

Tau immunophenotypes in chronic traumatic encephalopathy recapitulate those of ageing and Alzheimer's disease

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Traumatic brain injury (TBI) is a risk factor for neurodegenerative disease, including chronic traumatic encephalopathy (CTE). Preliminary consensus criteria define the pathognomonic lesion of CTE as patchy tau pathology within neurons and astrocytes at the depths of cortical sulci. However, the specific tau isoform composition and post-translational modifications in CTE remain largely unexplored. Using immunohistochemistry, we performed tau phenotyping of CTE neuropathologies and compared this to a range of tau pathologies, including Alzheimer's disease, primary age-related tauopathy, ageing-related tau astroglialopathy and multiple subtypes of frontotemporal lobar degeneration with tau inclusions. Cases satisfying preliminary consensus diagnostic criteria for CTE neuropathological change (CTE-NC) were identified (athletes, $n = 10$; long-term survivors of moderate or severe TBI, $n = 4$) from the Glasgow TBI Archive and Penn Neurodegenerative Disease Brain Bank. In addition, material from a range of autopsy-proven ageing-associated and primary tauopathies in which there was no known history of exposure to TBI was selected as non-injured controls ($n = 32$). Each case was then stained with a panel of tau antibodies specific for phospho-epitopes (PHF1, CP13, AT100, pS262), microtubule-binding repeat domains (3R, 4R), truncation (Tau-C3) or conformation (GT-7, GT-38) and the extent and distribution of staining assessed. Cell types were confirmed with double immunofluorescent labelling. Results demonstrate that astroglial tau pathology in CTE is composed of 4R-immunoreactive thorn-shaped astrocytes, echoing the morphology and immunophenotype of astrocytes encountered in ageing-related tau astroglialopathy. In contrast, neurofibrillary tangles of CTE contain both 3R and 4R tau, with post-translational modifications and conformations consistent with Alzheimer's disease and primary age-related tauopathy. Our observations establish that the astroglial and neurofibrillary tau pathologies of CTE are phenotypically distinct from each other and recapitulate the tau immunophenotypes encountered in ageing and Alzheimer's disease. As such, the immunohistochemical distinction of CTE neuropathology from other mixed 3R/4R tauopathies of Alzheimer's disease and ageing may rest solely on the pattern and distribution of pathology.

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Abbreviations: ARTAG = ageing-related tau astroglialopathy; CBD = corticobasal degeneration; CTE = chronic traumatic encephalopathy; FTLD-tau = frontotemporal lobar degeneration characterized by tau inclusions; NFT = neurofibrillary tangle; PART = primary age-related tauopathy; PSP = progressive supranuclear palsy; TBI = traumatic brain injury; TSA = thorn-shaped astrocytes

Introduction

There is increasing recognition of the association between exposure to traumatic brain injury (TBI) and risk of subsequent neurodegenerative disease, in particular chronic traumatic encephalopathy (CTE). First described as ‘punch drunk syndrome’ in 1928 (Martland, 1928) and later ‘dementia pugilistica’ (Millsbaugh, 1937), progressive neurodegeneration in the context of TBI was historically considered a consequence of participation in boxing. However, more recent descriptions of the neurodegenerative pathology of dementia pugilistica, now recognized as CTE, in non-boxer athletes exposed to repetitive mild TBI (Corsellis *et al.*, 1973; Geddes *et al.*, 1999; Omalu *et al.*, 2011; Saing *et al.*, 2012; McKee *et al.*, 2013, 2016; Smith *et al.*, 2013, 2019; Stewart *et al.*, 2016; Johnson *et al.*, 2017; Mez *et al.*, 2017; Wilson *et al.*, 2017; Lee *et al.*, 2019) and late survivors of moderate or severe TBI (Johnson *et al.*, 2012; Zanier *et al.*, 2018; Kenney *et al.*, 2018) have brought the potential life-long consequences of TBI exposure to wider attention. Nevertheless, the current consensus criteria for CTE neuropathological assessment remain preliminary and there are few detailed accounts of its similarities and differences with other tauopathies.

While intraneuronal tau aggregates in the form of neurofibrillary tangles (NFTs) have long been described in dementia pugilistica/CTE, more recent studies also note the presence of pathological astroglial tau accumulation in the form of thorn-shaped astrocytes (TSA) (Ikeda *et al.*, 1995, 1998; Schmidt *et al.*, 2001; McKee *et al.*, 2009, 2013; Saing *et al.*, 2012; Kanaan *et al.*, 2016; Mez *et al.*, 2017). Indeed, the importance of this mixed neuronal and astroglial pathology in CTE is reflected in preliminary consensus diagnostic criteria for the disease, which propose the pathognomonic lesion as ‘phosphorylated-tau aggregates in neurons, astrocytes, and cell processes around small vessels in an irregular pattern at the depths of the cortical sulci’ (McKee *et al.*, 2016). This sulcal lesion is suggested as sufficiently unique to distinguish CTE from other tau-associated neurodegenerative diseases including Alzheimer’s disease, frontotemporal lobar degeneration characterized by tau inclusions (FTLD-tau)—including progressive supranuclear palsy (PSP), corticobasal degeneration (CBD) and Pick’s disease (Cairns *et al.*, 2007; Montine *et al.*, 2012; Kovacs, 2015), as well as the pathologies of primary age-related tauopathy (PART) and

ageing-related tau astroglialopathy (ARTAG) (Crary *et al.*, 2014; Kovacs *et al.*, 2016). However, there remains debate regarding whether tau found at sulcal depths is solely found in individuals with a history of head impacts, or if it can occur in individuals with no contact sport or TBI history (Iverson *et al.*, 2019).

Beyond the potentially unique distribution of tau pathologies in CTE, little is known regarding tau composition and post-translational modifications. Of the six tau isoforms containing either 3 or 4 microtubule-binding repeat domains (3R versus 4R tau) (Goode *et al.*, 2000), NFTs in Alzheimer’s disease and PART contain both 3R and 4R tau (Espinoza *et al.*, 2008; Santa-Maria *et al.*, 2012; Crary *et al.*, 2014). In contrast, while the Pick bodies and ramified astrocytes of Pick’s disease contain primarily 3R tau (Irwin *et al.*, 2016), the astrocytic tau pathologies of PSP, CBD and ARTAG are composed of 4R tau (Cairns *et al.*, 2007; Ferrer *et al.*, 2014). Regarding CTE pathology, biochemical analysis of material from two former boxers with dementia pugilistica/CTE reported NFTs comprising both 3R and 4R tau, similar to Alzheimer’s disease (Schmidt *et al.*, 2001; McKee *et al.*, 2013). A further case series indicates that tau in CTE displays conformations and phospho-epitopes comparable to those observed early in NFT maturation in Alzheimer’s disease (Kanaan *et al.*, 2016). Nonetheless, comprehensive assessment of the phosphorylation, truncation and conformation of both glial and neuronal tau pathologies in CTE has not previously been assessed, particularly within the context of other established tauopathies (Ferrer *et al.*, 2014; Kovacs, 2015; Irwin *et al.*, 2016).

Here, we report immunohistochemical observations on extensive tau phenotypic analysis of autopsy-derived material from individuals with known CTE neuropathologies and histories of exposure to either repetitive sport-related mild TBI, or moderate/severe TBI, when compared to ageing-associated and primary neurodegenerative tauopathies. Specifically, applying antibodies specific for tau isoform, multiple phospho-epitopes, caspase-mediated truncation, and novel monoclonal antibodies capable of differentiating 3R/4R tau in Alzheimer’s disease from the conformationally distinct FTLD-tau (Gibbons *et al.*, 2018, 2019), we demonstrate that while the NFTs and astroglial tau pathologies of CTE differ immunohistochemically from each other, they recapitulate features of Alzheimer’s disease/PART and ARTAG, respectively.

Materials and methods

Cohort demographics

All cases were obtained from the Glasgow TBI Archive, Department of Neuropathology, Queen Elizabeth University Hospital, Glasgow, UK or the University of Pennsylvania Center for Neurodegenerative Disease Research (CNDR) Brain Bank, Philadelphia, PA, USA. Brain tissue was acquired by means of planned donation or at routine diagnostic autopsy. Ethical approval for use of tissue in this study was provided by the West of Scotland Research Ethics Committee (Project ID 225271); and the Greater Glasgow and Clyde Biorepository (Application Number 340), as well as the institutional review board of the University of Pennsylvania.

Cases were selected with a history of participation in contact sports: American football ($n = 4$), rugby ($n = 3$), soccer ($n = 2$), or boxing ($n = 1$), or a remote history of single moderate or severe TBI caused by assault ($n = 1$), motor vehicle collision ($n = 1$) or fall ($n = 1$). One additional case sustained one mild and one moderate TBI caused by falls ($n = 1$). Cases were selected with previously confirmed CTE neuropathological change (CTE-NC) (Lee *et al.*, 2019) based on the preliminary consensus criteria for the neuropathological evaluation of CTE (McKee *et al.*, 2016). Notably, in addition to CTE neuropathology, multiple cases also displayed co-morbid neuropathologies ($n = 8$) as has previously been reported (Lee *et al.*, 2019). Clinical, demographic and neuropathological information, including integrated clinicopathological diagnoses (Lee, 2018) for all cases is presented in Table 1.

To permit comparisons of CTE tau pathologies with those of established neurodegenerative disease, material from patients without documented history of TBI or participation in contact sport was selected that met neuropathological criteria for the diagnosis of Alzheimer's disease (Braak stage V or VI; $n = 6$) or FTLD-tau as Pick's disease ($n = 6$), PSP ($n = 6$), or CBD ($n = 6$). In addition, non-demented control cases without history of TBI and with known ageing-related tau pathologies were selected including: PART ($n = 1$), ARTAG and PART ($n = 3$) and ARTAG with low Alzheimer's disease neuropathological change ($n = 4$) (Cairns *et al.*, 2007; Montine *et al.*, 2012; Crary *et al.*, 2014; Kovacs *et al.*, 2016; McKee *et al.*, 2016) (Table 1).

Brain tissue handling

Whole brains from the Glasgow TBI archive were fixed in 10% formol saline at autopsy for a minimum of 2 weeks prior to dissection. Standardized anatomical sampling, tissue processing and paraffin embedding were performed as previously described (Graham *et al.*, 1995). From the University of Pennsylvania CNDR Brain Bank, tissue blocks cut from fresh brains were fixed overnight in 70% ethanol and 150 mM sodium chloride or 10% neutral buffered formalin and processed to paraffin as previously described (Toledo *et al.*, 2014). From each case, 8- μ m tissue sections were prepared from regions with stereotypical CTE neuropathology at the depths of cortical sulci. For comparison, regions displaying hallmark, disease-specific pathologies were selected from non-trauma control cases of Alzheimer's disease (NFTs), PART (NFTs), Pick's disease (Pick bodies and ramified astrocytes), PSP (tufted astrocytes), CBD

(astrocytic plaques) and ARTAG (TSA) (Cairns *et al.*, 2007; Montine *et al.*, 2012; Crary *et al.*, 2014; Kovacs *et al.*, 2016).

Single immunohistochemical labelling

Serial tissue sections for all cases were subjected to deparaffinization and rehydration to H₂O before being immersed in 3% aqueous H₂O₂ (15 min) to quench endogenous peroxidase activity. Antigen retrieval was performed via microwave pressure cooker in either Tris/EDTA or citrate buffer, with or without formic acid pretreatment, as optimized for each antibody (Supplementary Table 1). Sections were blocked using normal horse serum (Vector Labs) in OptimaxTM buffer (BioGenex) for 30 min followed by incubation in the primary antibody overnight at 4°C. Specifically, a panel of tau antibodies (Supplementary Table 1) was applied targeting multiple phospho-epitopes including S202 (CP13) (Jicha *et al.*, 1999), S396/S404 (PHF1) (Greenberg *et al.*, 1992; Otvos *et al.*, 1994), S212/T214 (AT100) (Hoffmann *et al.*, 1997; Zheng-Fischhofer *et al.*, 1998) and S262; 3 or 4 microtubule-binding domain repeats (RD3 and RD4) (de Silva *et al.*, 2003); and caspase-cleaved tau at Asp421 (Tau-C3) (Gamblin *et al.*, 2003). In addition we applied the recently characterized antibodies GT-7 and GT-38 (Gibbons *et al.*, 2018, 2019). Evidence from co-immunofluorescence studies in human tissue with FTLD-tau and Alzheimer's disease-tau suggest that GT-38 requires the presence of both 3R and 4R tau. Moreover, it was demonstrated that GT-38 binding requires a pathological conformation of Alzheimer's disease-tau since chemical denaturation leads to a reduction of GT-38 binding (Gibbons *et al.*, 2018).

After rinsing, sections were incubated in a biotinylated universal secondary antibody (Vector Labs) for 30 min, followed by the avidin-biotin complex for 30 min (Vector Labs). Visualization was achieved using the DAB peroxidase substrate kit (Vector Labs). Sections were counterstained with haematoxylin, followed by rinsing, dehydration, and coverslipping using CytosealTM 60. Tissue sections from a case with neuropathologically confirmed Alzheimer's disease were included as a positive control in all staining procedures. Omission of the primary antibody using the same Alzheimer's disease case was performed in parallel to control for non-specific binding. Notably, three cases [Cases 9 (CTE neuropathology), 11 (CTE neuropathology) and 15 (ARTAG/PART)] (Table 2) failed to demonstrate immunoreactivity to antibodies specific for 3R and 4R tau due to fixation sensitivity, as has been reported previously with these antibodies (Espinoza *et al.*, 2008; Ferrer *et al.*, 2014).

Double immunofluorescent labelling

Serial tissue sections from a subset of cases (CTE $n = 5$, ARTAG $n = 2$, Alzheimer's disease $n = 2$) were selected for double labelling immunofluorescence to confirm and validate morphological identification of cell types (astrocytes versus neurons) as identified by both PHF1 and GT-38. Specifically, sections were labelled with combinations of tau antibodies (PHF1 or GT-38) and cell-type specific markers, namely MAP2 for neurons and GFAP for astrocytes using established protocols (Johnson *et al.*, 2016). Briefly, following deparaffinization and rehydration, antigen retrieval was performed as described above and tissue blocked in the relevant species-specific serum (1%) (Vector

Table 1 Demographic and clinical summary of cases

Case	Group	Age	Sex	Sport / TBI exposure	Integrated clinicopathologic diagnosis	PMI, h	Source	Anatomical region examined
1	CTE-NC	40s	M	Football	CBD	7	Penn-CNDR	Frontal
2	CTE-NC	60s	M	Football	CBD	3	Penn-CNDR	Frontal
3	CTE-NC	70s	M	Football	DLB	18	Penn-CNDR	Frontal
4	CTE-NC	80s	M	Football	FTLD-TDP	7	Penn-CNDR	Frontal
5	CTE-NC	60s	M	Boxing	CTE	24	GTBIA	Temporal
6	CTE-NC	70s	M	Rugby	CTE	12	GTBIA	Insular
7	CTE-NC	70s	M	Rugby	AD	48	GTBIA	Frontal
8	CTE-NC	70s	M	Rugby	Mixed AD/VaD	48	GTBIA	Frontal
9	CTE-NC	50s	M	Soccer	CTE	Unknown	GTBIA	Insular
10	CTE-NC	80s	M	Soccer	AD	24	GTBIA	Frontal
11	CTE-NC	50s	M	sTBI	Remote TBI - no NDD	108	GTBIA	Frontal
12	CTE-NC	60s	M	sTBI	Remote TBI - no NDD	24	GTBIA	Temporal
13	CTE-NC	70s	M	sTBI	CTE	24	GTBIA	Temporal
14	CTE-NC	70s	M	Mild and moderate TBI	PDD	7.5	Penn-CNDR	Angular
15	ARTAG, PART	50s	M	No	No NDD	80.5	GTBIA	Frontal
16	ARTAG, PART	80s	M	No	PART	19	Penn-CNDR	Temporal
17	ARTAG, PART	70s	M	No	No NDD	17	Penn-CNDR	Temporal
18	PART	60s	M	No	No NDD	11	Penn-CNDR	Temporal
19	ARTAG, low ADNC	70s	F	No	No NDD	18	Penn-CNDR	Amygdala
20	ARTAG, low ADNC	70s	F	No	No NDD	19	Penn-CNDR	Amygdala
21	ARTAG, low ADNC	70s	F	No	No NDD	18	Penn-CNDR	Temporal
22	ARTAG, low ADNC	80s	M	No	No NDD; CVD	7	Penn-CNDR	Amygdala
23	AD	60s	M	No	AD	13.5	Penn-CNDR	Angular
24	AD	60s	M	No	AD	5	Penn-CNDR	Frontal
25	AD	70s	M	No	AD	4	Penn-CNDR	Angular
26	AD	70s	M	No	AD	8.5	Penn-CNDR	Angular
27	AD	70s	F	No	AD	11	Penn-CNDR	Temporal
28	AD	80s	F	No	AD	6	GTBIA	Cingulate
29	FTLD-Tau	50s	M	No	PiD	11	Penn-CNDR	Temporal
30	FTLD-Tau	60s	M	No	PiD	Unknown	Penn-CNDR	Temporal
31	FTLD-Tau	70s	M	No	PiD	22	Penn-CNDR	Frontal
32	FTLD-Tau	70s	M	No	PiD	4	Penn-CNDR	Angular
33	FTLD-Tau	80s	F	No	PiD	24	Penn-CNDR	Frontal
34	FTLD-Tau	50s	M	No	PiD	10	Penn-CNDR	Frontal
35	FTLD-Tau	70s	M	No	PSP	14	Penn-CNDR	Frontal
36	FTLD-Tau	80s	M	No	PSP	23	Penn-CNDR	Angular
37	FTLD-Tau	70s	M	No	PSP	23	Penn-CNDR	Frontal
38	FTLD-Tau	60s	M	No	PSP	13	Penn-CNDR	Angular
39	FTLD-Tau	70s	M	No	PSP	17	Penn-CNDR	Temporal
40	FTLD-Tau	70s	M	No	PSP	19	Penn-CNDR	Frontal
41	FTLD-Tau	60s	F	No	CBD	16	Penn-CNDR	Temporal
42	FTLD-Tau	70s	M	No	CBD	11	Penn-CNDR	Angular
43	FTLD-Tau	50s	M	No	CBD	41	Penn-CNDR	Frontal
44	FTLD-Tau	70s	M	No	CBD	7.5	Penn-CNDR	Temporal
45	FTLD-Tau	80s	M	No	CBD	11	Penn-CNDR	Frontal
46	FTLD-Tau	70s	M	No	CBD	5	Penn-CNDR	Frontal

AD = Alzheimer's disease; ADNC = Alzheimer's disease neuropathological change; CTE-NC = CTE-neuropathological change; CVD = cerebrovascular disease; DLB = dementia with Lewy bodies; F = female; FTLD-TDP = frontotemporal lobar degeneration with TDP-43 inclusions; GTBIA = Glasgow TBI archive; M = male; NDD = neurodegenerative disease; PDD = Parkinson's disease dementia; Penn-CNDR = University of Pennsylvania Center for Neurodegenerative Disease Research; PiD = Pick's disease; PMI = post-mortem interval; sTBI = single moderate or severe TBI; VaD = vascular dementia.

Labs). Primary antibodies were applied serially for 20 h (4°C) specific for PHF1 (1:100) or GT-38 (1:100), followed by glial fibrillary acidic protein (GFAP) (Abcam; 1:200) or the microtubule-associated protein 2 (MAP2) (Abcam; 1:200). After

rising, the corresponding Alexa Fluor® (Invitrogen) secondary antibody was applied at 1:500 in a 2% species-specific blocking solution for 2 h at room temperature. Serial sections of positive control tissue (Alzheimer's disease) were subjected to the entire

Table 2 Tau phenotype semiquantitative scores: 3R, 4R and conformation

Case	3R		4R		GT-7		GT-38	
	TSA	NFT	TSA	NFT	TSA	NFT	TSA	NFT
1 CTE-NC	–	++	+++	+++	–	++	–	++
2 CTE-NC	–	++	+++	+++	–	++	–	++
3 CTE-NC	–	+++	+++	+++	–	+++	–	+++
4 CTE-NC	+	++	+++	+++	–	++	+	++
5 CTE-NC	–	+++	+++	+++	–	+++	–	+++
6 CTE-NC	–	+++	+++	+++	–	+++	–	+++
7 CTE-NC	+	+++	+++	+++	–	+++	–	+++
8 CTE-NC	–	+++	+++	+++	+	+++	+	+++
9 CTE-NC	x	x	x	x	x	x	x	x
10 CTE-NC	–	+++	+++	+++	–	+++	–	+++
11 CTE-NC	x	x	x	x	–	+++	–	+++
12 CTE-NC	–	++	++	++	–	++	–	++
13 CTE-NC	–	+++	++	++	+	+++	+	+++
14 CTE-NC	–	+++	+++	+++	+	++	–	++
15 ARTAG, PART	x	x	x	x	–	+++	–	+++
16 ARTAG, PART	–	+	+++	+++	–	+++	–	+++
17 ARTAG, PART	–	+++	+++	+++	–	+++	–	+++
18 PART	N/A	++	N/A	++	N/A	++	N/A	++
19 ARTAG, low ADNC	–	+++	+++	+++	–	++	–	+++
20 ARTAG, low ADNC	+	++	+++	+++	+	+++	+	+++
21 ARTAG, low ADNC	–	+++	+++	+++	–	+++	–	+++
22 ARTAG, low ADNC	–	+++	+++	+++	–	+++	–	+++
23 AD		+++		+++		+++		+++
24 AD		+++		+++		+++		+++
25 AD		+++		+++		+++		+++
26 AD		+++		+++		+++		+++
27 AD		+++		+++		+++		+++
28 AD		++		++		+++		+++
	Pick bodies	Ramified astro	Pick bodies	Ramified astro	Pick bodies	Ramified astro	Pick bodies	Ramified astro
29 PiD	+++	+++	–	++	–	–	–	–
30 PiD	+++	N/A	–	N/A	–	N/A	–	N/A
31 PiD	+++	+++	–	–	–	–	–	–
32 PiD	+++	N/A	–	N/A	–	N/A	–	N/A
33 PiD	+++	N/A	–	N/A	–	N/A	–	N/A
34 PiD	+++	+++	–	–	–	–	–	–
	Tufted astrocytes		Tufted astrocytes		Tufted astrocytes		Tufted astrocytes	
35 PSP			+++		–		–	
36 PSP			+++		–		–	
37 PSP			++		–		–	
38 PSP			+++		–		–	
39 PSP			++		–		–	
40 PSP			+++		–		–	
	Astrocytic plaques		Astrocytic plaques		Astrocytic plaques		Astrocytic plaques	
41 CBD			+++		–		–	
42 CBD			+++		–		–	
43 CBD			+++		–		–	
44 CBD			+++		–		–	
45 CBD			+++		–		–	
46 CBD			+++		–		–	

Semiquantitative scoring: – = absent, or nearly absent ($\leq 5\%$ concordance with extent of PHF1); + = minimal (5–30% concordance); ++ = moderate (30–70% concordance); +++ = extensive ($> 70\%$ concordance); x = no immunoreactivity.

AD = Alzheimer's disease; ADNC = Alzheimer's disease neuropathological change; CTE-NC = CTE neuropathological change; N/A = not applicable.

procedure with omission of subsets of primary antibodies to control for non-specific immunofluorescence. Following rinsing, all double fluorescent-immunolabelled sections were incubated in TrueView autofluorescence quenching reagent (Vector Labs) for 5 min at room temperature before being rinsed and cover-slipped using VECTASHIELD® mounting medium (Vector Labs).

Analysis of immunohistochemical findings

Using a standardized approach, the extent of immunoreactivity for each antibody was scored relative to that observed using an index antibody in each individual case, as has been described previously (Ferrer *et al.*, 2014). Specifically, a semi-quantitative score was used to denote the extent of immunoreactivity relative to that of PHF1 as: absent or nearly absent (<5% concordance with the extent of PHF1 immunoreactivity); minimal (5–30% concordance); moderate (30–70% concordance); or extensive (>70% concordance). PHF1 was selected as the index antibody given its widely reported use for the identification of tau pathologies across neurodegenerative diseases and recommended use for the identification of CTE pathology in preliminary consensus criteria (McKee *et al.*, 2016). Under this protocol, scoring does not reflect the number of positive cells or permit comparisons of the extent of pathology between cases, but rather reflects the relative extent to which a particular antibody recognizes the burden of tau pathology as identified via PHF1 in each field of interest. Glial versus neuronal pathologies were distinguished based on characteristic cellular morphologies. A subset of sections was reviewed and scored independently by two observers (J.D.A. and V.E.J.), with good interrater reliability (Cohen's $\kappa = 0.71$). Where there was a discrepancy in scoring, cases were jointly reviewed and a consensus score reached.

Data availability

The authors confirm that the data supporting the findings of this study are available within the article and its [Supplementary material](#).

Results

Consistent with prior descriptions of CTE neuropathology (McKee *et al.*, 2016), each TBI case displayed PHF1-positive neurons and astrocytes in a patchy and perivascular distribution concentrated at the depths of cortical sulci (Fig. 1). Astrocytes typically displayed thorn-shaped morphologies, with short, thickened processes (Ikeda *et al.*, 1995; Kovacs *et al.*, 2016, 2017a), frequently in the immediate subpial region at the sulcal depth, in addition to being observed in a patchy and perivascular distribution within deeper layers of cortex. Neuronal tau pathology at the depths of cortical sulci in cases with CTE neuropathology displayed the morphology of NFTs, consistent with historical and contemporary descriptions (Corsellis *et al.*, 1973; Geddes *et al.*, 1996, 1999; Omalu *et al.*, 2005; McKee *et al.*, 2009, 2013,

2016). The distinctive cellular morphologies of CTE astrocytes and neurons were confirmed via double immunofluorescence labelling on a subset of cases as described. Specifically, NFTs identified via PHF1 co-localized with MAP2, but not GFAP. Conversely, TSAs were observed to co-localize with GFAP but not MAP2, consistent with findings in Alzheimer's disease and ARTAG (Supplementary Figs 1–3).

As expected, neurodegenerative disease control subjects displayed the hallmark and cell-specific tau pathologies characteristic for each diagnosis as identified by PHF1. Specifically, FTLD-tau controls demonstrated Pick bodies and, in some cases, ramified astrocytes in Pick's disease, tufted astrocytes in PSP and astrocytic plaques in CBD. Alzheimer's disease cases had NFTs in a bilaminar cortical distribution in addition to diffuse neuritic threads. Controls with ageing-related pathologies displayed limited cortical NFT pathology consistent with PART or low Alzheimer's disease neuropathological change, as well as TSAs consistent with ARTAG in white matter, perivascular, grey matter, subpial, or subependymal distributions.

Astroglial pathology of CTE contains 4R tau only, whereas neurofibrillary tangles are composed of both 3R and 4R tau

Tau positive astrocytes within regions of CTE neuropathology were composed of 4R tau only in virtually all cases (Figs 2, 3 and Table 2). Specifically, in cases demonstrating adequate immunoreactivity, robust and consistent immunoreactivity to RD4 was observed in astrocytes within the subpial region, as well as those extending to deeper cortical layers. In contrast, astrocytic immunoreactivity to the RD3 antibody was absent in all but two cases where just minimal staining was observed. Notably, TSAs of CTE were morphologically indistinguishable from those within non-injured control cases with ARTAG, which were also composed almost entirely of 4R tau, consistent with previous reports (Lopez-Gonzalez *et al.*, 2013; Ferrer *et al.*, 2018) (Fig. 3). Several cases with CTE neuropathology displayed TSA elsewhere in the tissue sections examined, including within subcortical white matter, subependymal, and subpial regions, in keeping with descriptions of ARTAG. These astrocytes were morphologically indistinguishable from those in the sulcal depths associated with stereotypical CTE neuropathology and displayed the same pattern of selective 4R tau immunoreactivity.

In contrast, ramified astrocytes of Pick's disease were composed of 3R tau, with a subset of cells in just one of the six cases also displaying 4R immunoreactivity, as has been reported previously in a subset of cases (Arai *et al.*, 2001; Ferrer *et al.*, 2014; Irwin *et al.*, 2016). The tufted astrocytes in PSP and astrocytic plaques in CBD controls displayed immunoreactivity for 4R tau only (Fig. 3).

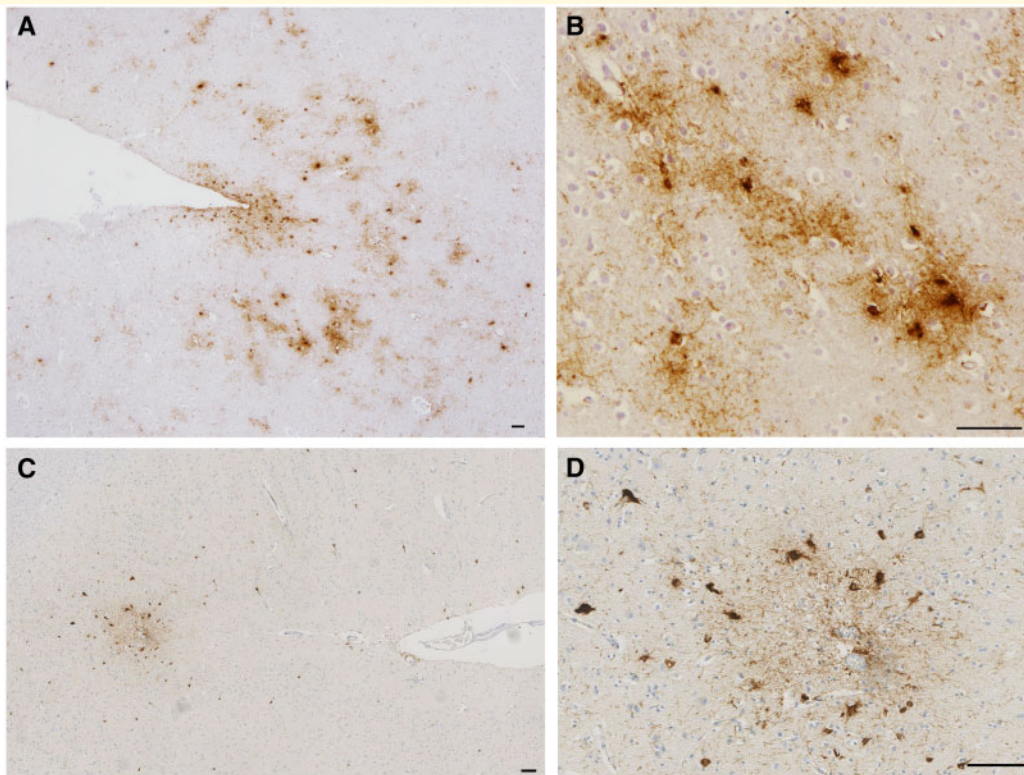


Figure 1 Sulcal depth astrocytic and neuronal tau pathologies in CTE. (A and B) PHF1 immunohistochemistry reveals NFT and TSA pathology concentrated at the depths of cortical sulci of a former American football player (Case 3) and (C and D) chronic survivor of a single severe TBI (Case 11). (B and D) High magnification images from the same sulci showing perivascular pathology composed of mixed neuronal and astrocytic populations consistent with the preliminary diagnostic criteria for CTE. Scale bars = 100 μ m.

While astrocytes in CTE were typically 4R tau-immunoreactive/3R tau-negative, NFTs in these regions were immunoreactive for both 3R and 4R tau isoforms, similar to those of Alzheimer's disease and PART (Fig. 2 and Table 2) and distinct from Pick bodies (3R positive only) (Fig. 2).

Post-translational modification of tau in CTE neuropathology is consistent with ARTAG and Alzheimer's disease

The post-translational modifications of tau within astrocytes and NFTs of CTE were observed to recapitulate those of ARTAG and Alzheimer's disease with respect to all antibodies assessed. Specifically, tau immunoreactive astrocytes in CTE exhibited phosphorylation at residues S202 (CP13), S212/T214 (AT100), S262, and S396/S404 (PHF1) (Fig. 4 and Table 3). Typically, CP13, AT100, and PHF1 displayed dense cytoplasmic staining throughout the cell body. In contrast, pS262 immunoreactivity displayed both robust and confluent immunoreactivity (Fig. 4), as well as a more dot-like pattern of immunoreactivity in a subset of cases/cells that was often concentrated in the peripheral processes of

the astrocyte (Supplementary Fig. 4). Caspase-mediated truncation at D421 (Tau-C3), however, was virtually never seen in astrocytes in CTE, with just minimal cells observed in a single case. Again, all findings were consistent across TSA in subpial and deeper cortical astrocytes within the sulcal depth. This profile of staining was also indistinguishable from that observed in non-injured, ARTAG control cases (Fig. 4). In contrast with the astroglial pathologies of CTE, ARTAG, Pick's disease and CBD, which did not typically demonstrate immunoreactivity to Tau-C3, subsets of tufted astrocytes in two PSP cases demonstrated Tau-C3 immunoreactivity, consistent with previous reports (Fig. 4) (Ferrer *et al.*, 2014).

Notably, NFTs in CTE also displayed robust cytoplasmic immunoreactivity for all tau phospho-epitopes including S202 (CP13), S212/T214 (AT100), S262 and S396/S404 (PHF1) (Fig. 4). Moreover, a subpopulation of NFTs in CTE demonstrated evidence of caspase-mediated truncation of tau at D421 (Tau-C3) (Fig. 4), consistent with that observed previously (Kanaan *et al.*, 2016), and in Alzheimer's disease here and in prior reports (Gamblin *et al.*, 2003).

FTLD-tau controls were consistent with prior characterizations performed using these antibodies (Buee and Delacourte, 1999; Guillozet-Bongaarts *et al.*, 2007; Ferrer

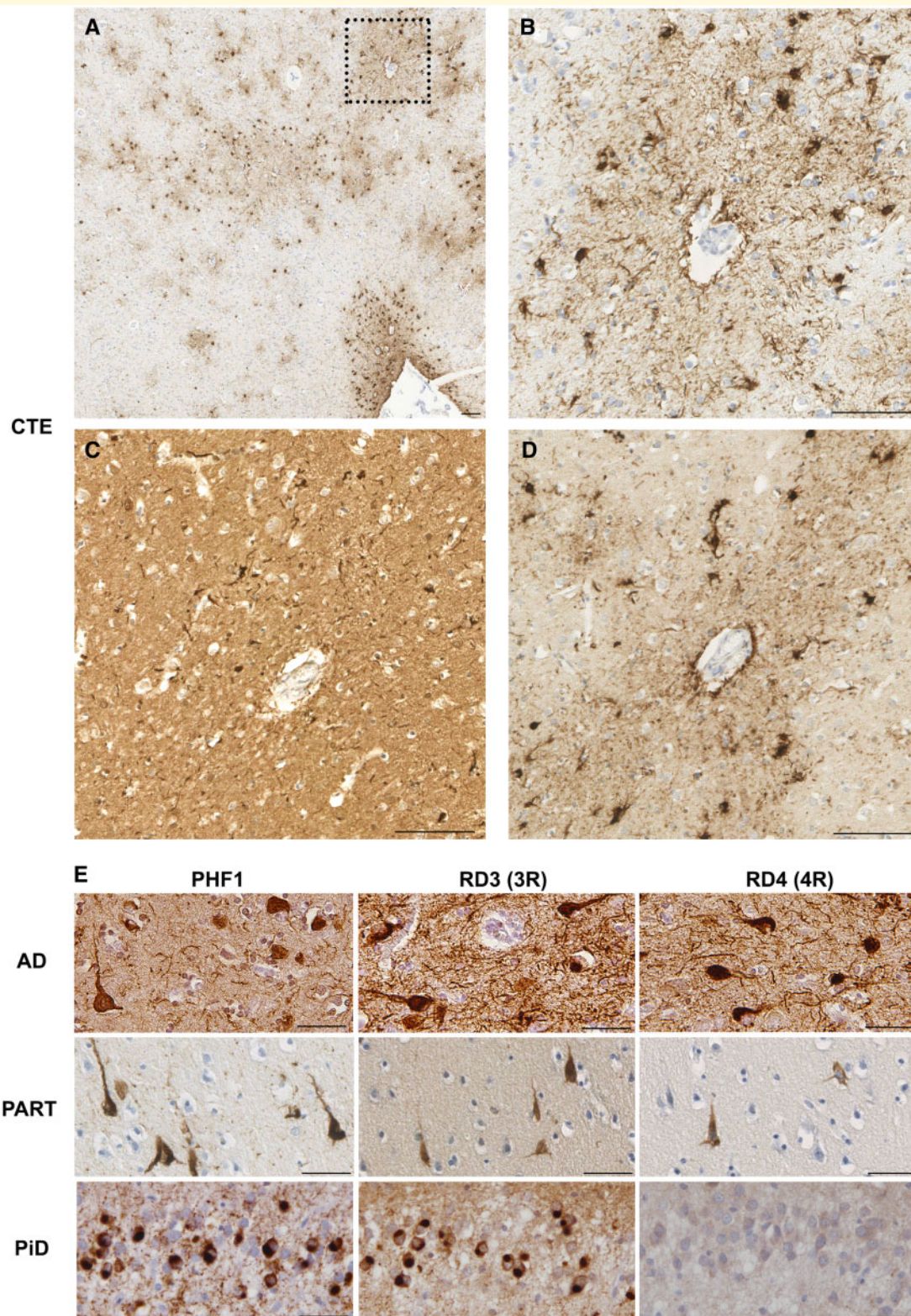


Figure 2 3R versus 4R tau immunoreactivity in CTE, Alzheimer's disease, PART and Pick's disease. (A) Sulcal depth CTE neuropathology with prominent subpial TSAs in addition to patchy and perivascular TSAs and NFTs within the deeper layers of cortex (Case 10; PHF1 staining). (B) Higher magnification of box in A displaying perivascular astrocytic and neuronal tau pathologies. (C) Immunoreactivity-specific for 3R tau in the same region as in B showing perivascular NFTs, but an absence of immunoreactivity within astrocytes. (D) In contrast, immunohistochemistry specific for 4R tau identified cells with both neuronal and astrocytic morphologies in the same region. (E) Consistent with previous descriptions, cases meeting diagnostic criteria for Alzheimer's disease (AD; Case 25) and PART (Case 15) displayed neurofibrillary tangles in the cortex that were immunoreactive for 3R and 4R tau. In contrast, Pick bodies of Pick's disease (PiD) within the dentate granule cells of Case 29 were composed of only 3R tau. Scale bars = 100 μ m in A–D, 50 μ m in E.

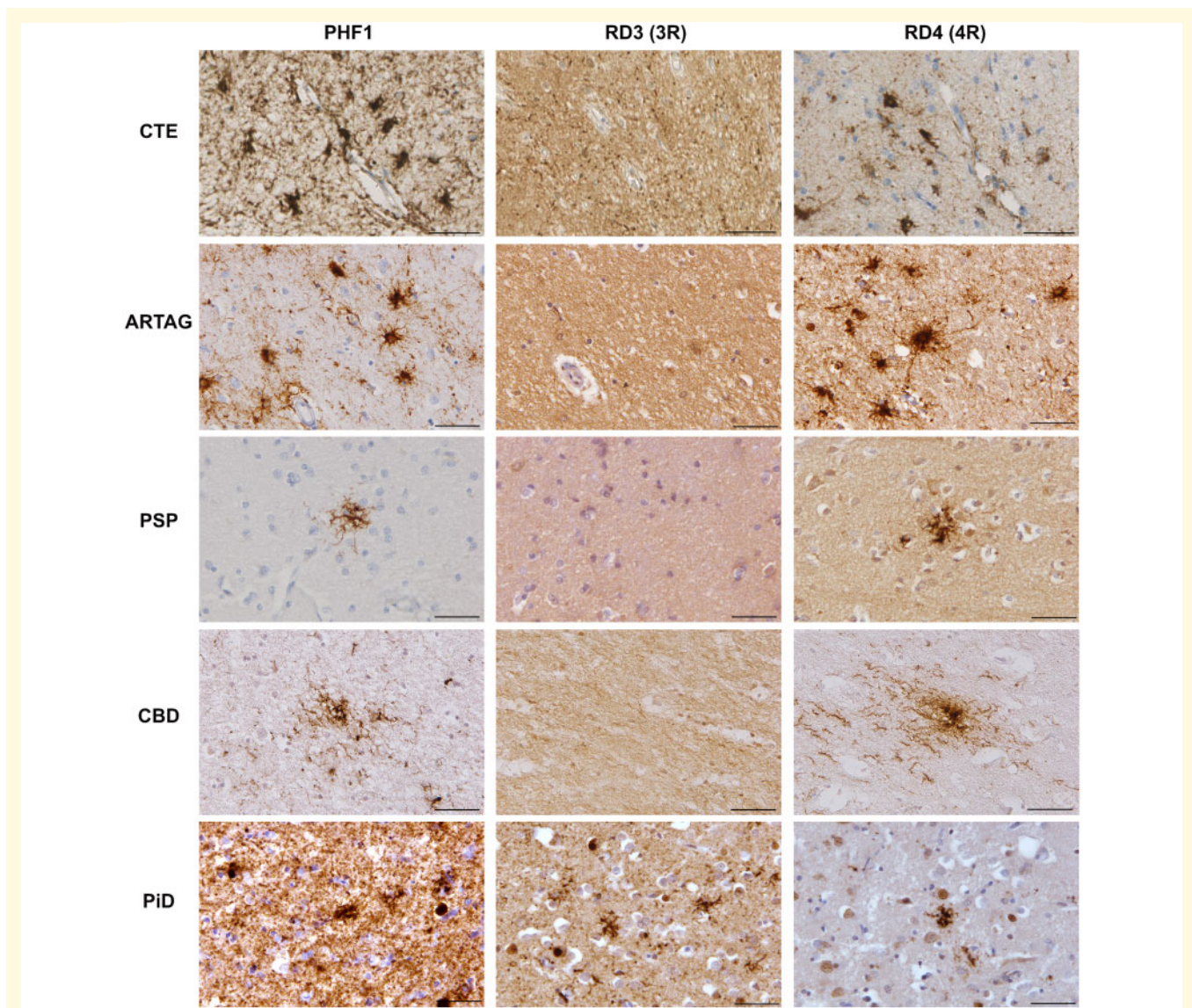


Figure 3 3R versus 4R tau immunoreactivity within the astrocytic pathologies of CTE, ARTAG, PSP, CBD and Pick's disease. Representative examples of serial sections showing immunoreactivity specific for 4R tau, but not 3R tau, in the TSAs in cases of both CTE (Case 6) and ARTAG (Case 21), tufted astrocytes in a case with PSP (Case 35) and astrocytic plaques within a case of CBD (Case 41). The ramified astrocytes observed in just one case of Pick's disease (PiD; Case 29) were immunoreactive for 3R and 4R tau. Scale bars = 50 μ m.

et al., 2014), and displayed immunoreactivity for antibodies recognizing phosphorylation at residues S202 (CP13), S212/T214 (AT100), S262 (p262), and S396/S404 (PHF1). Notably, in one case of Pick's disease, Pick bodies appeared weakly immunoreactive for p262 (Fig. 4). In contrast, neurons were otherwise negative for tau p262 in all other five Pick's disease cases. Notably, previous work has demonstrated conflicting results with regard to pS262 immunoreactivity in Pick's disease (Probst *et al.*, 1996; Delacourte *et al.*, 1998; Ferrer *et al.*, 2002; Zhukareva *et al.*, 2002; Irwin *et al.*, 2016; Falcon *et al.*, 2018). However, greater immunoreactivity has been reported in ethanol versus formalin fixed tissue (Irwin *et al.*, 2016), consistent with our observations. Moreover, the intensity of pS262 immunoreactivity was

reported as greater in cases with 4R tau inclusions (Zhukareva *et al.*, 2002).

Neurofibrillary tangles, but not astrocytes, in CTE show a similar conformational profile to those in Alzheimer's disease

Recently developed conformation-selective antibodies, GT-7 and GT-38, have been shown to detect a conformation-dependent epitope present in tau within the inclusions of Alzheimer's disease requiring both 3R and 4R tau, but not 3R or 4R tau only inclusions of other primary tauopathies,

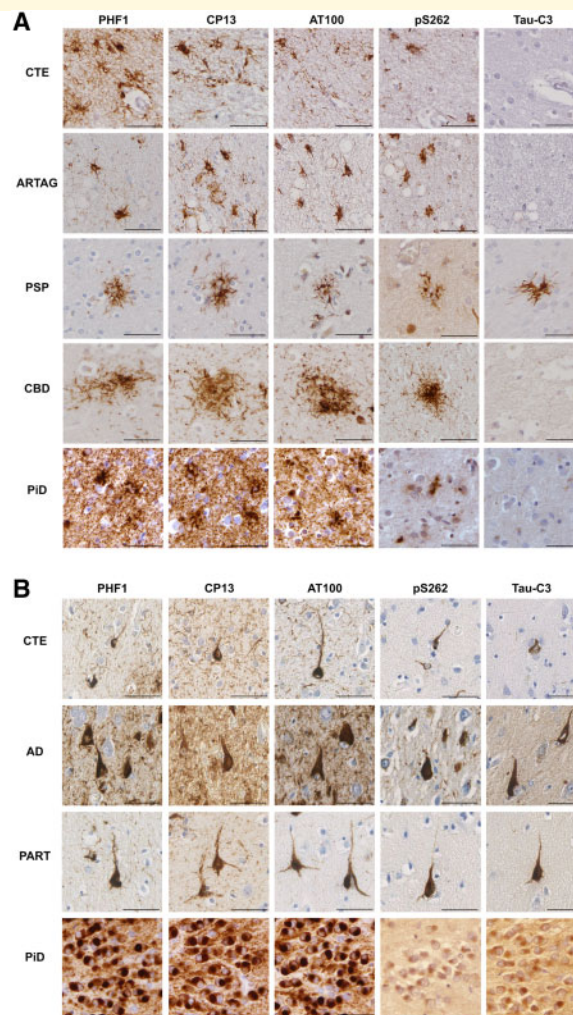


Figure 4 Post-translational modifications of tau within astrocytes and neurons in CTE, Alzheimer's disease, PART, ARTAG, PSP, CBD and Pick's disease. **(A)** Representative examples of serial sections showing the TSAs of CTE (Case 3) and ARTAG (Case 22), astrocytic plaques of CBD (Case 41) and ramified astrocytes in a case of Pick's disease (PiD; Case 29) all displayed robust immunoreactivity to phospho-epitope antibodies PHF1, CP13, AT100 and pS262, but not Tau-C3 (truncation at D421). In contrast with all other astrocytic tau pathologies examined, the tufted astrocytes of PSP in a subset of cases (Case 35 pictured here) were labelled with Tau-C3, indicative of truncation as described. **(B)** Representative examples of serial sections showing the neurofibrillary tangles of CTE (Case 5), Alzheimer's disease (AD; Case 28) and PART (Case 15) with immunoreactivity to the phospho-epitope antibodies PHF1, CP13, AT100 and pS262, as well as Tau-C3 indicating truncation at D421. In addition, while Pick bodies (Case 29) also demonstrated the same post-translational modifications, immunoreactivity to p262 and Tau-C3 was noted in just one case as shown here within the dentate granule cells, where immunoreactivity was notably less robust. Scale bars = 50 μ m.

with both antibodies labelling Alzheimer's disease-tau in a phosphorylation-independent manner (Gibbons *et al.*, 2018, 2019). In CTE neuropathology, both GT-7 and GT-38 showed moderate to strong labelling of NFTs at the depths of sulci, consistent with Alzheimer's disease (Fig. 5 and Table 2). However, in contrast with NFTs, astrocytes within CTE were negative for GT-7 or GT-38 in all but four cases in which there was occasional and minimal positivity to one or the other antibody (Table 2). These cells did not differ in their morphology, and no notable differences in fixation, clinical history or anatomic distribution distinguished them from the rest of the cohort. Double immunofluorescence

labelling in a subset of cases confirmed neuron-specific co-localization of GT-38 with MAP-2, and an absence of co-localization with GFAP (Supplementary Figs 1 and 3).

As anticipated, NFTs in non-injured Alzheimer's disease and PART control cases demonstrated robust immunoreactivity for both antibodies (Fig. 5 and Table 2), while Pick bodies and the glial profiles of ARTAG, CBD and PSP were negative for both GT-7 and GT-38 (Fig. 5). Notably, the ramified astrocytes within the single Pick's disease case that were immunoreactive for both 3R and 4R tau, were not immunoreactive for GT-7 or GT-38. Among TBI cases with co-morbid diagnoses of FTLT-tau, both GT-7 and GT-38

Table 3 Tau phenotype semiquantitative scores: post-translational modifications

Case	PHF-1		CPI3		AT100		pS262		Tau-C3	
	TSA	NFT	TSA	NFT	TSA	NFT	TSA	NFT	TSA	NFT
1 CTE-NC	+++	+++	+++	+++	+	+++	+++	+++	–	+
2 CTE-NC	+++	+++	+++	+++	+++	+++	+++	+++	–	+
3 CTE-NC	+++	+++	+++	+++	+++	+++	+++	+++	–	–
4 CTE-NC	+++	+++	+++	+++	++	++	+++	+++	–	++
5 CTE-NC	+++	+++	+++	+++	+++	+++	+	+++	–	+
6 CTE-NC	+++	+++	+++	+++	+++	+++	++	+++	–	+
7 CTE-NC	+++	+++	+++	+++	+++	+++	+	+++	–	+
8 CTE-NC	+++	+++	+++	+++	+++	+++	++	+++	–	+
9 CTE-NC	+++	+++	+++	+++	+++	+++	–	+	x	x
10 CTE-NC	+++	+++	+++	+++	+++	+++	++	+++	+	++
11 CTE-NC	+++	+++	+++	+++	+++	+++	–	+++	–	+
12 CTE-NC	+++	+++	+++	+++	++	+++	–	+++	–	+
13 CTE-NC	+++	+++	+++	+++	+++	+++	++	+++	–	++
14 CTE-NC	+++	+++	+++	+++	+++	+++	+++	+++	–	+
15 ARTAG, PART	+++	+++	+++	+++	+++	+++	+	++	+	++
16 ARTAG, PART	+++	+++	+++	+++	+++	+++	++	+++	–	+++
17 ARTAG, PART	+++	+++	+++	+++	+++	+++	+	+++	–	+
18 PART	N/A	+++	N/A	+++	N/A	++	N/A	++	N/A	–
19 ARTAG, low ADNC	+++	+++	+++	+++	+++	+++	+++	+++	–	+
20 ARTAG, low ADNC	+++	+++	+++	+++	+++	+++	+++	+++	–	++
21 ARTAG, low ADNC	+++	+++	+++	+++	+++	+++	++	+++	–	–
22 ARTAG, low ADNC	+++	+++	+++	+++	+++	+++	+++	+++	–	–
23 AD		+++		+++		++		+++		++
24 AD		+++		+++		+++		+++		++
25 AD		+++		+++		+++		+++		+
26 AD		+++		+++		+++		+++		+
27 AD		+++		+++		+++		+++		++
28 AD		+++		+++		+++		+++		+++
	Pick bodies	Ramified astro	Pick bodies	Ramified astro	Pick bodies	Ramified astro	Pick bodies	Ramified astro	Pick bodies	Ramified astro
29 PiD	+++	+++	+++	+++	+++	+++	++	+	++	–
30 PiD	+++	N/A	+++	N/A	+++	N/A	–	N/A	++	N/A
31 PiD	+++	+++	+++	+++	+++	+++	–	–	–	–
32 PiD	+++	N/A	+++	N/A	+++	N/A	–	N/A	–	N/A
33 PiD	+++	N/A	+++	N/A	+++	N/A	–	N/A	–	N/A
34 PiD	+++	+++	+++	+++	+++	+++	–	–	–	–
	Tufted astrocytes		Tufted astrocytes		Tufted astrocytes		Tufted astrocytes		Tufted astrocytes	
35 PSP	+++		+++		++		++		+	
36 PSP	+++		+++		+++		+		+++	
37 PSP	+++		+++		+++		+		–	
38 PSP	+++		+++		+++		+		–	
39 PSP	+++		+++		++		–		–	
40 PSP	+++		+++		+++		++		–	
	Astrocytic plaques		Astrocytic plaques		Astrocytic plaques		Astrocytic plaques		Astrocytic plaques	
41 CBD	+++		+++		+		+		–	
42 CBD	+++		+++		+++		–		–	
43 CBD	+++		+++		++		–		–	
44 CBD	+++		+++		++		++		–	
45 CBD	+++		+++		+++		+		–	
46 CBD	+++		+++		+++		+++		–	

Semi-quantitative scoring: – = absent, or nearly absent ($\leq 5\%$ concordance with extent of PHF1); + = minimal (5–30% concordance); ++ = moderate (30–70% concordance); +++ = extensive ($> 70\%$ concordance); x = no immunoreactivity.

AD = Alzheimer's disease; ADNC = Alzheimer's disease neuropathological change; CTE-NC = chronic traumatic encephalopathy neuropathological change; N/A = not applicable; PiD = Pick's disease.

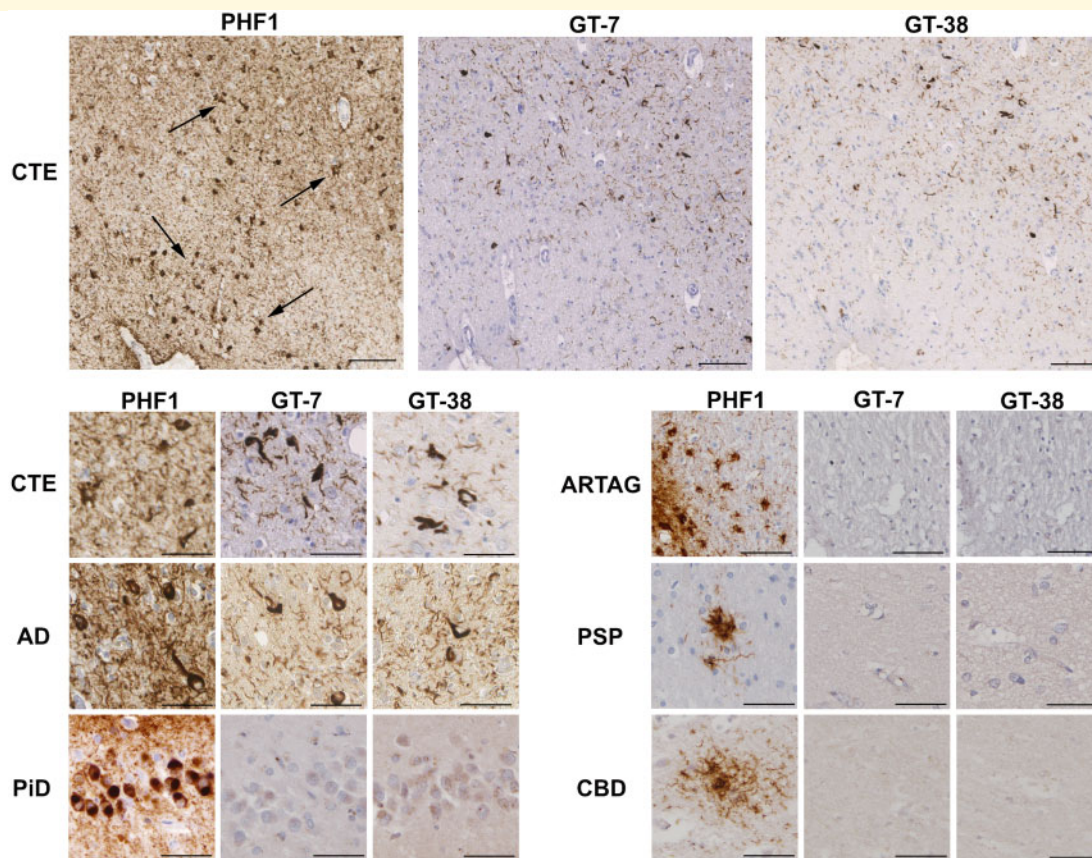


Figure 5 Neurofibrillary tangles, but not astrocytes, are immunoreactive for antibodies that detect a conformation-dependent epitope of tau in Alzheimer's disease. PHF1 immunohistochemistry revealed sulcal depth astrocytic and neuronal tau pathology in CTE (Case 8), including prominent clusters of perivascular and subpial TSAs (top left; black arrows). However, GT-7 and GT-38 antibodies labelled NFTs in CTE, but not TSAs, on serial tissue sections (top middle and top right). GT-7 and GT-38 reliably detected NFTs in Alzheimer's disease (Case 28), but failed to label the characteristic pathologies of ARTAG (Case 19), Pick's disease (PiD; Case 29), PSP (Case 35) or CBD (Case 41). Scale bars = 100 μ m in CTE (low power); = 50 μ m in CTE (high power), Alzheimer's disease (AD), Pick's disease, ARTAG, PSP and CBD.

antibodies stained the characteristic NFTs of CTE but not the adjacent FTLT-tau disease-associated pathologies within the same tissue section.

Discussion

Here, we performed immunohistochemical characterization of tau phenotypes within the cellular constituents of CTE in patients with known exposure to repetitive mild TBI or moderate/severe TBI. Intriguingly, it was found that the tau species within NFTs and immediately adjacent astrocytes in CTE are phenotypically distinct. Specifically, the NFTs of CTE displayed an immunophenotype that mirrored that seen in Alzheimer's disease or PART, including being both 3R and 4R immunoreactive and positive for antibodies previously shown to bind to a conformation-dependent epitope present within the tau inclusions of Alzheimer's disease. In contrast, the astroglial component of CTE was solely 4R immunoreactive and without evidence of Alzheimer's disease conformation, which was indistinguishable from ARTAG.

Notably, while the neuronal and glial pathologies of CTE showed similarity with those of Alzheimer's disease, PART and ARTAG, they were distinct from those of the primary FTLT tauopathies of Pick's disease, CBD and PSP. As such, our data suggest that while pattern and distribution of involvement might be distinct, the tau pathologies of CTE show considerable overlap with the immunoreactivity profiles of both ageing-related tau pathologies and Alzheimer's disease.

Typically, we observed tau immunoreactive astrocytes within CTE to have the characteristic morphology of TSA, in keeping with those encountered in ARTAG, in both the non-injured control material studied here and in multiple other reports (Lopez-Gonzalez *et al.*, 2013; Kovacs *et al.*, 2016, 2017a, 2018a). Notably, TSAs have long been described in material from patients with and without concomitant neurodegenerative disease. In particular, their appearance in association with increased age led to the recognition of the specific entity recently defined as ARTAG, wherein TSAs are described in subpial, subependymal and perivascular distributions (Ikeda *et al.*, 1995, 1998; Ikeda,

1996; Schultz *et al.*, 2004; Lace *et al.*, 2012; Kovacs *et al.*, 2013, 2016). While the clinical significance of ARTAG has yet to be fully explored (Kovacs *et al.*, 2017b), the morphological resemblance between astrocytes immunoreactive for phosphorylated tau in both CTE and ARTAG raises the possibility that these entities share common pathogenic mechanisms (Kovacs *et al.*, 2016, 2017b, 2018a; Liu *et al.*, 2016; Goldfinger *et al.*, 2018; Forrest *et al.*, 2019). Our data demonstrate that the astrocytes of CTE and ARTAG not only share similar morphologies, but display indistinguishable tau immunophenotypes with respect to the panel of antibodies applied. Specifically, consistent with previous reports characterizing ARTAG (Kovacs *et al.*, 2016; Ferrer *et al.*, 2018) and one limited description in CTE (McKee *et al.*, 2013), TSAs in CTE were typically 4R tau-immunoreactive/3R tau-negative, with only very occasional cells displaying immunoreactivity for 3R tau in a subset of cases. Further, astrocytes in CTE demonstrated a profile of immunoreactivity for multiple phospho-epitopes of tau consistent with descriptions of ARTAG here and previously (Lopez-Gonzalez *et al.*, 2013; Ferrer *et al.*, 2018).

Beyond this astroglial pathology, the regionally co-existing NFTs in CTE appeared morphologically and phenotypically consistent with those of Alzheimer's disease and PART, composed of both 3R and 4R tau, and immunoreactive for tau hyperphosphorylated at multiple phospho-epitopes, including S202, S212/T214, S262, and S396/S404. Notably, characterization by others has highlighted additional shared features of tau including phosphatase-activating domain exposed conformation (TNT1 antibody), tau oligomers (TOC1), and truncation at D421 (Tau-C3) (Kanaan *et al.*, 2016). Here we demonstrate that a subset of NFTs in Alzheimer's disease, PART and CTE demonstrate caspase-mediated truncation at D421. Interestingly, Tau-C3 positivity has previously been reported as relatively sparse in CTE neuropathology when compared with Alzheimer's disease (Kanaan *et al.*, 2016). However, the diminished total burden of Tau-C3 immunoreactivity may reflect a virtual absence of immunoreactivity in TSA, as observed here. Truncation of tau at D421 has been reported as an early event in the evolution of NFTs in Alzheimer's disease (Gamblin *et al.*, 2003; Rissman *et al.*, 2004; Cotman *et al.*, 2005), although it may not be essential for filament formation (Delobel *et al.*, 2008). Moreover, experimental data indicate caspase-mediated truncation may promote polymerization and seeding of full-length tau (Abraham *et al.*, 2000; Berry *et al.*, 2003; Gamblin *et al.*, 2003; Rissman *et al.*, 2004), as well as contribute to neurotoxicity via apoptosis (Fasulo *et al.*, 2000, 2005; Chung *et al.*, 2001). As such, the relative absence of Tau-C3 immunoreactivity in TSAs of CTE may have implications as to the potential pathological nature of tau immunoreactive TSAs in CTE.

The astrocytic tau pathologies of both CTE and ARTAG also failed to display immunoreactivity for the recently developed tau antibodies, GT-7 and GT-38, previously demonstrated to detect a conformation-dependent epitope of tau in Alzheimer's disease in a phosphorylation-independent

manner (Gibbons *et al.*, 2018, 2019). Furthermore, these antibodies failed to label pathologies composed of tau with either 3R or 4R isoforms only (Gibbons *et al.*, 2018, 2019). Thus, the observation that both GT-7 and GT-38 bind to NFTs in CTE, but not sulcal TSAs, further supports the observation that astrocytic tau is 4R only and differs from that of the adjacent NFTs. Indeed, while the characteristic pathologies of FTLD-tau were not immunoreactive for either GT-7 or GT-38, as previously characterized in detail (Gibbons *et al.*, 2018, 2019), the NFTs of CTE, Alzheimer's disease and PART were consistently immunoreactive for both.

The panel of antibodies examined herein were selected for their previous extensive characterization and reported differences across a range of neurodegenerative diseases. As no immunolabel examined thus far can morphologically or phenotypically differentiate individual cells in CTE from those of other established tauopathies, the distinguishing features of this pathology remain dependent on the overall pattern and distribution of pathology when using immunohistochemistry (McKee *et al.*, 2016). Notably, this panel is not exhaustive, and the use of immunohistochemistry to explore additional, and potentially distinguishing tau phenotypes, including ubiquitination and acetylation, will be important. Notably, a recent study based on cryo-electron microscopy (cryo-EM) analysis of temporal lobe tissue from three cases of known CTE suggests that the tau filament structure of CTE is distinct from that of Alzheimer's disease (Falcon *et al.*, 2019). However, it is noteworthy that all three cases demonstrated clinical or neuropathological evidence of other neurodegenerative disease, including Parkinson's disease, FTLD and/or motor neuron disease, in one instance associated with *C9orf72* mutation (Falcon *et al.*, 2019). Given the diversity and heterogeneity of comorbid pathologies reported with CTE here and elsewhere (Mez *et al.*, 2017; Lee *et al.*, 2019), it will be important to extend cryo-EM studies to a wider range of trauma-associated cases. Nonetheless, these data suggest that while tau in CTE might differ from that of other established neurodegenerative disease, in particular Alzheimer's disease, this may only be detectable by means beyond established immunohistochemical or biochemical approaches.

There is increasing recognition that mixed, often multiple pathologies might co-exist in patients with neurodegenerative disease, including in those with TBI-related neurodegeneration where CTE might serve as the primary pathology driving disease or as a co-morbidity in context of an alternate diagnosis (Lee *et al.*, 2019). Consistent with this, most cases with CTE examined here also met criteria for other tauopathies, including Alzheimer's disease, PSP, and CBD, as previously reported (Lee *et al.*, 2019). Notably, the morphologies and immunophenotype of CTE were highly consistent across cases, regardless of the presence or extent of co-morbid disease. Moreover, CTE was consistent in phenotype, regardless of the nature of TBI exposure, including in three cases with a remote history of single moderate or severe TBI. These cases add to the limited number of described

cases of single TBI associated with CTE neuropathologies, supporting the assertion that it is exposure to TBI rather than severity or number of injuries that serves as the primary risk factor for CTE (Smith *et al.*, 2013, 2019; Maroon *et al.*, 2015).

Collectively, these data indicate the co-existence of distinct tau phenotypes within neurons and astrocytes contributing to CTE neuropathology. Moreover, immunohistochemical observations were notably consistent across all cases examined, composed of diverse TBI exposure histories. While criteria for delineating the extent or stage of disease in CTE have yet to be adequately defined, future explorations of tau immunophenotype in association with disease progression will be of importance to examine.

While the mechanisms driving tau pathology following TBI exposure remain poorly understood, it is possible that the differential neuronal and astrocytic components reflect mechanistically independent pathological processes. Curiously, TSA in both ARTAG and CTE are frequently observed at brain parenchyma–fluid interfaces, including subpial, subependymal and perivascular regions (Ikeda *et al.*, 1995; Geddes *et al.*, 1996, 1999; Kovacs *et al.*, 2016, 2018a; McKee *et al.*, 2016). Moreover, regional correlation of ARTAG with astrocytic expression of connexin-43 and aquaporin 4 has led to speculation that blood–brain barrier dysfunction may be of pathological significance to the development of this pathology (Kovacs *et al.*, 2018b). While blood–brain barrier dysfunction is increasingly recognized as an important acute pathology of TBI, even following concussion (Weissberg *et al.*, 2014; Johnson *et al.*, 2018), recent data indicate blood–brain barrier permeability may persist chronically in some individuals after severe TBI (Hay *et al.*, 2015) and has also been described in cases of CTE (Doherty *et al.*, 2016). However, a potential mechanistic relationship between blood–brain barrier dysfunction and pathological astrocytic tau accumulation remains unexplored.

Here we provide new insights into the nature of tau in CTE neuropathological change directly within the context of other neurodegenerative pathologies. Moreover, these data highlight the potential challenges in distinguishing trauma-associated tau pathologies from those of other diseases at the individual cell level using immunohistochemistry alone. Nonetheless, morphological and phenotypic similarities between tau in CTE and those of ARTAG suggests the intriguing possibility of shared pathogenic mechanisms.

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Competing interests

The authors report no competing interests.

Supplementary material

Supplementary material is available at *Brain* online.

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