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# <sup>18</sup>F-flortaucipir PET to autopsy comparisons in Alzheimer's disease and other neurodegenerative diseases

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Few studies have evaluated the relationship between *in vivo* <sup>18</sup>F-flortaucipir PET and post-mortem pathology. We sought to compare antemortem <sup>18</sup>F-flortaucipir PET to neuropathology in a consecutive series of patients with a broad spectrum of neurodegenerative conditions. Twenty patients were included [mean age at PET 61 years (range 34–76); eight female; median PET-to-autopsy interval of 30 months (range 4–59 months)]. Eight patients had primary Alzheimer's disease pathology, nine had non-Alzheimer tauopathies (progressive supranuclear palsy, corticobasal degeneration, argyrophilic grain disease, and frontotemporal lobar degeneration with *MAPT* mutations), and three had non-tau frontotemporal lobar degeneration. Using an inferior cerebellar grey matter reference, 80–100-min <sup>18</sup>F-flortaucipir PET standardized uptake value ratio (SUVR) images were created. Mean SUVRs were calculated for progressive supranuclear palsy, corticobasal degeneration, and neurofibrillary tangle Braak stage regions of interest, and these values were compared to SUVRs derived from young, non-autopsy, cognitively normal controls used as a standard for tau negativity. W-score maps were generated to highlight areas of increased tracer retention compared to cognitively normal controls, adjusting for age as a covariate. Autopsies were performed blinded to PET results. There was excellent correspondence between areas of <sup>18</sup>F-flortaucipir retention, on both SUVR images and W-score maps, and neurofibrillary tangle distribution in patients with primary Alzheimer's disease neuropathology. Patients with non-Alzheimer tauopathies and non-tau frontotemporal lobar degeneration showed a range of tracer retention that was less than Alzheimer's disease, though higher than age-matched, cognitively normal controls. Overall, binding across both tau-positive and tau-negative non-Alzheimer disorders did not reliably correspond with post-mortem tau pathology. <sup>18</sup>F-flortaucipir SUVRs in subcortical regions were higher in autopsy-confirmed progressive supranuclear palsy and corticobasal degeneration than in controls, but were similar to values measured in Alzheimer's disease and tau-negative neurodegenerative pathologies. Quantification of <sup>18</sup>F-flortaucipir SUVR images at Braak stage regions of interest reliably detected advanced Alzheimer's (Braak VI) pathology. However, patients with earlier Braak stages (Braak I–IV) did not show elevated tracer uptake in these regions compared to young, tau-negative controls. In summary, PET-to-autopsy comparisons confirm that <sup>18</sup>F-flortaucipir PET is a reliable biomarker of advanced Braak tau pathology in Alzheimer's disease. The tracer cannot reliably differentiate non-Alzheimer tauopathies and may not detect early Braak stages of neurofibrillary tangle pathology.

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**Abbreviations:** ADNC = Alzheimer's disease neuropathological change; CBD = corticobasal degeneration; FTLN = frontotemporal lobar degeneration; FTP = <sup>18</sup>F-flortaucipir; NFT = neurofibrillary tangle; PSP = progressive supranuclear palsy; PVC = partial volume correction; SUVR = standardized uptake value ratio; TDP-43 = TAR DNA binding protein 43

## Introduction

Aggregation of hyper-phosphorylated tau characterizes various neurodegenerative diseases, collectively known as tauopathies (Lee *et al.*, 2001; Gibbons *et al.*, 2019). Alzheimer's disease is the most common tauopathy, and neurofibrillary tangles (NFTs) consisting of paired helical filaments and straight filaments of hyperphosphorylated tau are one of the pathologic hallmarks of this disease (Fitzpatrick *et al.*, 2017). Other tauopathies show disease-specific tau deposits, with distinct neuronal and glial vulnerabilities, tau ultrastructural conformations, and regional distributions in the brain. Tauopathies can be classified by the number of repeats of the tau microtubule-binding domain [i.e. three-repeat (3R), four-repeat (4R), or mixed (3R/4R)] and ultrastructural conformation (Fitzpatrick *et al.*, 2017; Falcon *et al.*, 2018, 2019; Goedert *et al.*, 2018; Scheres *et al.*, 2020; Zhang *et al.*, 2020). Alzheimer's disease inclusions contain 3R/4R tau isoforms in paired helical filaments. Inclusions in progressive supranuclear palsy (PSP), corticobasal degeneration (CBD), and argyrophilic grain disease primarily contain 4R tau isoforms in straight filaments (Komori, 1999; Takahashi *et al.*, 2002; Zhukareva *et al.*, 2002; Arima, 2006). Pick's disease inclusions consist of predominantly 3R tau isoforms in straight and twisted filaments (Murayama *et al.*, 1990; Komori, 1999; King *et al.*, 2001; Arima, 2006).

PET imaging tracers have been developed to enable *in vivo* imaging of pathological tau. One of the most widely used tau PET tracers, <sup>18</sup>F-flortaucipir (FTP) (formerly <sup>18</sup>F-AV1451 and <sup>18</sup>F-T807), has been shown by autoradiography to bind selectively to paired helical filament tau versus amyloid- $\beta$ ,  $\alpha$ -synuclein, and TAR DNA-binding protein 43 (TDP-43) deposits in post-mortem human brain tissue (Xia *et al.*, 2013; Marquié *et al.*, 2015; Lowe *et al.*, 2016; Sander *et al.*, 2016). FTP-PET accurately differentiates clinically diagnosed Alzheimer's disease from healthy controls and other neurodegenerative diseases (Chien *et al.*, 2013; Cho *et al.*, 2016; Ossenkoppele *et al.*, 2018). In non-Alzheimer tauopathies, *in vitro* autoradiography studies demonstrate absent-to-low binding in non-Alzheimer tauopathies

consisting of straight tau filaments (Marquié *et al.*, 2015, 2017a; Lowe *et al.*, 2016; Sander *et al.*, 2016; Ono *et al.*, 2017). *In vivo*, FTP shows tracer uptake in regions expected to contain tau pathology, which is less intense than in Alzheimer's disease and partly overlaps with areas of off-target (i.e. non-tau-related) binding (Cho *et al.*, 2017a, b; Coakeley *et al.*, 2017; Passamonti *et al.*, 2017; Schonhaut *et al.*, 2017; Smith *et al.*, 2017a, b; Jones *et al.*, 2018; Tsai *et al.*, 2019). Furthermore, low-level binding is also seen in patients expected to harbour tau-negative, TDP-43 pathology based on clinical phenotype or presence of a disease-causing mutation (Bevan-Jones *et al.*, 2018a, b; Makaretz *et al.*, 2018; Smith *et al.*, 2019a; Tsai *et al.*, 2019). Further FTP-to-autopsy correlation is needed in Alzheimer's disease, non-Alzheimer tauopathies, and non-tau frontotemporal lobar degeneration (FTLD).

*In vivo* FTP retention patterns match the topography of Braak NFT staging, suggesting that FTP could be used as a surrogate marker of tau burden and severity in Alzheimer's disease (Schöll *et al.*, 2016; Schwarz *et al.*, 2016; Marquié *et al.*, 2017b). Recently, Smith *et al.* (2019b) showed strong correlation between FTP retention and post-mortem tau pathology in a patient with Alzheimer's disease caused by presenilin 1 (*PSEN1*) mutation. Lowe *et al.* (2019) showed that patients with primary pathological diagnoses of Alzheimer's disease, who had Braak NFT stages IV–VI, exhibited elevated FTP retention in a temporal composite meta-region of interest, and found strong correlation between FTP binding and quantitative phosphorylated tau immunohistochemistry at autopsy. Additionally, a larger autopsy study, using a binary visual classification scheme to interpret antemortem FTP scans, found that positively read scans accurately identified patients with advanced tau (Braak NFT stages V–VI) and Alzheimer's disease [high Alzheimer's disease neuropathological change (ADNC)] pathology (Fleisher *et al.*, 2020).

Our primary aims in this study were to: (i) compare FTP-PET to pathology in patients with a primary autopsy diagnosis of Alzheimer's disease; (ii) compare FTP-PET to pathology in patients with a primary autopsy diagnosis of

non-Alzheimer tauopathy or tau-negative FTLN; and (iii) evaluate whether FTP-PET detects early-stage (Braak I–IV) neurofibrillary changes of Alzheimer’s disease.

## Materials and methods

### Participants

Twenty consecutive participants enrolled in research at the University of California, San Francisco (UCSF) Memory and Aging Center underwent antemortem FTP-PET and brain autopsy (Table 1). One patient underwent FTP-PET as part of a clinical trial sponsored by Avid Radiopharmaceuticals (Flortaucipir PET Imaging in Subjects with Frontotemporal Dementia, ClinicalTrials.gov Identifier: NCT03040713), while all other patients were enrolled in UCSF-specific protocols. Three patients had FTP-PET and autopsy results described in previous clinical studies (Schonhaut *et al.*, 2017; Tsai *et al.*, 2019). All participants underwent antemortem history, neurological exam, neuropsychological testing, genetic testing, brain MRI, and FTP-PET. Clinical diagnoses were made by consensus application of standard research criteria (Albert *et al.*, 2011; Gorno-Tempini *et al.*, 2011; McKhann *et al.*, 2011; Rascovsky *et al.*, 2011; Armstrong *et al.*, 2013; Höglinger *et al.*, 2017; McKeith *et al.*, 2017). Seventeen participants underwent antemortem <sup>11</sup>C-Pittsburgh compound-B (PiB) PET and one participant underwent <sup>18</sup>F-florbetapir PET for *in vivo* determination of brain amyloid- $\beta$  status. Written informed consent was obtained from participants or their designated surrogate decision-makers. Institutional review boards at UCSF, University of California, Berkeley, and Lawrence Berkeley National Laboratory (LBNL) approved the study.

### Genetic assessment

All participants received targeted sequencing of the most common causative genes for Alzheimer’s disease and FTLN, including *PSEN1* and *PSEN2*, amyloid precursor protein (*APP*), microtubule-associated protein tau (*MAPT*), chromosome 9 open reading frame 72 (*C9orf72*), transactive response DNA binding protein 43 kDa (*TARDBP*), and progranulin (*GRN*) (Kim *et al.*, 2018).

### MRI acquisition and preprocessing

T<sub>1</sub>-weighted magnetization prepared rapid gradient echo (MPRAGE) MRI sequences were acquired at UCSF, either on a 3 T Siemens Tim Trio ( $n = 15$ ) or a 3 T Siemens Prisma Fit ( $n = 5$ ) scanner. Detailed MRI acquisition parameters are described in the Supplementary material.

MRIs were segmented and parcellated using Freesurfer 5.3 (<https://surfer.nmr.mgh.harvard>). Statistical Parametric Mapping (SPM12; Wellcome Trust Center for Neuroimaging, London, UK, <https://www.fil.ion.ucl.ac.uk/spm>) was used to process PET data as described below.

### PET acquisition and preprocessing

FTP-PET images were acquired on a Siemens Biograph PET/CT scanner at LBNL ( $n = 19$ ) or a GE Discovery STE/VCT PET/CT

scanner at the UCSF Department of Radiology and Biomedical Imaging at China Basin ( $n = 1$ ). We analysed PET data acquired 80–100 min (four 5-min frames) after injection of  $\sim 10$  mCi of FTP. A low-dose CT scan was performed for attenuation correction, and data were reconstructed using an ordered subset expectation maximization algorithm with weighted attenuation and smoothed with a 4 mm Gaussian kernel with scatter correction.

PET frames were realigned, averaged, and coregistered onto their corresponding MRI. Standardized uptake value ratio (SUVR) images were created in native space using MRI-defined inferior cerebellum grey matter as a reference region (Maass *et al.*, 2017).

Eighteen patients also underwent amyloid PET with <sup>11</sup>C-PiB ( $n = 17$ ) or <sup>18</sup>F-florbetapir ( $n = 1$ ). Acquisition, preprocessing, and analysis of FTP and amyloid PET images are described in detail in the Supplementary material.

### Neuropathology

All participants underwent a standardized post-mortem assessment at the UCSF Neurodegenerative Disease Brain Bank (Schonhaut *et al.*, 2017; Kim *et al.*, 2018). Pathological assessment was performed blinded to FTP-PET. The fresh brains were fixed, and tissue blocks were obtained from neurodegenerative disease-related neuroanatomical regions of interest. Microvacuolation, astrogliosis, and neuronal loss were scored on haematoxylin and eosin stained sections. Immunohistochemistry was performed using antibodies against hyperphosphorylated tau (CP-13, S202, mouse, 1:250, courtesy of Dr P. Davies), amyloid- $\beta$  (1–16, clone DE2, mouse, 1:500, Millipore),  $\alpha$ -synuclein (LB509, mouse, 1:5000, courtesy of Drs J. Trojanowski and V. Lee), and TDP-43 (rabbit, 1:4000, Proteintech Group).

Neuropathological diagnosis followed currently accepted guidelines (Litvan *et al.*, 1996; McKeith *et al.*, 1996; Dickson *et al.*, 2002; Ferrer *et al.*, 2008; MacKenzie *et al.*, 2010; Montine *et al.*, 2012). Pathologies were designated as primary, contributing, or incidental, based on the neuropathologist’s interpretation of their contribution to the patient’s clinical syndrome. ADNC was rated according to the latest ‘ABC’ score criteria, which includes Braak staging for neurofibrillary changes (Braak and Braak, 1991; Montine *et al.*, 2012). Typically, intermediate-to-high ADNC was deemed a primary or contributing pathology associated with some component of the clinical syndrome, whereas low ADNC was always considered incidental in this cohort.

### Experimental design and statistical analyses

#### FTP-PET W-score maps

To obtain maps of abnormally elevated FTP-PET at the individual patient level (i.e. distinguish tracer binding beyond background level), we computed voxelwise FTP-PET W-score maps (Z-score maps adjusted for age) from SUVR images, which are also referred to as W-maps (O’Brien and Dyck, 1995; Jack *et al.*, 1997; La Joie *et al.*, 2012). See the Supplementary material for additional information on W-map analysis.

**Table 1 Patient characteristics**

Autopsy diagnosis category	ID	Autopsy diagnosis (primary, contributing)	Clinical diagnosis	Gender	Age at PET	Time: PET-to-autopsy, months	MMSE at PET	Amyloid PET	Amyloid quant. (centiloid SUVR)	Pathology: Braak stage; ABC score; ADNC level
Alzheimer's disease	A1	AD, LBD	AD (logopenic PPA)	Female	53	36	16	n/a	n/a	V1; A3, B3, C3; High
	A2	AD, LBD	AD/PD	Male	71	8	22	Positive	109	V1; A3, B3, C3; High
	A3	AD, LBD	AD (logopenic PPA)	Male	73	38	22	Positive	115	V1; A3, B3, C3; High
	A4	AD	AD	Male	68	31	8	Positive	106	V1; A3, B3, C3; High
	A5	AD	AD (logopenic PPA)	Female	60	59	27	Positive	89	V1; A3, B3, C3; High
	A6	AD	AD (PCA)	Female	59	40	9	Positive	124	V1; A3, B3, C3; High
	A7	AD	AD (logopenic PPA)	Female	63	25	28	Positive	172	V1; A3, B3, C3; High
	A8	AD	AD (PCA)	Male	53	44	26	Positive	96	V1; A3, B3, C3; High
Non-Alzheimer tauopathies	1	PSP	MCI (mixed)	Male	60	12	26	Negative	0	II; A1, B1, C1; Low
	2	PSP	Non-fluent PPA	Female	74	29	27	Negative	-6	II; A1, B1, C0; Low
	3	PSP	PSP	Male	76	50	24	Negative	18	II; A1, B1, C0; Low
	4	PSP, AD	PSP	Male	72	49	26	Positive	69	III; A3, B2, C2; Intermediate
	5	CBD	PSP	Female	63	9	16	Negative	-1	I; A0, B1, C0; Not ADNC
	6	CBD, HS (left), TDP-43 unclassifiable	bvFTD	Male	46	43	22	Negative	-4	III; A0, B2, C0; Not ADNC
	7	FTLD-tau (MAPT P301L), AD	FTLD (possible TES)	Male	67	36	5	Positive	69	IV; A3, B2, C3; Intermediate
	8	FTLD-tau (MAPT S301I), HS (right)	bvFTD	Female	37	26	4	Negative	12	0; A0, B0, C0; Not ADNC
	9	AGD, vascular	bvFTD	Male	68	4	29	Negative	-4	III; A0, B2, C0; Not ADNC
	10	TDP-43 type A (GRN NM_002087:c.708+6_708+9del)	bvFTD	Male	68	18	21	Negative	-11	I; A1, B1, C0; Low
Non-tau FTLD	11	TDP-43 type B (C9orf72)	bvFTD	Female	48	20	26	Negative	-6	I; A1, B1, C0; Low
	12	FTLD-FUS with MND	bvFTD	Male	34	12	22	n/a	n/a	0; A0, B0, C0; Not ADNC
Summary statistics			Male: 12 Female: 8	61	30	20				

ABC = Amyloid Braak CERAD; AD = Alzheimer's disease; AGD = argyrophilic grain disease; bvFTD = behavioural variant frontotemporal dementia; HS = hippocampal sclerosis; LBD = Lewy body disease; MCI = mild cognitive impairment; MMSE = Mini-Mental State Examination; MND = motor neuron disease; n/a = not applicable (these patients did not undergo antemortem amyloid PET); PCA = posterior cortical atrophy, PD = Parkinson's disease; PPA = primary progressive aphasia; TES = traumatic encephalopathy syndrome.



### FTP-PET region of interest analyses

Using Freesurfer segmentation, based on the Desikan atlas (Desikan *et al.*, 2006), the average cortical SUVR value, weighted by subregion volume, was extracted from each patient in a set of *a priori* regions of interest corresponding to Braak NFT stages (Maass *et al.*, 2017). Entorhinal cortex (Braak I) was used instead of a Braak III region of interest, because exclusion of the hippocampus was necessary to reduce signal spill-in due to off-target FTP retention in the adjacent choroid plexus (Marquie *et al.*, 2015, 2017c; Schöll *et al.*, 2016; Lowe *et al.*, 2016; Lemoine *et al.*, 2018; Baker *et al.*, 2019). The Braak III/IV and V/VI regions of interest have been previously described (Maass *et al.*, 2017).

SUVRs were also extracted from each patient at seven regions of interest (precentral gyrus, postcentral gyrus, putamen, globus pallidus, subthalamic nucleus, substantia nigra, dentate nucleus) corresponding to neuroanatomical regions that often have elevated tau burden in PSP and CBD (Hauw *et al.*, 1994; Dickson *et al.*, 2002). The tools used to define these cortical and subcortical regions of interest are described in the [Supplementary material](#).

As a standard for tau-negativity, mean and standard deviation (SD) of SUVR levels were calculated at each region of interest for 14 young (mean age 26, SD 5 years), non-autopsy, cognitively normal controls who underwent FTP-PET on the same scanner as the patients. Using these data, a mean + 2 SD SUVR threshold was created for each region of interest. Any FTP-PET SUVR above the threshold was considered potentially distinguishable from controls (i.e. higher than noise and off-target binding seen in tau-negative controls) at the region of interest. The FTP-PET entorhinal cortex and Braak stage region of interest data also underwent partial volume correction (PVC) based on the geometric transfer matrix approach tailored for FTP-PET (Rousset *et al.*, 1998; Baker *et al.*, 2017, 2019). The other regions of interest underwent PVC using a method combining geometric transfer matrix and geometric transfer matrix-derived region-based voxel-wise approaches to enhance accuracy given the inclusion of small subcortical regions of interest (Thomas *et al.*, 2011). All analyses were repeated with PVC data.

### Data availability

All data used in this study are available for review upon request.

## Results

### Participants and neuropathological diagnoses

**Table 1** summarizes demographic, clinical, and pathological characteristics. The patients were predominantly male ( $n = 12$ ), relatively young (mean age 61, SD 12 years), and most had dementia at the time of FTP-PET imaging. The time gap between FTP-PET and autopsy ranged from 4 to 59 months, with a median of 30 months. All patients with clinical syndromes that predict Alzheimer's disease, including four patients with logopenic variant primary progressive aphasia and two with posterior cortical atrophy, had high ADNC

(A3, B3, C3; Braak VI) pathology at autopsy. Patients who met clinical criteria for PSP had pathological diagnoses of PSP or CBD at autopsy. Patients who met clinical diagnostic criteria for behavioural variant frontotemporal dementia or non-fluent primary progressive aphasia, had either tau (PSP, CBD, argyrophilic grain disease, FTLD-tau due to *MAPT* mutation), TDP-43, or FUS pathology at autopsy.

### Comparison of FTP retention to pathology in patients with a primary neuropathological diagnosis of Alzheimer's disease

On visual inspection of SUVR images and W-maps, all patients with primary Alzheimer's disease autopsy diagnosis had intense tracer retention in the parietal lobes, especially the precuneus, and posterior cingulate (**Fig. 1** and **Supplementary Fig. 1**). Three of four patients with logopenic variant primary progressive aphasia had asymmetric, left greater than right hemisphere tracer retention (**Supplementary Fig. 1**). Both patients with posterior cortical atrophy had parieto-occipital tracer retention (**Fig. 1** and **Supplementary Fig. 1**). The location of tracer retention mirrored the clinical and neuroanatomical variability of the different Alzheimer's disease phenotypes.

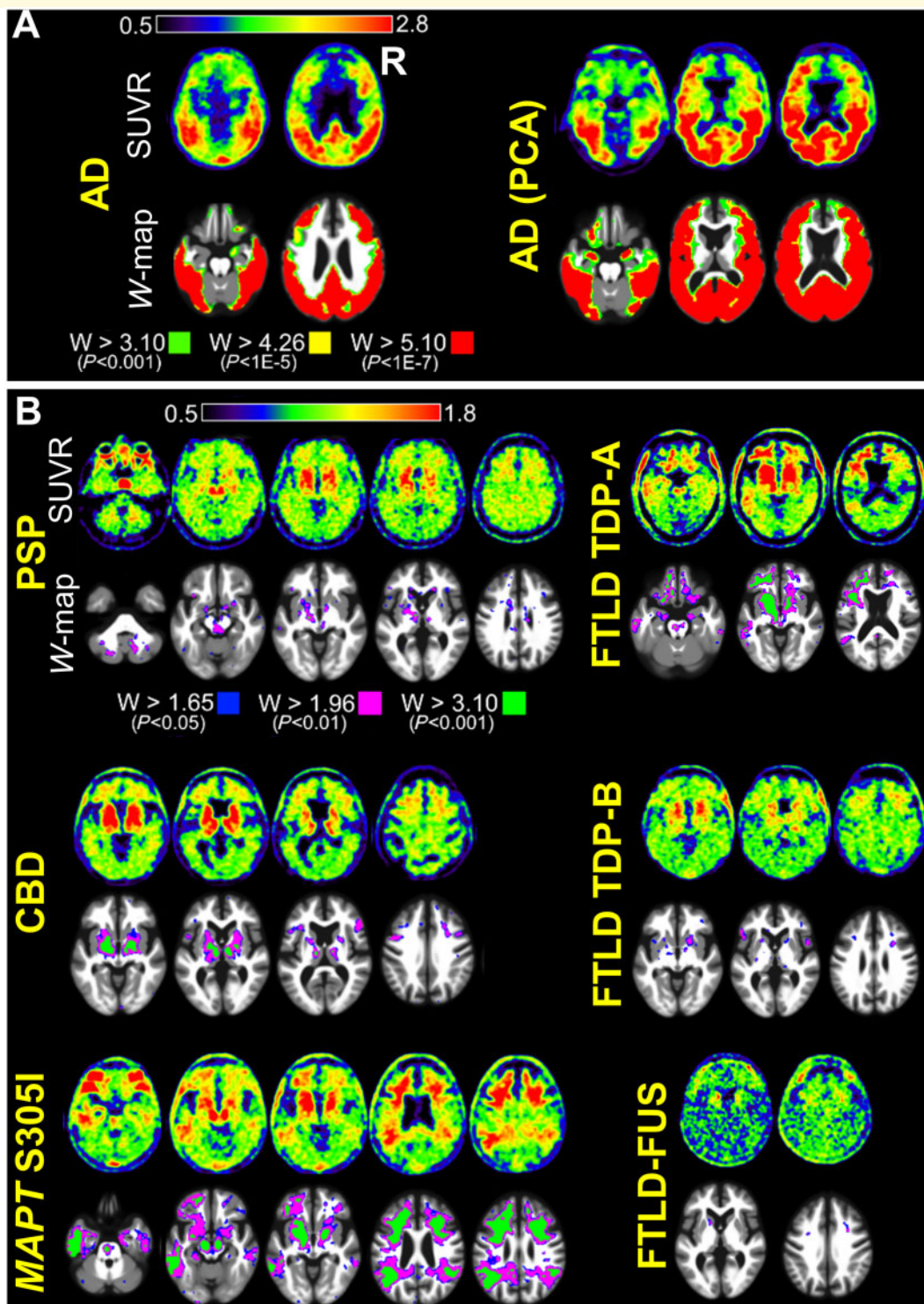
All patients with a primary neuropathological diagnosis of Alzheimer's disease had areas of tracer retention that corresponded to the distribution and severity of NFT pathology in both typical and atypical presentations of Alzheimer's disease (**Fig. 2**, **Supplementary Fig. 1** and **Supplementary Table 1**). For example, compared to typical amnesic Alzheimer's disease, the patients with posterior cortical atrophy had more NFT pathology in the calcarine cortex and less NFT pathology in the hippocampus (**Fig. 2**).

Compared to patients with other neuropathological diagnoses at autopsy, Alzheimer's disease patients had more intense tracer binding throughout the cortex, which necessitated the use of higher upper thresholds to visualize the full range of tracer binding (**Fig. 1** and **Supplementary Fig. 1**). Altogether, the distribution and intensity of tracer uptake in patients with a primary Alzheimer's disease autopsy diagnosis allowed for qualitative differentiation of these patients from those with other primary neurodegenerative pathologies at the single patient level on both SUVR images and W-maps (**Fig. 1** and **Supplementary Figs 1 and 2**).

### Comparison of FTP retention to pathology in patients with non-Alzheimer primary neuropathological diagnoses

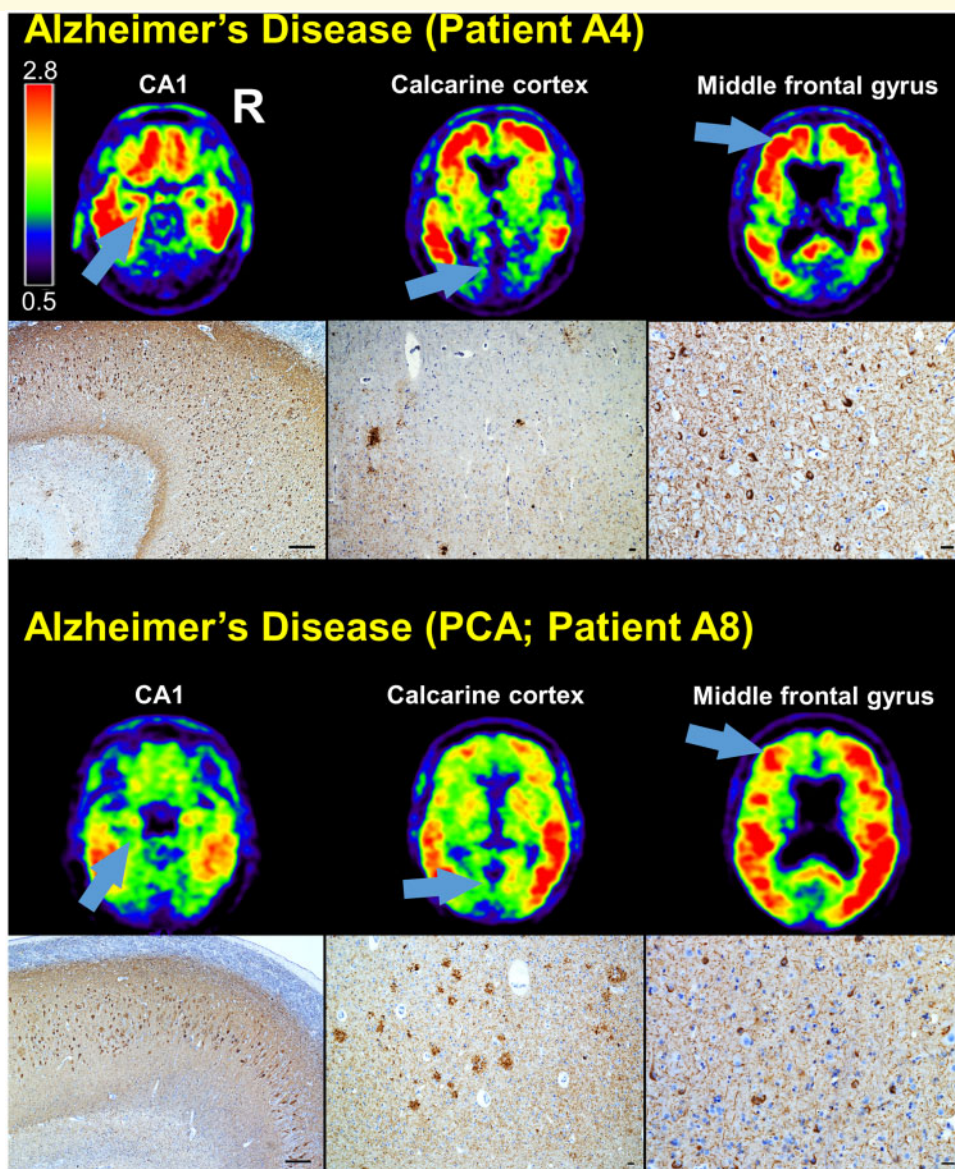
#### Progressive supranuclear palsy

All four patients with a primary neuropathological diagnosis of PSP showed tracer uptake in basal ganglia (particularly



**Figure 1** FTP-PET SUVR and W-score map images. **(A)** Example SUVR and W-map images are shown for patients with primary Alzheimer's disease (AD) neuropathological diagnosis. One patient had a typical amnesic clinical phenotype and the other patient had an atypical posterior cortical atrophy (PCA) phenotype. Patients with primary Alzheimer's disease pathology had marked FTP uptake, and high SUVR and W-map thresholds were chosen to highlight the full range of tracer retention in these patients. Images for all Alzheimer's disease patients are shown in [Supplementary Fig. 1](#). **(B)** Example SUVR and W-map images are shown for patients with primary autopsy diagnoses of non-Alzheimer tauopathies, FTLD-TDP, and FTLD-FUS. These patients had lower FTP uptake than patients with primary Alzheimer's disease autopsy diagnosis, and lower SUVR and W-map thresholds needed to be used to highlight the areas of tracer retention in these patients. To allow for comparison with the Alzheimer's disease cases, the W-map upper threshold ( $W > 3.10$ ) is the same as the lower threshold used in patients with primary Alzheimer's disease pathology. Images for all patients with non-Alzheimer's autopsy diagnoses are shown in [Supplementary Fig. 2](#).





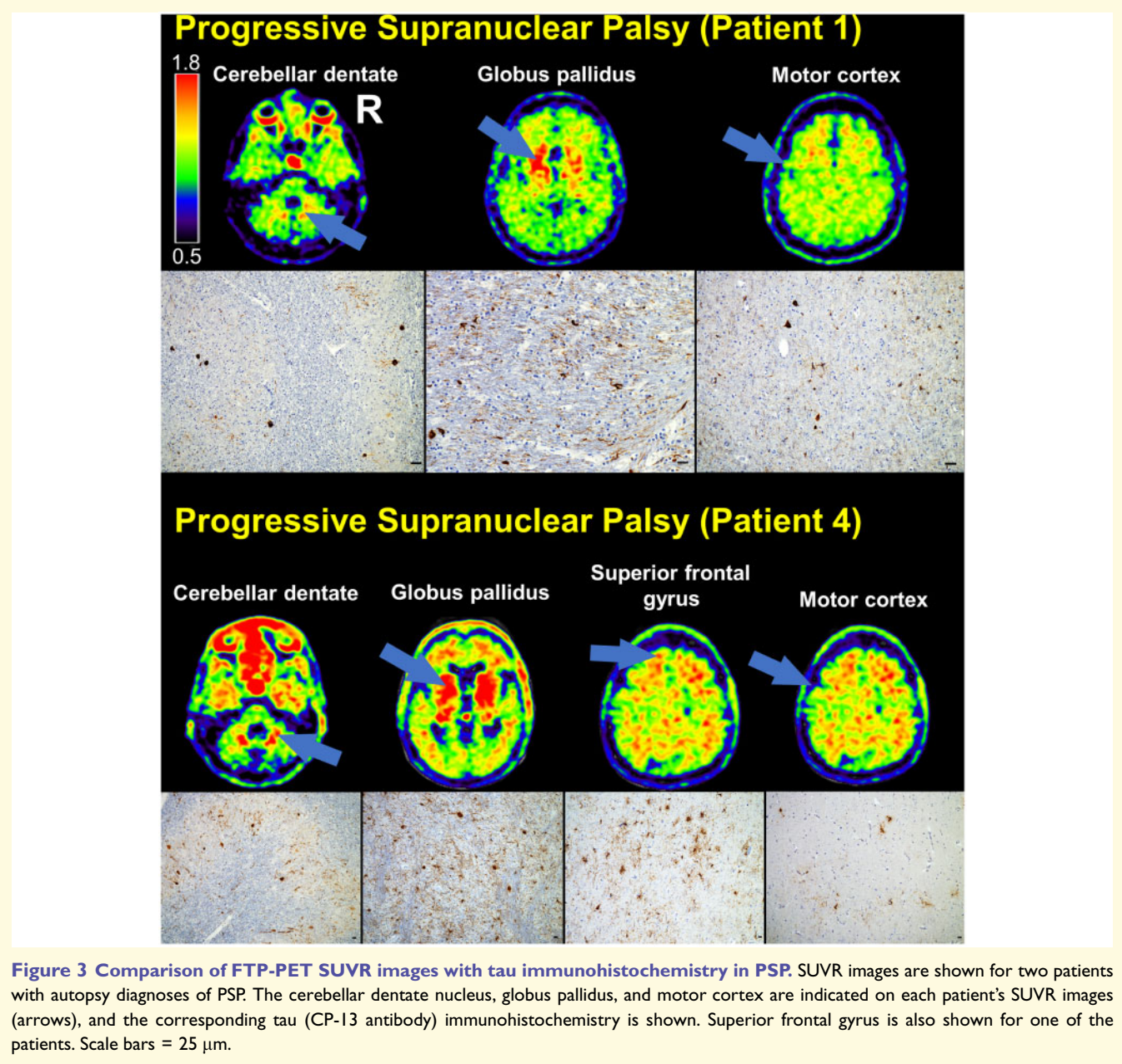
**Figure 2** Comparison of FTP-PET SUVR images to tau immunohistochemistry in Alzheimer's disease. SUVR images are shown for two patients with autopsy diagnoses of Alzheimer's disease. Both patients were assigned antemortem clinical diagnoses of Alzheimer's disease, but one patient had a typical amnesic phenotype and the other patient had an atypical posterior cortical atrophy (PCA) phenotype. CA1, calcarine cortex, and middle frontal gyrus are indicated on each patient's SUVR images (arrows), and the corresponding tau (CP-13 antibody) immunohistochemistry is shown. Scale bars = 250  $\mu$ m in the CA1 micrographs and 25  $\mu$ m in the other micrographs.

the globus pallidus), frontal white matter, and midbrain. Three patients showed significant tracer uptake in the cerebellar dentate nucleus (Fig. 1 and Supplementary Fig. 2). Though many of these regions overlap with areas of off-target FTP binding, W-maps confirmed the intensity of binding was greater than expected for age based on normative data from amyloid-negative controls. The intensity of binding was lower than in Alzheimer's disease.

At autopsy, all patients showed midbrain pathological changes typical of PSP, including tufted astrocytes, globose tangles, and oligodendroglial coiled bodies. The globus pallidus was examined in all four patients, and three patients

had tufted astrocytes and two patients had coiled bodies in this region. The cerebellar dentate nucleus was examined in three patients, and all had globose tangles and glial coiled bodies (Fig. 3 and Supplementary Tables 10–13). Comparing two patients (Patients 1 and 4) who had tau pathology in the globus pallidus and cerebellar dentate, the patient with the higher burden of tau pathology in these regions (Patient 4) had higher tracer retention in these regions (Fig. 3). Both patients had low FTP retention in the motor cortex, and there was less tau pathology in this region compared to globus pallidus and cerebellar dentate. Patient 2 had extensive PSP tau pathology in the inferior frontal





**Figure 3** Comparison of FTP-PET SUVR images with tau immunohistochemistry in PSP. SUVR images are shown for two patients with autopsy diagnoses of PSP. The cerebellar dentate nucleus, globus pallidus, and motor cortex are indicated on each patient's SUVR images (arrows), and the corresponding tau (CP-13 antibody) immunohistochemistry is shown. Superior frontal gyrus is also shown for one of the patients. Scale bars = 25  $\mu$ m.

gyrus, primary motor cortex, and brainstem, with relative sparing of the globus pallidus and dentate nucleus (the areas with lower tau pathology lacked FTP retention; [Supplementary Fig. 2](#)). Affected regions showed significant astrogliosis. Tracer retention in the frontal white matter did not correspond to tau pathology on stained tissue sections.

Temporal and limbic neuroanatomical structures contained moderate levels of PSP tau pathology and also NFTs. Patient 4 had contributing Alzheimer's disease pathology, with intermediate ADNC and Braak stage III, and the other patients had incidental Braak stage II. Despite the presence of both PSP tau pathology and NFTs of Alzheimer-type, there was no discernible tracer retention in the entorhinal

cortex, hippocampus, or amygdala on either SUVR images or W-maps. Most of the temporal tracer retention localized to the white matter ([Supplementary Fig. 2](#)).

#### Corticobasal degeneration

The two patients with CBD at autopsy had asymmetric tracer retention, more prominent in one hemisphere, mostly involving the perirolandic cortex, inferior frontal gyrus, frontal white matter, basal ganglia (especially the globus pallidus), thalamus, substantia nigra, and brainstem ([Fig. 1](#) and [Supplementary Fig. 2](#)). On inspection of SUVR images and W-maps, these patients had lower tracer retention than patients with primary Alzheimer's disease pathology, but

they generally had more tracer retention than other sporadic non-Alzheimer tauopathies, FTLTDP, or FTLTDP-FUS.

The areas of highest tracer retention (perirolandic cortex, frontal white matter, basal ganglia, substantia nigra, and brainstem) corresponded to regions of high CBD tau pathology burden (astrocytic plaques, coiled bodies, and white matter threads; [Supplementary Tables 14 and 15](#)). Patient 5 had incidental Braak I NFT pathology, but there was no entorhinal cortex tracer retention on the SUVR image or W-map. This patient had elevated FTP-PET signal in the temporal white matter ([Fig. 1](#) and [Supplementary Fig. 2](#)), which corresponded to underlying CBD tau pathology. Patient 6 had incidental Braak III NFT pathology and left hippocampal sclerosis due to TDP-43 pathology. This patient showed FTP tracer retention in left greater than right mesial temporal lobes and temporal white matter on the SUVR image, which was also apparent on the W-map ([Supplementary Fig. 2](#)). The mesial temporal lobes were atrophied and contained CBD tau pathology, NFTs, and TDP-43 pathology, making it difficult to attribute the elevated temporal FTP retention to a specific pathology.

### Argyrophilic grain disease

The patient with primary argyrophilic grain disease neuropathology showed tracer retention limited to basal ganglia, substantia nigra, and right temporal lobe (right inferior temporal gyrus, hippocampus) on the SUVR image, which was not significant on the W-map compared to age-matched controls ([Supplementary Fig. 2](#)). Tau-positive pretangles, neurofibrillary thread, and grains were found in the entorhinal cortex, CA1/subiculum, CA2, and amygdala, with sparse pre-tangles present in the inferior temporal gyrus ([Supplementary Table 18](#)). Incidental Braak III NFT pathology was also present. Very little tau pathology was observed in the basal ganglia, substantia nigra, or other neocortical regions assessed at autopsy.

### FTLTDP-tau pathology due to MAPT mutations

Both patients with FTLTDP-tau pathology due to *MAPT* mutations exhibited increased tracer retention in the temporal lobes, temporal white matter, and basal ganglia ([Fig. 1](#) and [Supplementary Fig. 2](#)). The patient carrying a P301L mutation was PiB-positive and showed elevated FTP-PET signal in right greater than left temporal and occipital cortex, as well as small clusters of elevated retention in posterior cingulate and lateral parietal cortex. At autopsy, the patient had intermediate ADNC (A3, B2, C3; Braak IV NFT pathology) and extensive FTLTDP-tau pathology (neuronal cytoplasmic inclusions, bushy astrocytes) throughout the hippocampus, amygdala, cortex, and basal ganglia ([Supplementary Tables 1 and 16](#)).

The patient carrying an S305I mutation showed marked tracer retention in the frontal and temporal white matter. At autopsy, FTLTDP-tau pathology was abundant in cortical and subcortical regions with features reminiscent of diffuse neocortical argyrophilic grain disease with granular

tau-positive neuronal cytoplasmic inclusions, often with a perinuclear ring-like pattern, coiled oligodendroglial cytoplasmic inclusions, threads, and grains. There was extensive thread and grain pathology in the subcortical white matter ([Fig. 4](#) and [Supplementary Table 17](#)). The frontal and temporal tracer retention was localized in areas in which there was considerable FTLTDP-tau pathology ([Fig. 4](#)). There was no Alzheimer-type NFT pathology (Braak 0). Hippocampal sclerosis was observed in the absence of TDP-43 inclusions.

### FTLTDP TDP-43 type A due to GRN mutation

The patient with FTLTDP TDP-43 type A due to *GRN* (NM\_002087:c.708+6\_708+9del) mutation exhibited a moderate level of tracer retention throughout the frontal cortex (superior frontal gyrus, middle frontal gyrus, orbitofrontal cortex, and frontal pole), frontal white matter, basal ganglia (especially, globus pallidus and putamen), lateral temporal cortex, and temporal poles ([Fig. 1](#) and [Supplementary Fig. 2](#)). At autopsy, tau NFTs were limited to the entorhinal cortex (Braak I). Notably, tau immunohistochemistry was negative in the sampled cortical regions, which had high FTP signal ([Fig. 5](#), [Supplementary Tables 1 and 19](#)). Extensive TDP-43 immunoreactive neuronal cytoplasmic inclusions and dystrophic neurites were present throughout the frontal lobes, temporal lobes, and basal ganglia.

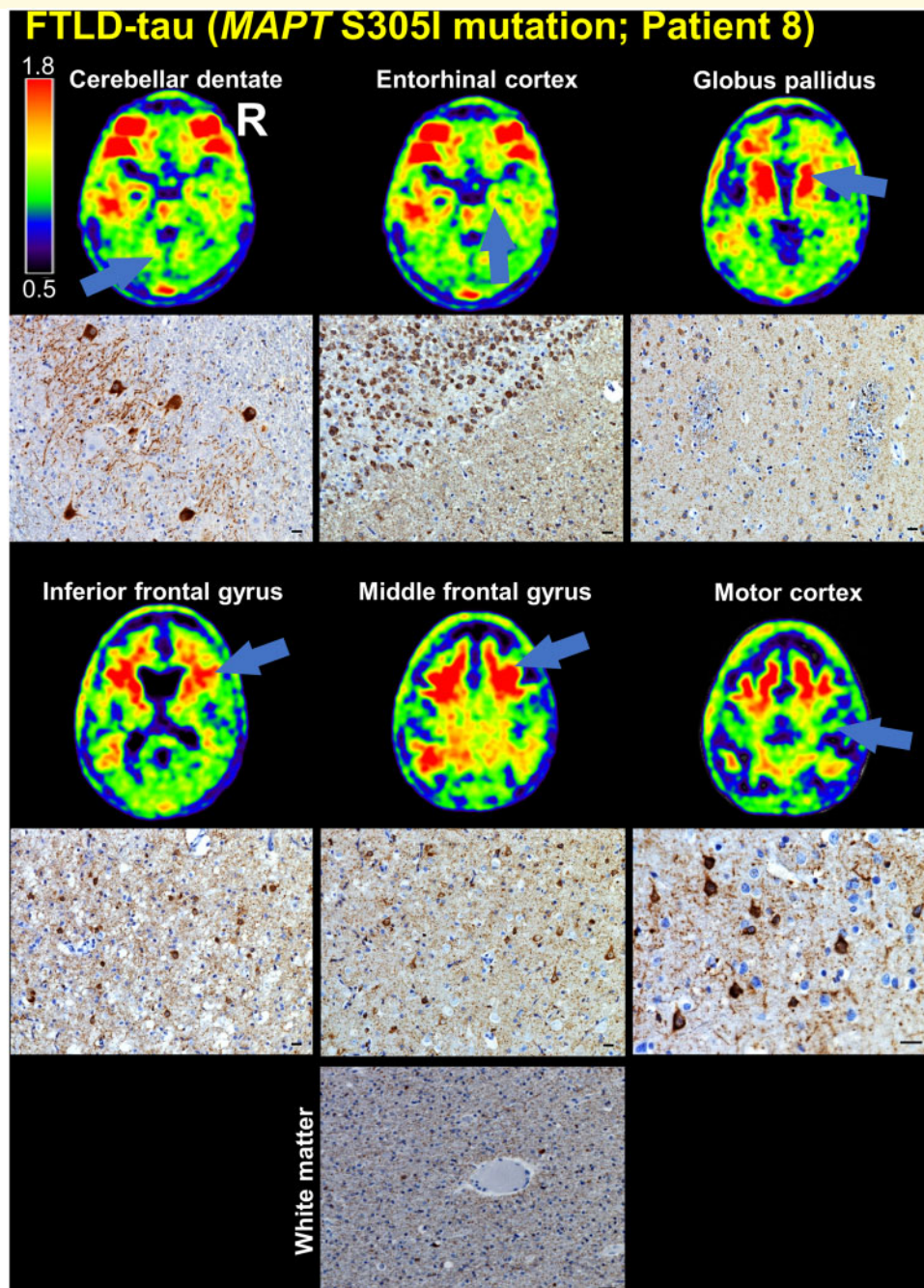
### FTLTDP TDP-43 type B due to C9orf72 expansion

The patient with FTLTDP TDP-43 type B due to a *C9orf72* expansion exhibited small clusters of elevated tracer uptake in the frontal white matter, striatum, thalamus, substantia nigra, and right half of the pons ([Fig. 1](#) and [Supplementary Fig. 2](#)). At autopsy, there was a small amount of tau pathology in the middle frontal gyrus, and extensive TDP-43 inclusions throughout the frontal lobes ([Supplementary Table 20](#)). The frontal tau pathology did not correspond to FTP-PET signal in the frontal lobes. The striatum, thalamus, and substantia nigra only contained TDP-43 inclusions. The right half of the pons did not undergo immunohistochemistry.

### FTLTDP-FUS

The patient with FTLTDP-FUS showed minimal tracer retention in the frontal lobes and frontal white matter, which corresponded to areas of significant atrophy ([Fig. 1](#) and [Supplementary Fig. 2](#)). Tracer retention was significantly higher than amyloid-negative healthy controls in the left caudate, as seen in the W-map, but this region was not sampled for pathological analysis. FUS pathology was present in the frontal lobes, and tau pathology was absent in this area. Braak stage was 0. Neurofibrillary threads were present in the entorhinal cortex, dorsal raphe, and locus coeruleus but no tracer retention was evident in these areas ([Supplementary Table 21](#)).





**Figure 4** Comparison of FTP-PET SUVR images with tau immunohistochemistry in patient with FTLN-tau due to *MAPT* S305I mutation. The cerebellar dentate nucleus, entorhinal cortex, globus pallidus, inferior frontal gyrus, middle frontal gyrus (cortex and white matter), and motor cortex are indicated on the patient's SUVR images (arrows), and the corresponding tau (CP-13 antibody) immunohistochemistry is shown. Scale bars = 10  $\mu$ m.

### Evaluating FTP-PET SUVR in precentral gyrus, postcentral gyrus, and subcortical regions of interest

All patients with PSP and CBD showed elevated FTP-PET SUVR levels in the putamen, globus pallidus, and subthalamic nucleus regions of interest that were above the mean +

2 SD SUVR threshold for these regions of interest in young, cognitively normal controls (Fig. 6). Patients with Alzheimer's disease, other non-Alzheimer tauopathies, and FTLN-TDP pathology had similarly elevated tracer retention in these regions compared to controls. SUVR in precentral and postcentral gyrus were highest in patients with Alzheimer's disease, and tracer retention in these regions



**Figure 5 Comparison of FTP-PET SUVR images with tau and TDP-43 immunohistochemistry in a patient with FTLD TDP-43 type A due to GRN (NM\_002087:c.708+6\_708+9del) mutation.** The left middle frontal gyrus is indicated on the patient's SUVR image (arrow). Left middle frontal gyrus tau immunohistochemistry (CP-13 antibody, haematoxylin counterstain, left) was negative, whereas TDP-43 immunohistochemistry (pan-TDP-43 antibody, right) showed short, thin neurites, crescentic neuronal cytoplasmic inclusions, and other typical features of FTLD-TDP type A, consistent with the patient's known mutation in GRN. Scale bars = 500  $\mu$ m and 100  $\mu$ m for the tau and TDP-43 immunohistochemistry micrographs, respectively.

were similar in patients with CBD, PSP, and controls. SUVR in the dentate nucleus region of cerebellum was similar across tauopathies and in controls. PVC was performed to account for the potential loss of sensitivity due to atrophy, which can influence FTP uptake measured in the regions of interest. PVC data yielded similar results (Supplementary Fig. 3).

## Evaluating the Braak neurofibrillary tangle stages detectable by FTP-PET quantification

All patients with primary Alzheimer's disease autopsy diagnosis, who had Braak VI NFT pathology, showed markedly elevated FTP-PET SUVR levels in the Braak stage regions of interest compared to young, cognitively normal controls (Fig. 7). Patient 4, who had primary PSP pathology and contributing Alzheimer's disease pathology [intermediate ADNC (A3, B2, C2), Braak III] showed elevated FTP-PET SUVR in the entorhinal cortex (Braak I) and Braak III/IV regions of interest. Patient 7, who had primary FTLD-tau pathology due to *MAPT* P301L mutation and contributing Alzheimer's disease pathology [intermediate ADNC (A3, B2, C3), Braak IV], did not show elevated FTP-PET SUVR in the entorhinal cortex or Braak III/IV regions of interest. Patient 8, who had primary FTLD-tau pathology due to *MAPT* S305I showed elevated FTP-PET SUVR in the entorhinal cortex region of interest, but this patient did not have any NFT pathology. Patients with primary non-Alzheimer autopsy diagnoses and Braak I–III NFT incidental pathology (low ADNC or not ADNC) did not have FTP-PET SUVR levels that were distinguishable from young, cognitively normal controls (Fig. 7). Overall, this quantitative method detected advanced Alzheimer's disease pathology (Braak VI) but did not reliably detect early (Braak I–IV) NFT pathology.

PVC recovered signal in Braak III/IV regions of interest in the CBD patient with incidental Braak III NFT pathology (Patient 6) and the *MAPT* P301L patient with contributing Alzheimer's disease pathology and Braak IV NFTs (Patient 7) (Supplementary Fig. 4). However, PVC also led to several false positive values in patients with non-Alzheimer tauopathies.

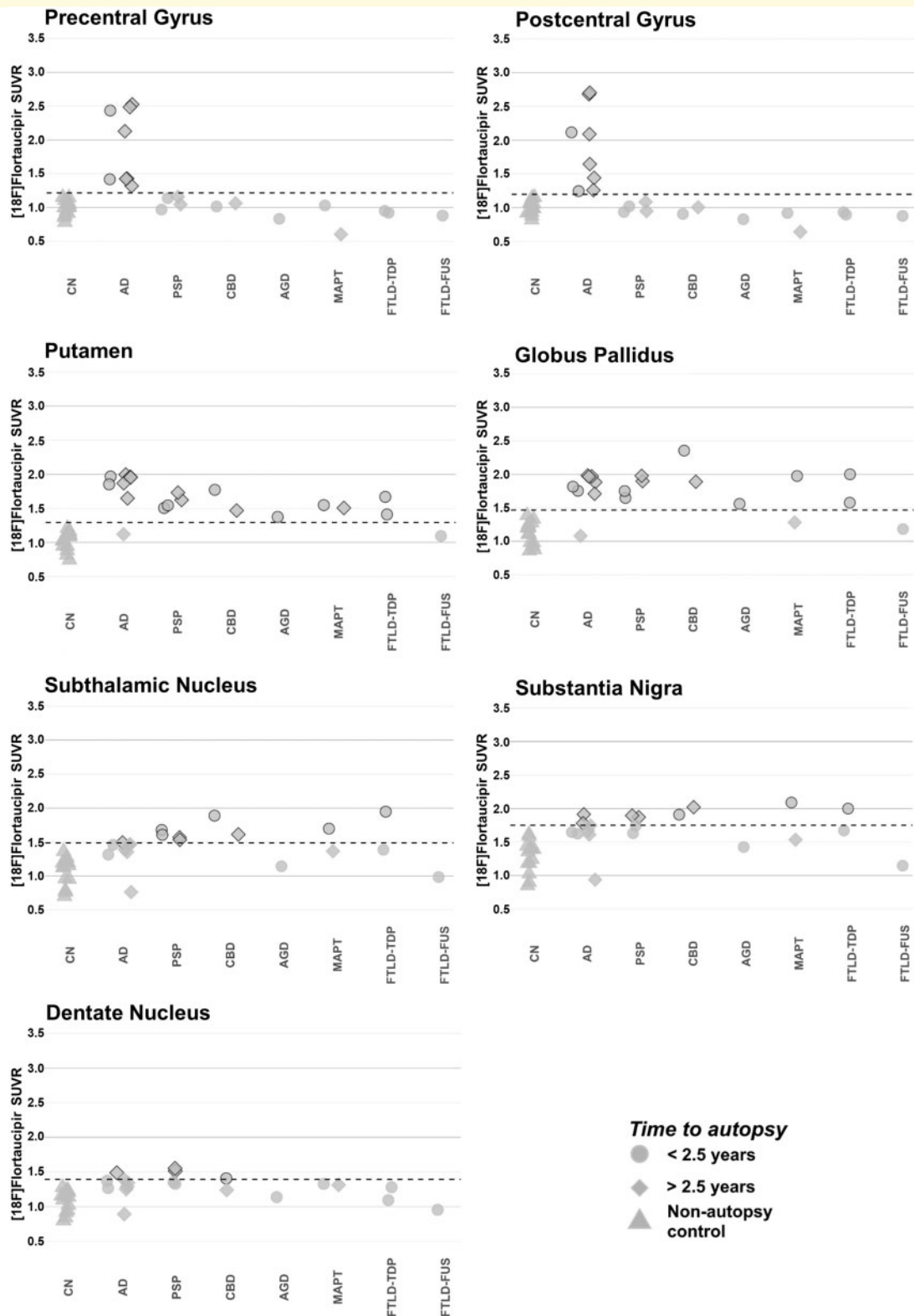
## Discussion

We assessed the relationship between antemortem FTP-PET and autopsy in 20 patients with a broad range of primary neurodegenerative pathologies including Alzheimer's disease, non-Alzheimer tauopathies, FTLD-TDP, and FTLD-FUS. First, we found an excellent correspondence between areas of FTP retention and NFT pathology in patients with primary Alzheimer's disease pathology. Second, patients with non-Alzheimer tauopathies showed a broad range of tracer retention that was elevated compared to controls on W-maps (even within known regions of 'off-target' tracer binding in the basal ganglia and midbrain), though lower in intensity compared to the level of retention characteristic of Alzheimer's disease. This signal usually (but not always) corresponded to locations of non-Alzheimer tau pathology found at autopsy. Patients with FTLD-TDP had a similar level and range of tracer retention compared to patients with non-Alzheimer tauopathies, but the tracer retention was in tau-negative regions. Patients with FTLD-FUS also showed small foci of tracer retention in tau-negative regions. FTP-PET SUVR in subcortical regions susceptible to tau pathology in PSP and CBD were similar in these conditions compared to other neurodegenerative pathologies. Therefore, FTP appears to have poor sensitivity and specificity for non-Alzheimer tauopathies. Third, quantification of FTP-PET SUVR in Braak stage regions of interest could reliably detect increased tracer retention in patients with Braak VI NFT pathology, but there was lower sensitivity for detection of earlier Braak stage (I–IV) pathology, at least in this cohort of patients who died during the later stages of neurodegenerative disease.

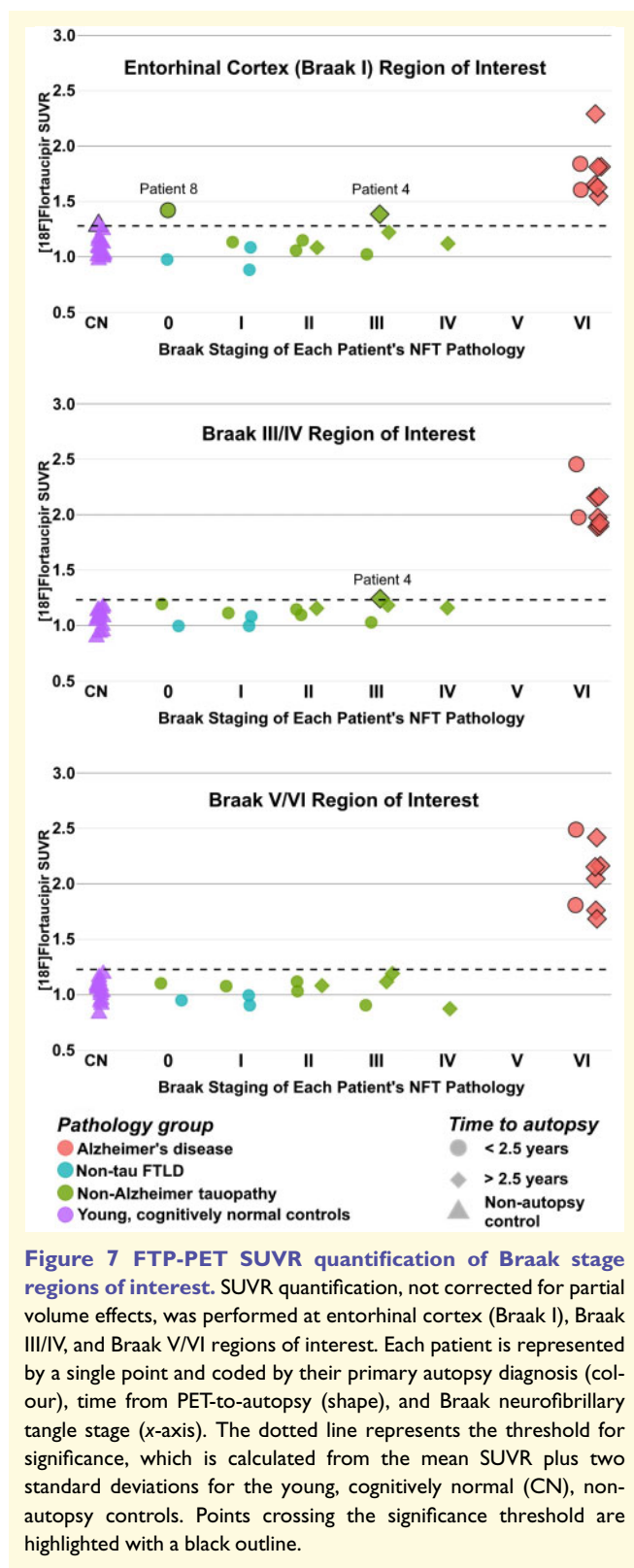
Patients with primary Alzheimer's disease pathology had intense FTP retention in the parietal lobes and posterior cingulate on SUVR images, and W-maps showed significant increases in these areas compared to amyloid-negative healthy controls. Patients with typical and atypical Alzheimer's disease phenotypes had patterns of tracer retention that mirrored the clinical and neuroanatomical variability of these phenotypes (Ossenkoppele *et al.*, 2016), and NFT pathology was found in the areas of tracer retention. All Alzheimer's disease patients had high ADNC (Braak VI) pathology, which was reliably detected on visual (SUVR images and W-maps) and quantitative assessment of FTP-PET.

CBD was associated with asymmetric tracer retention involving the frontal white matter, perirolandic cortex, basal ganglia, and substantia nigra. These areas contained CBD





**Figure 6** FTP-PET SUVR quantification at precentral gyrus, postcentral gyrus, and subcortical regions of interest. SUVR quantification, not corrected for partial volume effects, was performed at precentral gyrus, postcentral gyrus, putamen, globus pallidus, subthalamic nucleus, substantia nigra, and dentate nucleus regions of interest. Each patient is represented by a single point and coded by time from PET-to-autopsy (shape) and primary neuropathological diagnosis (x-axis). The dotted line represents the threshold for significance, which is calculated from the mean SUVR plus two standard deviations for the young, cognitively normal (CN), non-autopsy controls. Points crossing the significance threshold are highlighted with a black outline. AD = Alzheimer's disease; AGD = argyrophilic grain disease.



**Figure 7** FTP-PET SUVR quantification of Braak stage regions of interest. SUVR quantification, not corrected for partial volume effects, was performed at entorhinal cortex (Braak I), Braak III/IV, and Braak V/VI regions of interest. Each patient is represented by a single point and coded by their primary autopsy diagnosis (colour), time from PET-to-autopsy (shape), and Braak neurofibrillary tangle stage (x-axis). The dotted line represents the threshold for significance, which is calculated from the mean SUVR plus two standard deviations for the young, cognitively normal (CN), non-autopsy controls. Points crossing the significance threshold are highlighted with a black outline.

tau pathology. These findings are consistent with previous case reports, which found good PET-to-autopsy comparison for FTP in CBD (Josephs *et al.*, 2016; McMillan *et al.*, 2016; Schonhaut *et al.*, 2017). Patients with CBD showed

lower tracer retention than patients with primary Alzheimer's disease pathology, but more extensive tracer retention than other sporadic non-Alzheimer neurodegenerative pathologies (Smith *et al.*, 2017b). However, the pattern of uptake in CBD was very similar to the pattern observed in a patient with FTLD-TDP type A (Fig. 1), suggesting that FTP-PET has limited ability to differentiate CBD from FTLD-TDP pathology.

PSP was associated with tracer retention in the frontal white matter, globus pallidus, midbrain, and cerebellar dentate nucleus that corresponded with underlying tau pathology. These results are consistent with previous case reports (Marquie *et al.*, 2017a; Passamonti *et al.*, 2017). Interestingly, comparison of two PSP patients with tau pathology in the globus pallidus and cerebellar dentate revealed that the patient with higher tau burden in these regions also had higher corresponding tracer retention. Altogether, there is some concordance between the regional tau burden and FTP retention in PSP, but this may not be a clinically meaningful marker of pathological burden and disease severity. Compared to CBD, patients with PSP usually had more restricted tracer retention on SUVR images and restricted significance on W-maps. The low overall intensity, limited spatial extent, and significant overlap with regions showing 'off-target' tracer uptake in healthy controls suggests that FTP-PET has limited utility for imaging tau pathology in PSP.

Patients with FTLD-tau due to *MAPT* P301L and S305I mutations showed increased tracer retention in several common regions including the temporal lobes, temporal white matter, and basal ganglia. While attribution of tracer uptake in the patient with a P301L mutation was challenging given significant Alzheimer's disease co-pathology, this signal corresponded to regions showing tau inclusion pathology in the patient with the S305I mutation. Mutations in *MAPT* can yield different combinations of tau isoforms and different forms of tau fibrils (Jones *et al.*, 2018; Tsai *et al.*, 2019). Previous studies showed convincing elevated FTP binding in patients with the R406W or V337M mutations, which yield 'Alzheimer's-like' paired helical tau filaments with a mix of 3R/4R isoforms (Smith *et al.*, 2016; Spina *et al.*, 2017; Jones *et al.*, 2018; Tsai *et al.*, 2019). Here we report for the first-time elevated binding in the S305I mutation, corresponding to 4R tau, and showing pathological features similar to argyrophilic grain disease (Kovacs *et al.*, 2008; Tacik *et al.*, 2017). Notably, the burden of post-mortem tau pathology in this case was very high, corresponding to clearly elevated antemortem FTP retention.

Patients with FTLD-TDP and FTLD-FUS showed a range of tracer binding in areas containing non-tau FTLD pathology. These findings in the FTLD-TDP patients are consistent with previous reports of FTP binding in patients with semantic variant primary progressive aphasia and *GRN* and *C9orf72* mutation carriers (Bevan-Jones *et al.*, 2018a, b; Makarets *et al.*, 2018; Smith *et al.*, 2019a; Tsai *et al.*, 2019). The substrate of tracer binding in these cases remains unclear. FTP may show low-level binding to TDP-43 or FUS

aggregates *in vivo* that does not persist *in vitro* following tissue preparations for autoradiography (Marquié *et al.*, 2015; Lowe *et al.*, 2016; Sander *et al.*, 2016). Alternatively, the tracer could bind to other local pathological changes, such as monoamine oxidase B expressed in reactive astrocytes or iron (or associated proteins) in degenerating axons (Lockhart *et al.*, 2017; Lemoine *et al.*, 2018; Baker *et al.*, 2019; Drake *et al.*, 2019). We also cannot exclude the possibility that such cross-reactivity underlies tracer binding in non-Alzheimer tauopathies; however, the broad range of binding observed in patients with a similar degree of neurodegeneration (prominent in some, minimal in others) suggests that FTP binding is not simply a non-specific correlate of neurodegeneration in non-Alzheimer conditions.

Our findings are consistent with a recent large, multi-site study that demonstrated that FTP-PET accurately discriminates Alzheimer's disease from other neurodegenerative disorders (Ossenkoppele *et al.*, 2018), while variable uptake is seen in clinical syndromes associated with both tau-positive and tau-negative neuropathologies. Another 3R/4R tau disorder, chronic traumatic encephalopathy, has modest binding with lower SUVR values than Alzheimer's disease patients (Lesman-Segev *et al.*, 2019; Stern *et al.*, 2019), and a recent case report found only moderate correlations with autopsy (Mantyh *et al.*, 2020). There are several factors that may contribute to difficulties differentiating non-Alzheimer tauopathies from other pathologies and controls. First, FTP autoradiography studies have reported absent-to-low, displaceable binding in non-Alzheimer tauopathies and FTLTDP, with much lower binding than observed to tau NFTs in Alzheimer's disease under the same conditions (Marquié *et al.*, 2015; Lowe *et al.*, 2016; Sander *et al.*, 2016). Notably, we found that FTP-PET SUVR in regions susceptible to PSP and CBD was not elevated in patients with these primary autopsy diagnoses compared to other tauopathies and FTLTDP. Recent cryo-electron microscopy studies have uncovered unique tau ultrastructural differences in Alzheimer's disease, Pick's disease, CBD, and chronic traumatic encephalopathy (Fitzpatrick *et al.*, 2017; Falcon *et al.*, 2018, 2019; Scheres *et al.*, 2020; Zhang *et al.*, 2020), and tau ultrastructure may be a key reason for differential binding of the same tracer across tauopathies. Second, the density and overall mass of tau pathology affects the level of FTP retention. Sporadic 4R tauopathies, in particular PSP, are typically characterized by relatively low overall tau burden compared to Alzheimer's disease (Ono *et al.*, 2017). Notably, FTP signal was quite high in the patient with the *MAPT* S305I mutation, corresponding to a higher burden of tau pathology than in other cases with 4R tau pathology. Lastly, there is a great degree of variability in cortical, white matter, and basal ganglia FTP retention, which could be related to off-target tracer retention (Baker *et al.*, 2019). It will be interesting to see how second-generation tau PET tracers perform across the spectrum of tau neurodegenerative diseases, given their different binding characteristics and pharmacokinetic properties (Leuzy *et al.*, 2019).

All patients with primary Alzheimer's disease autopsy diagnosis, who had Braak VI NFT pathology (high ADNC), showed significantly elevated FTP-PET SUVR levels across Braak stage regions of interest. This quantitative method could reliably detect advanced Alzheimer's disease pathology, but there was difficulty detecting Braak I–III NFT pathology, even after correcting for partial volume effects. Given there was only one patient in our series with Braak IV and no patients with Braak V NFTs, we could not assess the tracer in these clinically meaningful stages. Using a different quantitative method (temporal lobe region of interest), Lowe *et al.* (2019) reliably detected patients who had Braak V and VI NFT pathology. Their study included two patients with Braak IV NFT pathology (one with primary Alzheimer's pathology and another with secondary Alzheimer's pathology) who narrowly crossed their significance threshold. Similar to our results, they were unable to detect Braak I–III NFTs. Additionally, a large autopsy study, with mean PET-to-autopsy time of 2.6 months, found that visual interpretation of FTP scans accurately detected advanced tau (Braak NFT stages V–VI) and Alzheimer's disease (high ADNC) pathology. Patients with earlier Braak NFT pathology on autopsy (Braak NFT stages I–IV) were often read as negative (Fleisher *et al.*, 2020). Based on these results, the United States Food and Drug Administration (FDA) recently approved the use of FTP 'to estimate the density and distribution of aggregated tau neurofibrillary tangles in adult patients with cognitive impairment who are being evaluated for Alzheimer's disease'. The FDA specified that FTP is not indicated for the evaluation of patients with chronic traumatic encephalopathy (FDA News Release, 2020). Our findings further suggest that FTP should not be used in the evaluation of other non-Alzheimer tauopathies in the FTLTDP spectrum. In interpreting results, clinicians should also be aware of potential low sensitivity for lower Braak stages of Alzheimer's disease that may still be clinically meaningful. Fluid biomarkers, such as CSF and plasma phosphorylated tau (pTau181, pTau217) could aid in earlier detection of tau in Alzheimer's disease, and could also differentiate Alzheimer's disease from non-Alzheimer tauopathies and other neurodegenerative diseases (Barthélemy *et al.*, 2020; Janelidze *et al.*, 2020; Mattsson-Carlsson *et al.*, 2020; Thijssen *et al.*, 2020).

The key strength of our study was the broad sample of well-characterized patients who had diverse underlying neuropathologies. A major difference between our sample and a recently published case series is the large number of patients with tau and non-tau FTLTDP pathology (Lowe *et al.*, 2019). Additionally, many patients had Braak I–IV incidental or contributing NFT pathology, even though they had other primary neurodegenerative disorders, which allowed for examination of FTP-PET tracer retention in these early Braak stages.

There were several limitations to this study. First, our sample size was modest, though more diverse compared to previous reports. Second, the quantitative technique used to detect *in vivo* early stage (Braak I–IV) NFT pathology can be affected by the density of NFTs and the size of regions of interest, which probably limits detection of low-level NFT

pathology in atrophic brain tissue contained within the large, bilateral Braak stage regions of interest. Third, we attempted to detect low-level NFT pathology in patients who had other pathological changes and atrophy; thus, our quantitative method may produce different results in other potential applications (e.g. detecting early Braak stage pathology in cognitively normal subjects being evaluated for pre-symptomatic tau accumulation). Developing more sensitive methods and defining the threshold of detection of tau in Alzheimer's disease are important goals for future research. Fourth, PET-to-autopsy interval was variable, but on average, over 2 years. Additional neurodegenerative pathology could have accumulated between imaging and autopsy, which could impact the comparison between regions of tracer retention and autopsy pathology. Fifth, we did not perform quantitative immunohistochemistry, which could be helpful to understand the correlation between the density of different tau pathologies and FTP retention, and also to identify potential non-tau related binding targets (e.g. monoamine oxidase A or B, iron, or metal-associated proteins) (Lockhart *et al.*, 2017; Baker *et al.*, 2019; Drake *et al.*, 2019). Finally, we did not compare FTP binding across tauopathies to binding of other tau tracers that may show different selectivity patterns (Okamura *et al.*, 2018; Betthausen *et al.*, 2019; Leuzy *et al.*, 2019; Lohith *et al.*, 2019; Mormino *et al.*, 2020). Comparing the sensitivity across tracers both *in vitro* and *in vivo* is another important area for future research.

In conclusion, FTP reliably detected high ADNC (Braak VI) pathology on SUVR images, W-maps, and SUVR quantification of Braak stage regions of interest. FTP-PET had poor sensitivity and specificity for non-Alzheimer tauopathies. Quantification of FTP-PET SUVR in Braak stage regions of interest had low sensitivity for early Braak stage NFT pathology.

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## Competing interests

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## Supplementary material

Supplementary material is available at *Brain* online.

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