

BIOMARKERS

PODIUM PRESENTATION

NEUROIMAGING

Heterogeneous tau patterns in atypical AD are explained by connectivity-associated tau progression

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Abstract

Background: The link between regional tau load and clinical manifestation of Alzheimer's disease (AD) highlights the importance of characterizing spatial tau distribution. In typical (memory-predominant) AD, the spatial progression of tau pathology mirrors the functional connections from temporal lobe epicenters. However, atypical (non-amnesic-predominant) AD variants with heterogeneous tau patterns provide a key opportunity to assess the universality of connectivity as a scaffold for tau progression.

Method: We included tau-PET data from 320 subjects with atypical AD, characterized by highly heterogeneous tau patterns ($n = 139$ posterior cortical atrophy/PCA-AD; $n = 103$ logopenic variant primary progressive aphasia/lvPPA-AD; $n = 35$ behavioural variant AD/bvAD; $n = 43$ corticobasal syndrome/CBS-AD) from 14 sites, with a subset of patients ($n = 78$) having longitudinal tau-PET data. As an independent sample, we further included regional post-mortem tau stainings from 93 atypical AD patients from two sites ($n = 19$ PCA-AD, $n = 32$ lvPPA-AD, $n = 23$ bvAD, $n = 19$ CBS-AD). Gaussian mixture modeling was used to harmonize different tau-PET tracers by transforming tau-PET standardized uptake value ratios to tau positivity probabilities (a uniform scale ranging from 0% to 100%). Using linear regression, we assessed whether 1) brain regions with stronger functional connectivity showed greater covariance in cross-sectional and longitudinal tau-PET and post-mortem tau pathology, and 2) functional connectivity of tau-PET epicenters and tau-PET accumulation epicenters was associated with cross-sectional and longitudinal tau patterns.

Result: Tau-PET epicenters—defined as the 5% brain regions with the highest tau load—aligned with clinical variants, e.g. a posterior pattern in PCA-AD (“visual AD”) and left-hemispheric temporal predominance in lvPPA-AD (“language AD”) (Figure 1). More strongly functionally connected regions showed correlated concurrent tau-PET levels, which was confirmed with post-mortem data (Figure 2). Moreover, the connectivity profile of tau-PET epicenters and accumulation epicenters corresponded to tau-PET progression patterns (Figure 3).

Conclusion: Our data are consistent with the hypothesis that tau propagation occurs along functional connections originating from local epicenters, across all AD clinical variants. Since tau proteinopathy is a key driver of neurodegeneration and cognitive decline, this finding may advance personalized medicine and participant-specific endpoints in clinical trials.

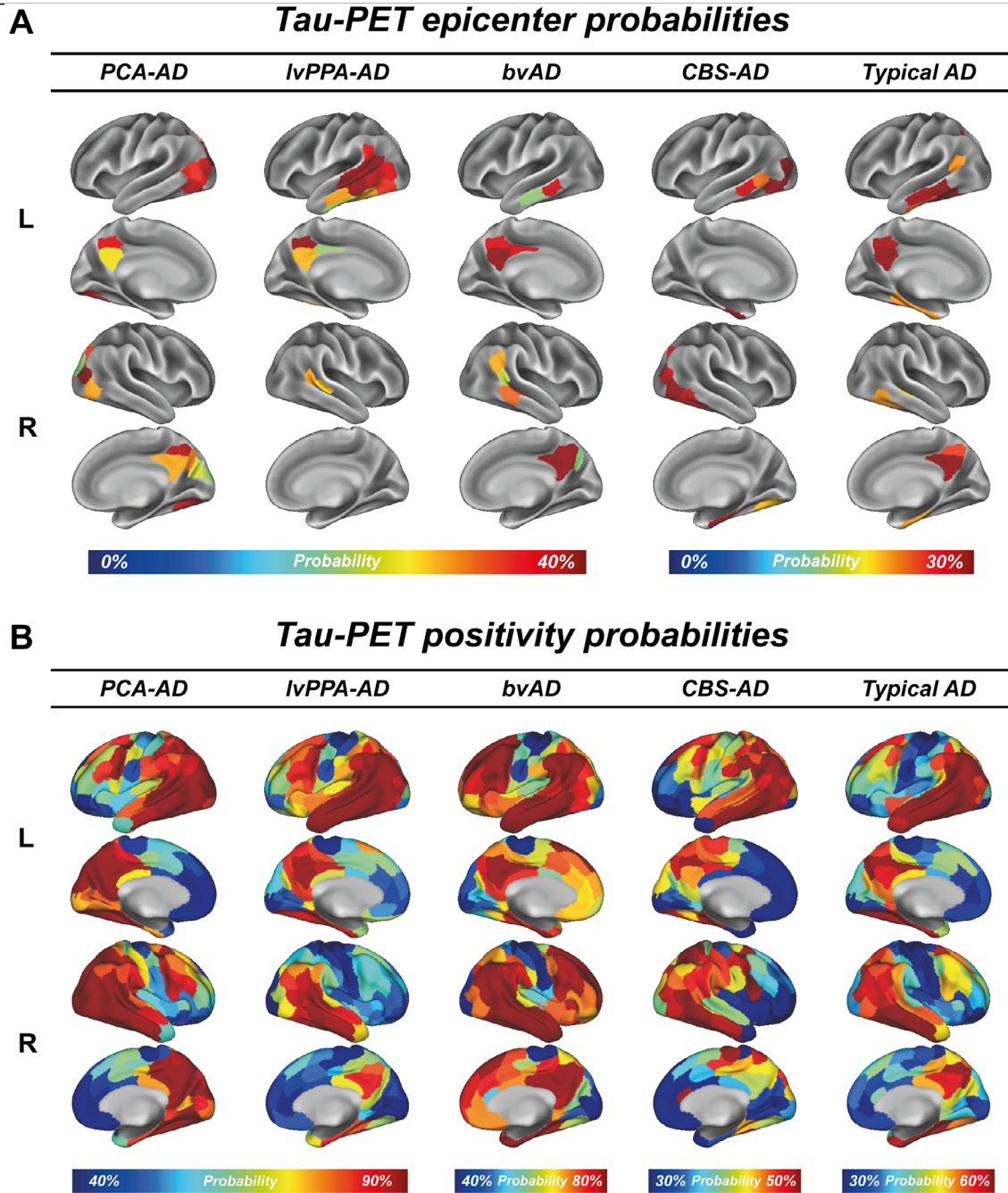


Figure 1: Tau-PET epicenters and positivity across AD variants. Tau epicenters (i.e., the regions with the assumed earliest and greatest tau burden) were defined at the subject level as the 5% regions with the highest tau-PET SUVRs. Group-average epicenter probabilities (A) indicate the likelihood of a region being part of the epicenter, with only epicenter probabilities $\geq 20\%$ shown. Group-average tau-PET positivity probabilities (a uniform tau-PET scale ranging from 0% to 100%) across AD variants are shown in (B). AD = Alzheimer's disease; bvAD = behavioural variant Alzheimer's disease; CBS = corticobasal syndrome; L = left; lvPPA = logopenic variant primary progressive aphasia; PCA = posterior cortical atrophy; PET = positron emission tomography; R = right; SUVR = standardized uptake value ratio.

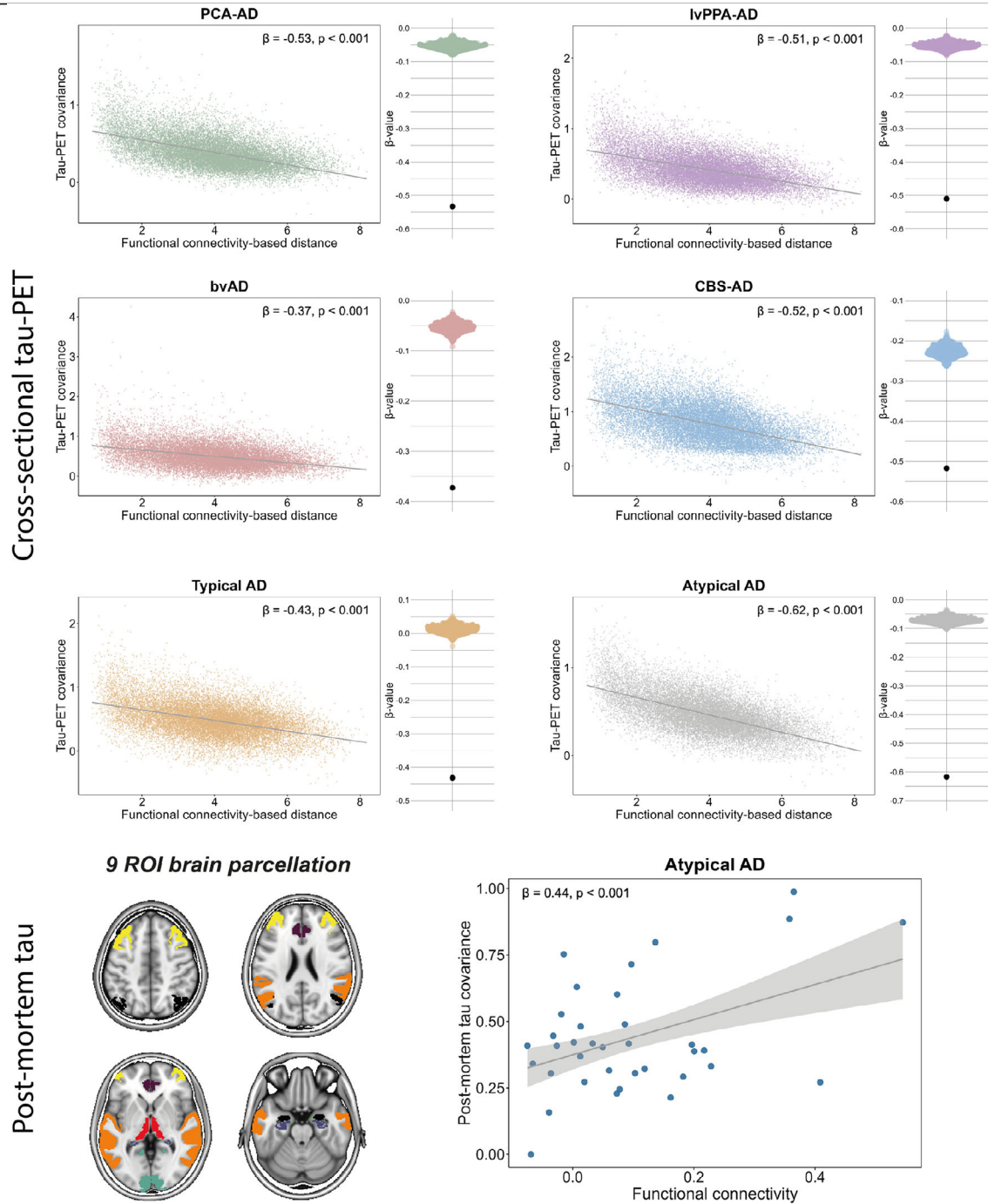


Figure 2: Association between functional connectivity and covariance in tau across variants of AD. Functional connectivity was derived from 42 CN A β -negative individuals from ADNI. Functional connectivity-based distance was used for tau-PET analyses, and functional connectivity for post-mortem analyses. Tau covariance was defined as average Fisher z-transformed partial correlations between tau positivity probabilities (tau-PET) or semi-quantitative tau pathology ratings (post-mortem tau) of all possible ROI pairs (200 Schaefer atlas for tau-PET, concatenated 9 ROI brain parcellation for post-mortem tau), while adjusting for age, sex, and site. A β = amyloid- β ; AD = Alzheimer's disease; ADNI = Alzheimer's disease neuroimaging initiative; bvAD = behavioural variant Alzheimer's disease; CBS = corticobasal syndrome; CN = cognitively normal; lvPPA = logopenic variant primary progressive aphasia; PCA = posterior cortical atrophy; PET = positron emission tomography; ROI = region of interest.

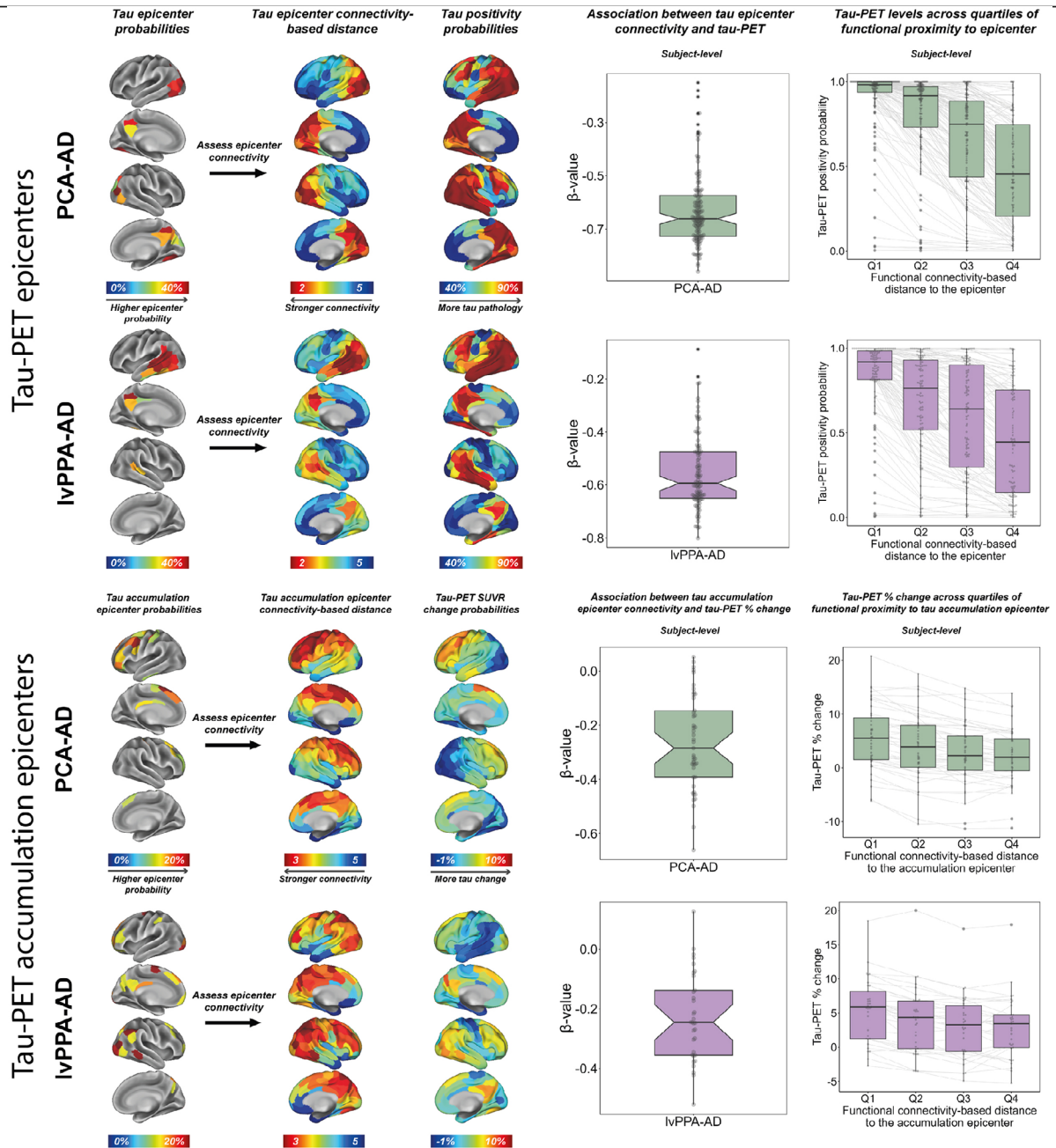


Figure 3: Association between tau (accumulation) epicenter connectivity and tau-PET (change) in PCA-AD and IvPPA-AD. Tau epicenters were defined as the top 5% of ROIs with the highest tau-PET SUVRs, and tau accumulation epicenters as the top 5% of ROIs with the highest rates of percentage change in tau-PET SUVR, both at the subject level. Tau (accumulation) epicenter connectivity was determined by taking the functional connectivity-based distance of each non-(accumulation-)epicenter ROI ($n=190$) to the (accumulation) epicenter ($n=10$). For each individual, linear regression was used to assess the association between functional connectivity-based distance to the tau (accumulation) epicenter and tau-PET SUVR/annual percentage change. Subject-level β -values are visualized per AD variant in the notched boxplots. Additionally, all non-(accumulation-)epicenter regions were grouped into quartiles based on their functional proximity to the (accumulation) epicenter (quartile 1 = close, quartile 4 = distant), and tau positivity probabilities/tau-PET SUVR percentage change rates across quartiles were significantly different by paired Wilcoxon signed-rank tests. AD = Alzheimer's disease; IvPPA = logopenic variant primary progressive aphasia; PCA = posterior cortical atrophy; PET = positron emission tomography; Q = quartile; ROI = region of interest; SUVR = standardized uptake value ratio.