassays for both outcomes.
12 Overall, our findings indicate that certain MS-based methods and immunoassays might be
13 suitable for implementation in drug trials and clinical practice whereas others require substantial
14 improvement. An important consideration is that compared with immunoassays, currently
15 available research-based MS analytical technologies are more labor intensive and time
16 consuming with less throughput. However, with the development of commercial fully automated
17 MS platforms which have already increased capacity and speed with automated systems, MS
18 platforms can provide reasonable clinical access.

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28 **Supplementary material**

29 Supplementary material is available at *Brain* online.

30

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19
20

1 **Figure legends**

2
3 **Figure 1 ROC curve analysis for abnormal CSF A β 42/40 status and progression to**
4 **Alzheimer's disease dementia.** Receiver operating characteristic (ROC) curve analysis for
5 differentiating (A) mild cognitive impairment (MCI) participants with abnormal CSF amyloid- β
6 (A β)42/40 from those with normal CSF A β 42/40 and (B) MCI patients who progressed to
7 Alzheimer's disease dementia during follow-up from those who did not (stable MCI patients and
8 MCI patients who progressed to other types of dementia).

9
10 **Figure 2 Plasma p-tau biomarkers in amyloid-negative and -positive MCI patients.** Plasma
11 levels of phosphorylated tau (p-tau)217 (A-C), p-tau181 (D-E, G-J) and p-tau231 (F) measured
12 using different assays in the amyloid- β (A β) negative (A-) and A+ mild cognitive impairment
13 (MCI) groups. A β status was defined based on the CSF A β 42/40 ratio. Data are presented as a
14 fold change from the mean of the A- MCI group. Two p-tau217^{WashU} and p-tau217^{Janss} outliers in
15 the A+ group and one p-tau181^{ADx} outlier in the A- group are not shown in (A), (C) and (D) but
16 these data were included in the statistical analysis. F-values and p-values are from univariate
17 general linear models adjusted for age and sex. Boxes show interquartile range, the horizontal
18 lines are medians and the whiskers and outliers were plotted using the Tukey method.

19
20 **Figure 3 Plasma p-tau biomarkers in MCI participants who progressed to Alzheimer's**
21 **disease dementia during follow-up and amyloid-negative and -positive non-progressors.**
22 Plasma levels of phosphorylated tau (p-tau)217 (A-C), p-tau181 (D-E, G-J) and p-tau231 (F)
23 measured using different assays in patients with mild cognitive impairment (MCI) who
24 progressed to Alzheimer's disease dementia during follow-up (MCI-ADD), amyloid- β negative
25 (A-) and A+ non-progressor MCI patients. A β status was defined based on the CSF A β 42/40
26 ratio. Data are presented as a fold change from the mean of the A- MCI group. Two p-
27 tau217^{WashU} and p-tau217^{Janss} outliers in the MCI-ADD group and one p-tau181^{ADx} outlier in the
28 A- group are not shown in (A), (C) and (D) but these data were included in the statistical
29 analysis. F-values and p-values are from univariate general linear models adjusted for age, sex

1 and follow-up time. Boxes show interquartile range, the horizontal lines are medians and the
2 whiskers and outliers were plotted using the Tukey method.

3

4 **Figure 4 Correlations between CSF and plasma p-tau.** Heatmap showing Spearman
5 coefficients for correlations between plasma CSF and plasma p-tau measured using different
6 assays (p-tau181^{UGOT}, N=72; p-tau181^{Splex}, N=52; all other biomarker N=78). Correlations
7 between plasma and CSF p-tau measured with the same assay are highlighted in orange except
8 plasma p-tau181^{Lilly} and p-tau181^{Splex} for which corresponding CSF assay data were not
9 available.

10

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1 **Table I Demographic and clinical characteristics**

	Overall	Non-progressors A^{-a}	Non-progressors A⁺a	MCI-ADD
N	135	64	26	45
Age, years	74.0 (66.0–79.0)	70.5 (63.0–76.8)	72.0 (65.0–76.0)	78.0 (73.5–81.0)
Female, n (%)	82 (60.7)	37 (57.8)	11 (42.3)	34 (75.6)
MMSE	28.0 (26.0–29.0)	28.0 (27.0–29.0)	28.0 (27.0–29.3)	26.0 (25.0–27.0)
APOE ε4 positivity, n (%)	75 (55.6)	19 (29.7)	20 (76.9)	36 (80.0)
Follow-up time, years	4.6 (3.3–6.6)	6.21 (4.02–7.21)	5.16 (3.90–6.64)	3.64 (2.68–4.65)
Plasma p-tau				
p-tau217 ^{WashU} , %	1.36 (0.742–3.25)	0.753 (0.614–0.951)	1.88 (1.27–2.73)	3.49 (2.91–4.73)
p-tau217 ^{Lilly} , pg/ml ^b	0.247 (0.170–0.404)	0.177 (0.146–0.201)	0.275 (0.200–0.359)	0.442 (0.330–0.532)
p-tau217 ^{Janss} , pg/ml ^b	0.055 (0.030–0.105)	0.034 (0.020–0.049)	0.066 (0.036–0.104)	0.109 (0.077–0.173)
p-tau181 ^{ADx} , pg/ml	29.7 (19.3–46.3)	19.5 (10.4–27.3)	30.0 (22.8–45.0)	46.3 (38.8–63.7)
p-tau181 ^{WashU} , %	23.5 (19.8–28.7)	20.3 (18.2–22.7)	24.5 (20.7–29.0)	28.4 (25.7–32.1)
p-tau231 ^{UGOT} , pg/ml	20.9 (15.7–27.3)	16.8 (12.7–21.4)	22.0 (17.6–27.2)	26.9 (22.6–33.1)
p-tau181 ^{Lilly} , pg/ml	1.90 (1.42–2.59)	1.57 (1.20–1.90)	1.77 (1.49–2.26)	2.59 (2.04–3.30)
p-tau181 ^{UGOT} , pg/ml ^c	2.46 (1.72–3.55)	1.88 (1.49–2.58)	2.43 (1.89–3.45)	3.38 (2.58–4.07)
p-tau181 ^{Fuji} , pg/ml	4.80 (3.64–5.75)	3.83 (3.01–5.14)	4.73 (3.74–5.57)	5.61 (4.77–6.25)
p-tau181 ^{Splex} , pg/ml ^c	1.07 (0.859–1.55)	0.999 (0.792–1.22)	0.927 (0.754–1.73)	1.28 (1.05–2.16)

2 A=amyloid-β; ADD=Alzheimer's disease dementia; MCI=mild cognitive impairment; MMSE=Mini Mental State Examination; p-
3 tau=phosphorylated tau.

4 ^aAβ status was defined using the CSF Aβ42/40 cutoff (0.07) as described in the methods. Data are shown as median (IQR) unless otherwise
5 specified.

6 ^bp-tau217^{Lilly} and p-tau217^{Janss} data in overlapping sample have been reported previously.¹⁹

7 ^cp-tau181-UGOT and p-tau181-Splex data were available for 124 and 101 participants, respectively.

8

9

1 **Table 2 Analytical performance of plasma p-tau assays**

Plasma biomarkers	Required plasma volume, ml	Intra-assay CV, %	Inter-assay CV, %	Samples below LLOD, %	LLOD, pg/ml
p-tau217 ^{WashU}	1 ^a	3.3 ^b	3.5 ^b	0	NA ^c
p-tau217 ^{Lilly}	0.07	6.8	10.1	15.6	0.150
p-tau217 ^{Janss}	0.2	23.7	12.4	0	0.013
p-tau181 ^{ADx}	0.1	11.1	3.8	16.3	2.312
p-tau181 ^{WashU}	1 ^a	3.7 ^b	0.4 ^b	0	NA ^c
p-tau231 ^{UGOT}	0.8	7.6	8.5	0	1
p-tau181 ^{Lilly}	0.07	6.0	11.2	0	0.864
p-tau181 ^{UGOT}	0.8	8.2	10.9	0	0.5
p-tau181 ^{Fuji}	0.13	NA ^d	NA ^d	0	0.052
p-tau181 ^{plex}	0.06	4.8	13.5	0	0.190

2 CV=coefficient of variation; LLOD=lower limit of detection; p-tau=phosphorylated tau.

3 ^a1 ml was required for the entire multiplex assay.

4 ^bCVs were estimated using quality control samples; study samples were tested in singlicate.

5 ^cNot applicable for phosphorylation occupancy measures.

6 ^dNot applicable, samples in this study were tested in singlicate in one run.

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1 **Table 3 Associations of plasma p-tau with CSF A β 42/40**

Plasma p-tau	AUC (95% CI)	P-value versus p-tau217 ^{WashU}	P-value versus p-tau217 ^{Lilly}	Specificity (95% CI)	Sensitivity (95% CI)	Accuracy (95% CI)
p-tau217 ^{WashU}	0.947 (0.907–0.987)	NA	0.015	90.6 (82.8–98.4)	94.4 (84.5–98.6)	92.6 (88.1–96.3)
p-tau217 ^{Lilly}	0.886 (0.827–0.944)	0.015	NA	84.4 (71.9–96.9)	85.9 (67.6–95.8)	84.4 (78.5–90.4)
p-tau217 ^{Inss}	0.858 (0.795–0.920)	0.004	0.38	87.5 (65.6–95.3)	74.6 (60.6–91.5)	80.0 (73.3–86.7)
p-tau181 ^{ADx}	0.841 (0.768–0.913)	<0.001	0.24	85.9 (68.8–95.3)	77.5 (66.2–93.0)	81.5 (74.8–87.4)
p-tau181 ^{WashU}	0.835 (0.765–0.906)	<0.001	0.20	87.5 (73.4–95.3)	76.1 (64.8–88.7)	81.5 (74.8–87.4)
p-tau231 ^{UGOT}	0.784 (0.703–0.864)	<0.001	0.029	73.4 (46.9–87.5)	78.9 (64.8–98.6)	76.3 (69.6–82.2)
p-tau181 ^{Lilly}	0.759 (0.676–0.841)	<0.001	<0.001	78.1 (65.6–89.1)	71.8 (60.6–84.5)	75.6 (68.1–82.2)
p-tau181 ^{UGOT^a}	0.743 (0.652–0.833)	<0.001	0.005	70.2 (50.9–86.0)	79.1 (59.7–92.5)	74.2 (66.9–81.5)
p-tau181 ^{Huji}	0.694 (0.604–0.784)	<0.001	<0.001	56.3 (40.6–85.9)	84.5 (50.7–93.0)	69.6 (62.2–76.3)
p-tau181 ^{Splex^a}	0.642 (0.533–0.751)	<0.001	<0.001	79.6 (22.4–98.0)	53.8 (26.9–100.0)	65.3 (58.4–73.3)

2 Data are from ROC curve analysis. MCI participants were classified as amyloid-negative (n=64) or as amyloid-positive (n=71) using CSF
 3 A β 42/40 as described in the methods. AUC=area under the curve; CSF = cerebrospinal fluid; CI=confidence interval; MCI=mild cognitive
 4 impairment; ROC=Receiver Operating Characteristic; p-tau=phosphorylated tau.

5 ^ap-tau181-UGOT and p-tau181-Splex data were available for 124 (57 amyloid-negative, 67 amyloid-positive) and 101 (49 amyloid-negative, 52
 6 amyloid-positive) participants, respectively.

7
 8

1 **Table 4 Associations of plasma p-tau with future progression to Alzheimer's disease dementia**

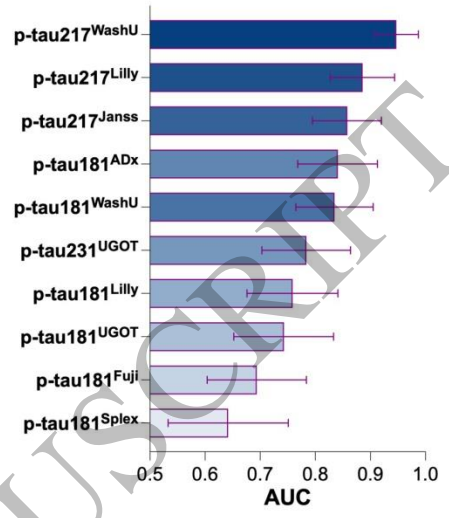
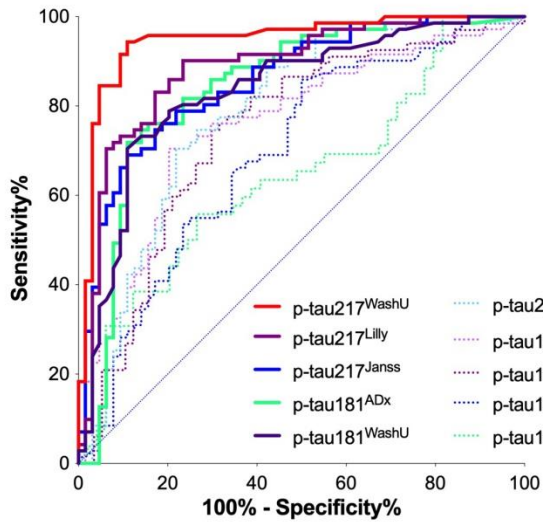
Plasma p-tau	AUC (95% CI)	P-value versus p-tau217 ^{WashU}	P-value versus p-tau217 ^{Lilly}	Specificity (95% CI)	Sensitivity (95% CI)	Accuracy (95% CI)
p-tau217 ^{WashU}	0.932 (0.891–0.974)	NA	0.027	86.7 (77.8–93.3)	95.6 (84.4–100.0)	88.9 (83.7–94.1)
p-tau217 ^{Lilly}	0.889 (0.833–0.946)	0.027	NA	83.3 (65.6–93.3)	88.9 (73.3–100.0)	84.4 (75.6–90.4)
p-tau217 ^{Janss}	0.872 (0.814–0.931)	0.027	0.53	74.4 (61.1–91.1)	91.1 (71.1–100.0)	80.0 (71.9–87.4)
p-tau181 ^{ADx}	0.846 (0.777–0.916)	0.007	0.16	81.1 (72.2–88.9)	91.1 (80.0–97.8)	84.4 (77.8–90.4)
p-tau181 ^{WashU}	0.835 (0.764–0.906)	0.001	0.09	76.7 (64.4–86.7)	88.9 (77.8–97.8)	80.7 (72.6–86.7)
p-tau181 ^{Lilly}	0.813 (0.734–0.892)	0.002	0.013	74.4 (60.0–86.7)	86.7 (71.1–97.8)	77.8 (70.4–85.2)
p-tau231 ^{UGOT}	0.777 (0.699–0.856)	<0.001	0.009	68.9 (57.8–81.1)	86.7 (73.3–95.6)	74.8 (67.4–81.5)
p-tau181 ^{UGOT a}	0.775 (0.692–0.858)	<0.001	0.014	65.9 (52.4–82.9)	88.1 (69.0–97.6)	73.4 (64.5–81.5)
p-tau181 ^{Huji}	0.735 (0.649–0.821)	<0.001	0.002	70.0 (40.0–86.7)	75.6 (53.3–97.8)	71.1 (57.8–79.3)
p-tau181 ^{Splex a}	0.688 (0.579–0.796)	<0.001	<0.001	66.7 (50.0–90.9)	74.3 (42.9–91.4)	69.3 (59.4–78.2)

2 Data are from ROC curve analysis. 45 MCI participants progressed to Alzheimer's disease dementia during follow-up and 90 remained stable or
 3 progressed to non-Alzheimer's disease dementia. AUC=area under the curve; CI=confidence interval; MCI=mild cognitive impairment;
 4 ROC=Receiver Operating Characteristic; p-tau=phosphorylated tau

5 ^ap-tau181-UGOT and p-tau181-Splex data were available for 124 (82 non-progressors, 42 MCI-ADD) and 101 (66 non-progressors, 35 MCI-
 6 ADD) participants, respectively.

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A A- MCI vs A+ MCI



B Non-progressors vs progressors

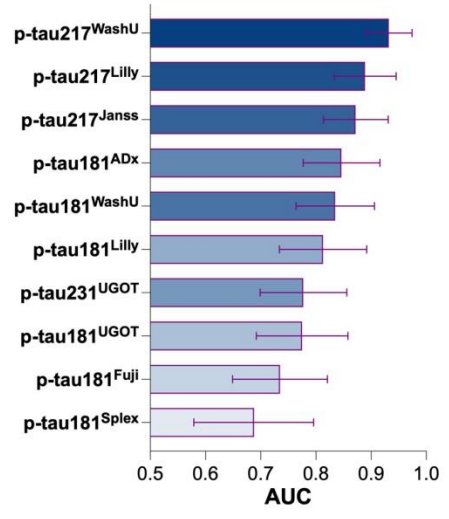
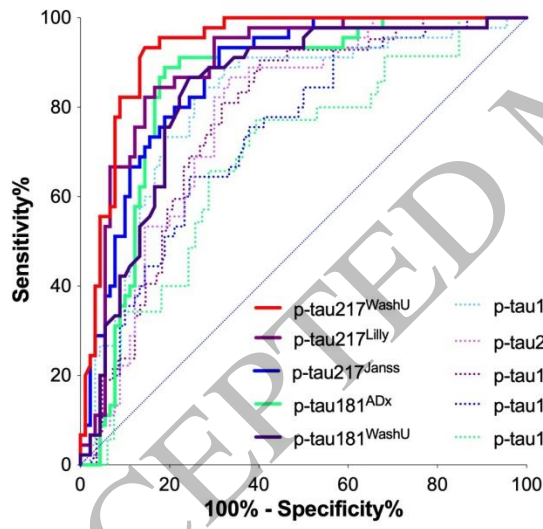


Figure 1
199x204 mm (.33 x DPI)

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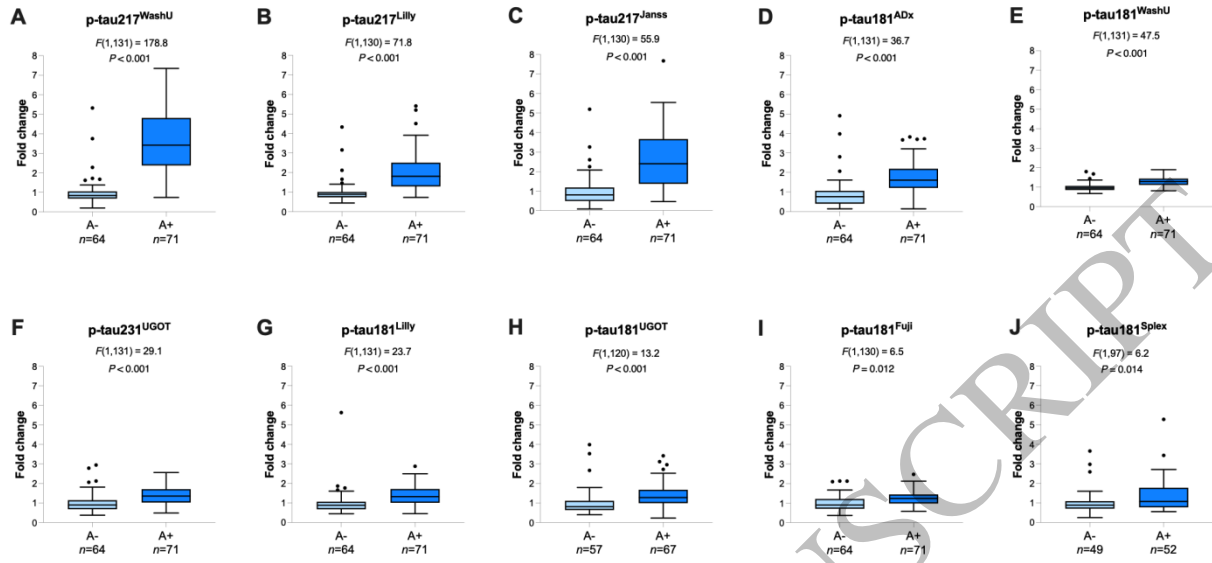
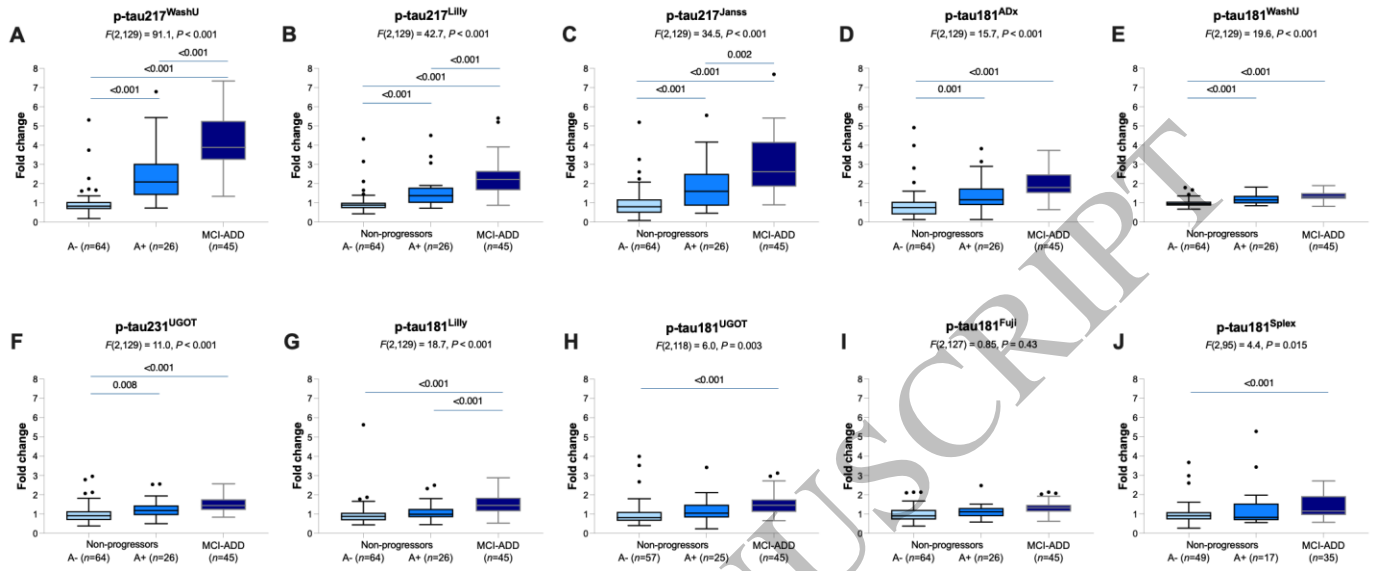


Figure 2
185x85 mm (.33 x DPI)

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Figure 3
185x76 mm (.33 x DPI)

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		CSF							
		p-tau217 ^{WashU}	p-tau217 ^{Lilly}	p-tau217 ^{Imms}	p-tau181 ^{ADx}	p-tau181 ^{WashU}	p-tau231 ^{UGOT}	p-tau181 ^{UGOT}	p-tau181 ^{Fuji}
Plasma	p-tau217 ^{WashU}	0.891	0.858	0.837	0.836	0.789	0.682	0.696	0.766
	p-tau217 ^{Lilly}	0.805	0.755	0.747	0.722	0.754	0.630	0.643	0.654
	p-tau217 ^{Imms}	0.700	0.682	0.659	0.619	0.645	0.487	0.513	0.589
	p-tau181 ^{ADx}	0.685	0.665	0.667	0.669	0.633	0.597	0.592	0.571
	p-tau181 ^{WashU}	0.716	0.681	0.659	0.648	0.580	0.521	0.544	0.621
	p-tau231 ^{UGOT}	0.427	0.421	0.406	0.376	0.366	0.320	0.320	0.368
	p-tau181 ^{UGOT}	0.436	0.449	0.469	0.420	0.369	0.414	0.410	0.441
	p-tau181 ^{Fuji}	0.421	0.384	0.368	0.362	0.329	0.257	0.288	0.372
	p-tau181 ^{Lilly}	0.535	0.537	0.512	0.490	0.547	0.427	0.423	0.445
	p-tau181 ^{Splex}	0.279	0.327	0.291	0.308	0.255	0.316	0.340	0.321

Figure 4
90x87 mm (.33 x DPI)

ACCEPTED MANUSCRIPT