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Cross-sectional and longitudinal atrophy is preferentially associated with tau rather than amyloid β positron emission tomography pathology

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Abstract

Introduction: Structural magnetic resonance imaging is a marker of gray matter health and decline that is sensitive to impaired cognition and Alzheimer's disease pathology. Prior work has shown that both amyloid β (A β) and tau biomarkers are related to cortical thinning, but it is unclear what unique influences they have on the brain.

Methods: A β pathology was measured with [¹⁸F] AV-45 (florbetapir) positron emission tomography (PET) and tau was assessed with [¹⁸F] AV-1451 (flortaucipir) PET in a population of 178 older adults, of which 123 had longitudinal magnetic resonance imaging assessments (average of 5.7 years) that preceded the PET acquisitions.

Results: In cross-sectional analyses, greater tau PET pathology was associated with thinner cortices. When examined independently in longitudinal models, both A β and tau were associated with greater antecedent loss of gray matter. However, when examined in a combined model, levels of tau, but not A β , were still highly related to change in cortical thickness.

Discussion: Measures of tau PET are strongly related to gray matter atrophy and likely mediate relationships between A β and gray matter.

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Keywords:

Thickness; Atrophy; Tau; Amyloid; PET; Positron emission tomography; MRI

1. Introduction

One of the earliest established and most widely replicated neuroimaging findings in Alzheimer's disease (AD) is a reduction in gray matter and greater rates of longitudinal atrophy [1–5] seen with magnetic resonance imaging (MRI).

Similarly, lower baseline levels and greater gray matter atrophy have been found in participants with mild cognitive impairment [6], subjective memory complaints [7], and cognitively normal individuals with abnormal levels of amyloid β (A β) [4,5,8–10] relative to those without any AD pathology. In a clinical setting, structural pathological MRI measures and rates of change have prognostic value, predict conversion from mild cognitive impairment to AD [11–14], and an increased risk of later AD dementia in cognitively normal individuals [15].

While much of the focus has understandably been on the medial temporal lobes, loss of gray matter is not restricted to

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these regions. Analyses looking at whole-brain spatial patterns see loss of gray matter prominently throughout the temporal, parietal, frontal, and occipital regions [3,4,6]. Prior work shows that such structural atrophy is related to levels of A β [5,10,16,17], but increased levels of A β may simply be a proxy for other AD-related pathologies such as neurofibrillary tangles (NFTs).

Positron emission tomography (PET) ligands that bind to NFTs [18–21] have provided a new biomarker to understand AD. Tau PET is elevated in AD [22–28] and in cognitively normal individuals with elevated A β pathology [23,25,29,30]. Elevated tau PET binding predicts other neurodegenerative biomarkers such as hypometabolism [31–33] and structural atrophy [34–36].

Owing to the accessibility and prevalence of structural MRI imaging, it is critical to understand the relationship between tau PET and gray matter integrity. Prior work has examined cross-sectional relationships between tau PET using AV-1451 and cortical thickness [34–37], finding that increased levels of tau PET binding were associated with thinner cortices, particularly in temporal, lateral parietal, and occipital regions. There is also initial evidence that prior longitudinal changes in structural MRI over a limited time window (~ 3 years) predict current levels of NFT pathology measured with PET [35]. This cross-sectional and longitudinal work reveals that there is a clear relationship between in vivo measures of tau pathology and structural integrity but has only evaluated the influence of tau pathology without also additionally considering measures of A β .

In a population of cognitively normal and mildly impaired older adults, the current work examines the influence of A β and NFT pathology on structural integrity. We examine both the concurrent relationship between cortical thickness and pathology and the relationship between current levels of pathology and antecedent longitudinal change in cortical thickness. By considering both pathologies simultaneously, it is possible to estimate the unique influence each has on structural atrophy. This is critical not only to understand how tau pathology is related to another marker of neurodegeneration but also to explore whether tau pathology mediates prior observed relationships between A β and gray matter health.

2. Methods

2.1. Participants

Participants were selected from ongoing studies on aging and dementia from the Knight Alzheimer's Disease Research Center at Washington University. Cognitive status was assessed using the clinical dementia rating (CDR) [38]. Participants were required to have tau PET imaging and structural MRI at their most recent neuroimaging assessment. The cross-sectional cohort consisted of 178 individuals (age 46–91 years) with either no cognitive impairment ($n = 156$, CDR = 0) or very mild dementia ($n = 22$,

CDR = 0.5). From this initial population, 123 individuals had at least one MRI session that preceded the acquisition of tau PET (age 55–91 years; $n = 111$, CDR = 0; $n = 12$, CDR = 0.5). Full demographics for the cohort at the time of the tau PET session are presented in Table 1.

2.2. MRI acquisition and processing

T1-weighted images were acquired using a magnetization-prepared rapid gradient-echo sequence on a 3T Siemens scanner. Scans had a resolution of either $1 \times 1 \times 1.25$ mm or $1 \times 1 \times 1$ mm. Structural scans were processed with FreeSurfer [39]. For each hemisphere, cortical thickness values were obtained for all FreeSurfer cortical regions of interest (ROIs), and volumes were obtained for all FreeSurfer subcortical ROIs. Cortical thickness was calculated as the shortest distance between the cortical gray/white boundary and the gray/cerebrospinal fluid (CSF) boundary [40]. Cortical surface meshes are placed into a common anatomical space within FreeSurfer for vertex-wise analyses. Parcellations of the T1-weighted image into cortical and subcortical regions were also performed for utilization in the processing of PET data. Longitudinal data were processed through the FreeSurfer 5.3 longitudinal stream [41]. There were a total of 390 sessions in the longitudinal analyses (mean 3.1 visits, 5.7 years).

2.3. PET acquisition and processing

A β PET imaging was completed using [18 F] AV-45 (florbetapir). Data from the 50- to 70-minute postinjection window were analyzed using an in-house pipeline using ROIs derived from FreeSurfer [39,42] (PET Unified Pipeline, <https://github.com/ysu001/PUP>). Tau PET imaging was performed using [18 F] AV-1451 (flortaucipir). Data from the 80- to 100-minute postinjection window were analyzed. For both tracers, regional estimates were transformed into standardized uptake value ratios using a cerebellar cortex reference region, although alternate

Table 1
Sample demographics

Variable	Cross-sectional cohort (N = 178)	Longitudinal cohort (N = 123)
Age, years	70.1 (8.2)	70.8 (8.4)
Female (%)	106 (60)	74 (60)
MMSE	29.0 (1.8)	29.1 (1.9)
CDR-sum	0.3 (0.8)	0.2 (0.7)
AV-45 summary (SUVR)	1.27 (0.64)	1.29 (0.66)
AV-1451 summary (SUVR)	1.29 (0.35)	1.26 (0.30)
Months between AV-45 and AV-1451	3.6 (6.7)	4.3 (7.8)
Follow-up (years)		5.7 (2.4)
Follow-up (visits)		3.1 (1.1)

Abbreviations: CDR-sum, Clinical Dementia Rating–sum of boxes; MMSE, Mini-Mental Status Examination; SUVR, standardized uptake value ratio.

NOTE. Unless otherwise noted values represent means and standard deviations.

reference regions are also used in the literature [43]. Data were partial volume corrected using a regional spread function technique [44,45]. This approach corrects for the spillover signal from adjacent ROIs based upon the scanner point spread function and the relative distance between ROIs. ROI PET data were averaged across hemispheres for statistical analyses. A summary global A β [42] and tau [30] measures were calculated as previously published. On average, PET data were acquired within 3.6 months of one another.

2.4. Statistical analyses

Cross-sectional vertex-wise analyses were performed using the FreeSurfer Qdec application (www.surfer.nmr.mgh.harvard.edu). Three statistical models were examined using general linear models. The first related the summary measure of tau [30], as measured with flortaucipir, to vertex measures of cortical thickness. The second model related the A β summary measure, as measured with florbetapir, to vertex measurements of cortical thickness. Finally, to assess the unique influence of each pathology, the last model included both the A β and tau summary measures. All three models were run separately for the left and right hemispheres and included current age, gender, and scan resolution ($1 \times 1 \times 1$ or $1 \times 1 \times 1.25$) as covariates. Vertices were considered significant after a false discovery rate correction considering both hemispheres.

Longitudinal analyses were performed two ways. The first set of analyses examined relationships of pathology with antecedent longitudinal change in cortical thickness at the cortical surface level using spatiotemporal linear mixed-effects (LMEs) models [14,46] implemented in MATLAB. LMEs are highly flexible approaches for analyzing longitudinal neuroimaging data because they can handle unequal numbers of data points and temporal spacing between assessments across subjects. The spatiotemporal approach is an extension of LMEs for mass-univariate analyses but also takes into account pooled covariance structure across neighboring vertices of homogeneous regions. Three models were again run and were similar in structure to the cross-sectional models. In the first model, preceding longitudinal change in cortical thickness was predicted by the main effects of our current tau PET summary measure, gender, current age, time, scan resolution ($1 \times 1 \times 1$ or $1 \times 1 \times 1.25$), and the interaction between tau PET and time. Models included random subject-specific slopes and intercepts. The second model implemented was identical in structure but used A β PET rather than tau as a predictor. The third, combined model examined the main effects and interactions with time of both global tau and A β PET. All three models were implemented separately for the left and right hemispheres, and vertices were considered significant after surviving false discovery rate correction accounting for comparisons in both hemispheres.

Finally, a series of LME data were run using ROIs derived from FreeSurfer to examine the effects of local pathology on local atrophy. Regional data were averaged between left and right hemispheres. Instead of examining the association between global summaries of A β and tau and cortical thickness, these analyses specifically modeled A β and tau PET binding within each ROI (e.g., precuneus) to thickness in that same ROI. This was done to avoid any bias that could be introduced by using summary measures formed from specific ROIs. The first model looked at the relationship between longitudinal measurements of thickness in that ROI predicted by main effects of current tau in that specific ROI, gender, current age, time, scan resolution, and the time by tau interaction. Models contained subject-specific random-effects intercepts and slopes. Similar models were run using A β rather than tau PET as a predictor, and finally combined models including both tau and A β and their interactions with time. These three models were run for all 34 cortical regions identified by FreeSurfer. Models were implemented using the lme4 package in R, version 3.4.1, and ROIs were considered significant after false discovery rate correction.

3. Results

3.1. Cross-sectional associations with global pathology

The vertex-wise results are presented in the left-hand column of Fig. 1. Fig. 1A depicts those vertices that had a significant relationship with the global tau PET measure. Fig. 1B depicts those vertices significantly related to the global A β PET measure. Fig. 1C shows those vertices significantly related to tau while additionally controlling for levels of A β . There were no vertices significantly related to A β when entered alone (Fig. 1B) or when tau was also included in the model (not shown). For simplicity, only the results from the left hemisphere are presented. Results for the right hemisphere are highly similar and presented in Supplementary Fig. 1.

3.2. Longitudinal associations with global pathology

Results from the spatiotemporal LMEs are presented in the right-hand column of Fig. 1. Fig. 1D shows those areas of the cortex that showed a significant global tau burden by time interaction, that is, how current levels of tau predict antecedent thickness change in cortical thickness. Fig. 1E depicts the A β by time interaction. Fig. 1F shows the tau by time interaction after additionally controlling for the main effect of global A β burden and A β by time interaction. There were no significant A β by time effects once tau was simultaneously considered in the model.

3.3. Longitudinal associations with local pathology

Results examining the relationships between antecedent structural change and local AD pathologies are

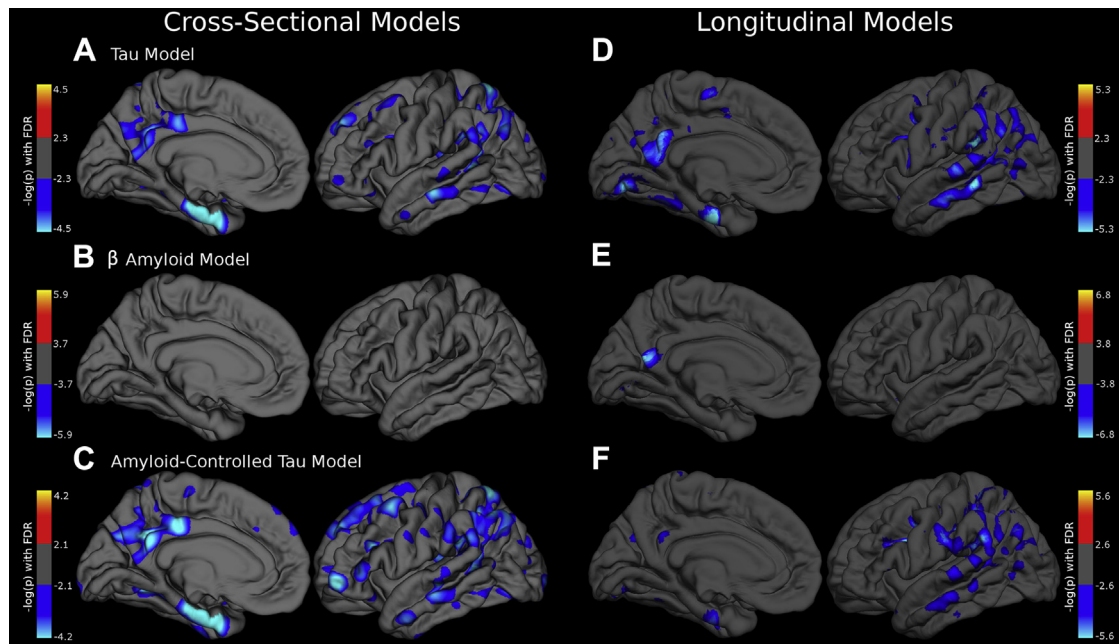


Fig. 1. The relationship between global AD pathology and gray matter. Three GLMs were implemented to estimate vertex-wise relationships between (A) global tau PET summary measure, (B) A β mean cortical SUVR, or (C) global tau controlling for A β , and left-hemisphere cross-sectional cortical thickness. Spatiotemporal LMEs models were used to investigate relationships between (D) global tau and time from initial MRI, (E) global A β and time, or (F) global tau and time controlling for A β and time, and antecedent longitudinal change in cortical thickness. All models controlled for the main effects of gender, current age, and scan resolution. Cross-sectional models used the Qdec multiple comparisons correction (FDR) procedure at $P < .025$ level to approximate comparisons in both hemispheres. Longitudinal models used the LME FDR2 comparisons correction in MATLAB and accounted for comparisons in both hemispheres. Values depicted from each model are thresholded at the level of significance identified by their respective FDR procedures. Abbreviations: A β , amyloid β ; AD, Alzheimer's disease; FDR, false discovery rate; GLM, general linear model; LMEs, linear mixed-effects; MRI, magnetic resonance imaging; PET, positron emission tomography; SUVR, standardized uptake value ratio.

shown in Fig. 2 and presented in Table 2. To be visually consistent with Fig. 1, results are displayed on the left hemisphere but represented the statistical results from the average of data from left and right hemispheres. Fig. 2A depicts the significant regional tau burden by time interaction (e.g., the relationship between precuneus tau and change in precuneus cortical thickness). Fig. 2B depicts the local A β by time interaction. Fig. 2C depicts the local tau by time interaction when both tau and A β main effects and interactions were considered in the model simultaneously. There were no significant associations between local A β and change in cortical thickness once tau was additionally considered in the model.

4. Discussion

Prior work has demonstrated that structural MRI is a sensitive marker to AD dementia [1–5] and preclinical AD [4,5,8–10]. The advent of tau PET imaging has provided another *in vivo* biomarker of the molecular pathology in AD. The current work examined the association between cortical thickness and tau pathology measured with flortaucipir PET. We found that current levels of tau PET binding, rather than A β PET, were related to concurrent cortical thinning and preceding structural atrophy.

We found that both global and local levels of flortaucipir were associated with greater antecedent atrophy. This is consistent with prior work in the field demonstrating structural declines in preclinical AD [4,5,8–10]. Early work has also indicated a significant relationship between tau PET binding and gray matter health [34–37]. When simultaneously considering both biomarkers, levels of tau PET mediated the effects of A β on cortical thickness. This suggests that in prior work, the association between gray matter and A β may have been a proxy for emerging NFT pathology. This also suggests that the relationship between tau PET and cortical thickness is distinct from the influence of A β .

The association between increasing levels of tau PET and cortical thinning was not restricted to the medial temporal lobe but was widespread throughout the temporal, occipital, parietal, and even portions of the frontal lobes. The effects in the temporal and parietal regions were particularly reminiscent of spatial signatures seen when examining whole-brain atrophy maps in AD [3–5], indicating a stable network of brain regions that also demonstrate tau-related atrophy. The pattern of areas demonstrating atrophy is also more similar to the spatial topography of tau rather than A β PET [22–28,31]. This effect was not however simply a product of the regions going into our summary measures. Analyses looking only at local tau and A β accumulation

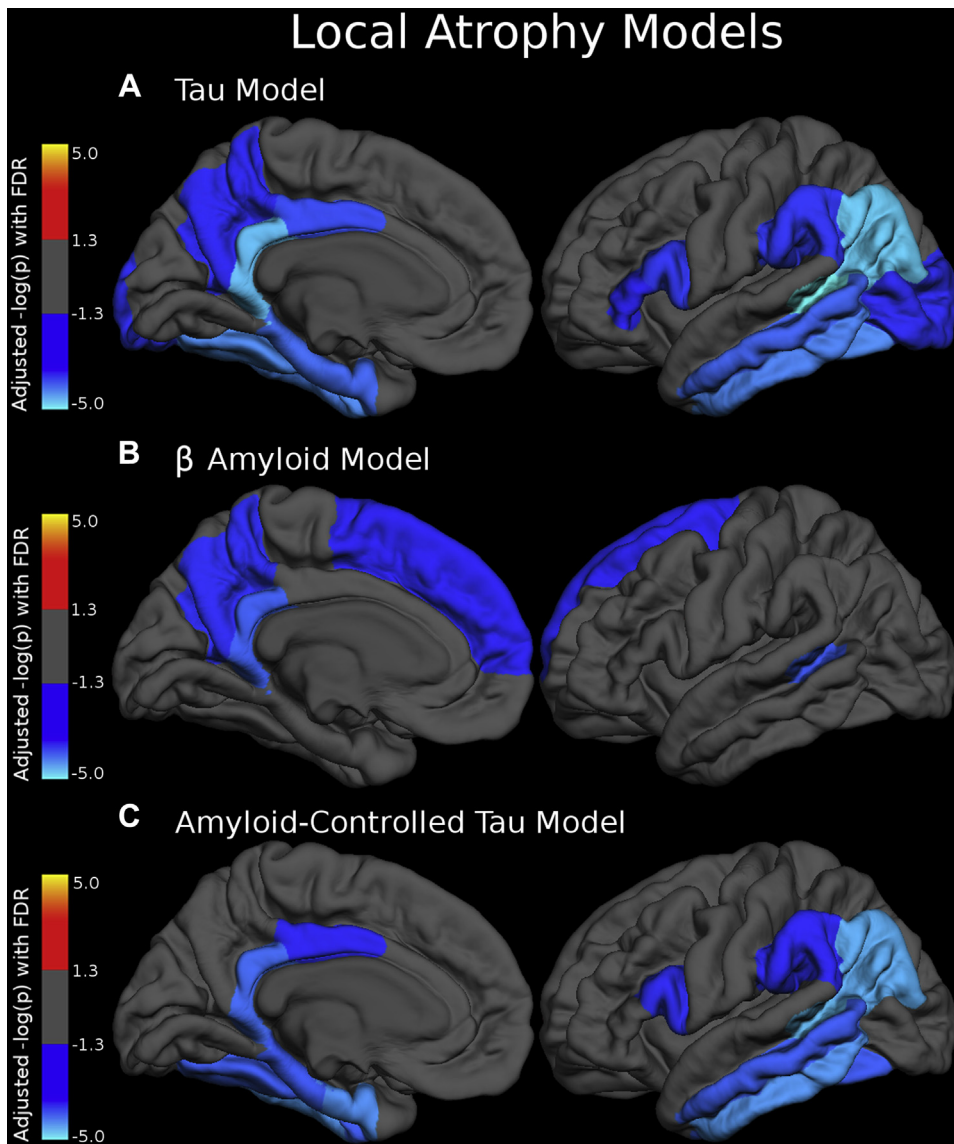


Fig. 2. The relationship between local AD pathology and gray matter. Linear mixed-effects models were implemented to model region-wise relationships between (A) local tau and time, (B) local A β and time, or (C) local tau and time controlling for local A β and time, on longitudinal local cortical thickness in all 34 FreeSurfer cortical regions. All models controlled for the main effects of gender, age, and scan resolution. Both hemispheres' values were averaged together. Values depicted are adjusted $-\log_{10}(P)$ values after FDR correction for multiple comparisons. Abbreviations: AD, Alzheimer's disease; FDR, false discovery rate.

recapitulated similar regional maps, suggesting a stronger local relationship between structural atrophy and tau than atrophy and A β .

The current work and prior results [35] indicate that NFT pathology measured with flortaucipir is related to cross-sectional cortical thinning, as well as atrophy that has occurred over the previous years. In models of AD pathophysiology [47], as well as work with autosomal-dominant AD [48], changes in tau pathology occur before structural changes seen with MRI. Such frameworks would suggest that tau pathology would be a stronger predictor of *future* rather than *retrospective* atrophy. Such a temporal relationship, for example, tau leading MRI with a time lag, could

still account for the current results. For example, we also found that rates of MRI change were also related to current levels of A β , although there is strong evidence that elevations in A β plaque deposition precedes all other pathologies [48]. Rates of structural atrophy may simply serve as a rough marker of the disease stage. As a result, greater antecedent atrophy could predict current A β or tau PET, even if that is not the correct causative temporal direction.

Alternatively, the temporal ordering of tau relative to structural MRI has been based on CSF total tau and phosphorylated tau. Although CSF and PET measures of tau are modestly correlated [23,29,37], these two markers likely measure distinct but related aspects of tau. Just as there appears to be

Table 2
Regional effects relating tau and A β to longitudinal rates of atrophy

Region	A β \times time interaction			Tau \times time interaction			Tau \times time controlling for A β \times time		
	B	t-value	Adjusted P value	B	t-value	Adjusted P value	B	t-value	Adjusted P value
Banks of the superior temporal sulcus	-0.0039	-2.84	.00583	-0.0109	-5.15	.00000	-0.0109	-4.06	.00010
Caudal anterior cingulate cortex	0.0004	0.2709	.78705	-0.0049	-0.85	.39591	-0.0052	-0.89	.37662
Caudal middle frontal gyrus	-0.0025	-1.6629	.10095	-0.0074	-1.51	.13530	-0.0044	-0.78	.43531
Cuneus	0.0002	0.0719	.94276	-0.0042	-0.67	.50606	-0.0042	-0.67	.50548
Entorhinal cortex	0.0008	0.0928	.92640	-0.0209	-3.09	.00267	-0.0269	-3.61	.00054
Frontal pole	-0.0006	-0.2986	.76560	0.0063	1.18	.24033	0.0069	1.26	.20816
Fusiform gyrus	-0.0038	-1.8799	.06381	-0.0147	-3.55	.00060	-0.0147	-2.91	.00454
Inferior parietal cortex	-0.0021	-1.8018	.07619	-0.013	-4.18	.00005	-0.0147	-3.85	.00016
Inferior temporal cortex	-0.001	-0.6511	.51701	-0.0103	-3.43	.00085	-0.0132	-3.64	.00040
Insula	-0.0039	-1.7147	.09024	-0.0035	-0.57	.56754	-0.0017	-0.27	.79159
Isthmus cingulate	-0.0062	-2.9631	.00403	-0.0191	-4.02	.00009	-0.0158	-3.06	.00256
Lateral occipital cortex	-0.0016	-0.9272	.35694	-0.0078	-2.17	.03314	-0.0084	-1.97	.05146
Lateral orbitofrontal cortex	-0.0014	-0.9398	.35083	-0.0009	-0.16	.87056	0.0004	0.08	.93691
Lingual gyrus	-0.0032	-1.1853	.23911	-0.0086	-1.39	.16707	-0.0079	-1.26	.21096
Medial orbital frontal cortex	-0.0027	-1.7311	.08784	-0.0123	-1.90	.06128	-0.0114	-1.76	.08258
Middle temporal gyrus	-0.0015	-1.0751	.28622	-0.0096	-3.01	.00313	-0.0111	-2.90	.00434
Paracentral gyrus	-0.0037	-1.4636	.14808	-0.0035	-0.49	.62762	-0.0019	-0.27	.79104
Parahippocampal cortex	-0.0047	-1.1655	.24740	-0.02	-3.14	.00233	-0.0203	-2.86	.00519
Pars opercularis	-0.0024	-1.5936	.11246	-0.0138	-2.32	.02142	-0.0128	-2.13	.03453
Pars orbitalis	0.0001	0.0628	.95017	0.006	1.25	.21444	0.0063	1.27	.20763
Pars triangularis	-0.0004	-0.3179	.75076	-0.0099	-1.98	.04832	-0.0096	-1.92	.05556
Pericalcarine cortex	-0.0003	-0.1267	.89929	0.0026	0.31	.75614	0.0027	0.32	.74731
Postcentral gyrus	0.0006	0.2626	.79360	-0.0014	-0.21	.83138	-0.0017	-0.26	.79910
Posterior cingulate cortex	-0.0024	-1.9147	.05899	-0.0141	-2.67	.00874	-0.012	-2.05	.04273
Precentral gyrus	-0.0029	-0.9078	.36737	-0.0087	-0.84	.40198	-0.0068	-0.64	.52336
Precuneus cortex	-0.0024	-2.1989	.03179	-0.0098	-2.11	.03777	-0.0067	-1.30	.19534
Rostral anterior cingulate cortex	-0.0001	-0.0734	.94167	0.0017	0.25	.80612	0.0017	0.25	.80556
Rostral middle frontal	-0.0008	-0.6686	.50571	-0.0026	-0.51	.60828	-0.0023	-0.45	.65569
Superior frontal cortex	-0.0024	-2.0056	.04637	-0.0059	-1.13	.26050	-0.0039	-0.74	.45983
Superior parietal cortex	-0.0002	-0.1315	.89578	-0.0022	-0.47	.63724	-0.0024	-0.44	.66145
Superior temporal gyrus	-0.0021	-1.2591	.21213	-0.0072	-1.49	.13919	-0.0058	-1.08	.28262
Supramarginal gyrus	-0.0014	-0.9435	.34642	-0.0119	-2.30	.02254	-0.0115	-2.05	.04189
Temporal pole	0.0001	0.0149	.98810	-0.0045	-0.50	.61688	-0.0055	-0.55	.58304
Transverse temporal gyrus	-0.0026	-0.8029	.42423	0.0052	0.58	.56118	0.0071	0.77	.44330

Abbreviation: A β , amyloid β .

NOTE. Bold values represent multiple comparison adjusted *P* values <.05.

a temporal lag between CSF and PET measures of A β [49,50], there is likely a temporal delay between changes in CSF and PET measures of tau. As a result, tau PET may be later in disease progression and more proximal or even later to changes seen with structural MRI. Neuronal dysfunction and reduced dendritic branching may initially occur and manifest on MRI before mature tangles [20] that are bound by floratacypir form. Prospective acquisition of both longitudinal MRI and tau PET along with work in autosomal-dominant AD populations will help to clarify the temporal dynamics of tau PET relative to other markers.

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Supplementary data

Supplementary data related to this article can be found at <https://doi.org/10.1016/j.dadm.2018.02.003>.

RESEARCH IN CONTEXT

1. Systematic review: The authors reviewed the literature using traditional sources (e.g. PubMed and Google Scholar) for articles examining structural decline in preclinical and clinical Alzheimer's disease as well as for work using tau positron emission tomography imaging. Although tau positron emission tomography imaging has been related to neurodegenerative biomarkers, it is still not understood how tau pathology relates to cortical atrophy in relation to amyloid β deposition.
2. Interpretation: Our work found that tau rather than amyloid β predicts concurrent and antecedent gray matter loss. This result is consistent with neuropathology work indicating that tau pathology is a better marker of cognitive impairment than amyloid β .
3. Future directions: The manuscript found relationships between preceding changes in cortical integrity and current levels of tau pathology. Future work using longitudinal measures of both tau and magnetic resonance imaging and will help to further clarify the temporal dynamic between these markers.

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