

Table 1: Clinical demographic information of cases used in the study

Case	Sex	Age (yr)	BS: Tau	Thal phase	BS:LB	ARTAG	PMD (hr)
CN	Female	82	1	0	0	-	7.45
CN	Female	89	1	0	0	-	13
	2F	85.5±1.75					10.2±1.3
PSP	Male	77	2	0	3	+	5.4
PSP	Female	84	3	3	0	+	8.4
	1M/1F	80.5±1.75					6.9±0.75

Note: Average of demographic values are presented as mean ± standard error of mean (SEM, bold values); BS: Tau, Braak staging for Tau positivity; BS:LB, Braak staging for Lewy body; Thal phase amyloid staging where 0= no amyloid and 5= extensive amyloid spread; ARTAG, age related tau astrogliaopathy; PMD, postmortem delay; yr, years; hr, hours.

Table 2: Histological analyses showing cell specific counting (density) of different astrocytic subtypes in different brain regions of CN and PSP cases (mean \pm SEM)

Brain regions	CN (# cell/field)	PSP (# cell/field)	p-value
Frontal cortex			
<i>Protoplasmic</i>	109.28 \pm 5.924	129.95 \pm 6.104	0.34
<i>Fibrous</i>	19 \pm 1.571	17.49 \pm 0.572	0.88
<i>IL-As</i>	212.50 \pm 11.530	257.50 \pm 9.969	0.20
<i>Vp-As</i>	18.49 \pm 1.515	29.99 \pm 0.901	0.05
<i>Tufted</i>	6.245 \pm 0.596	18 \pm 2.150	0.02
<i>AT8+ve</i>	17 \pm 1.075	107.49 \pm 2.246	0.02
<i>AT8+ve (subtype I)</i>	8.50 \pm 0.649	29 \pm 3.206	0.02
<i>AT8+ve (subtype II)</i>	10.99 \pm 0.467	81 \pm 3.49	0.02
<i>AT8-ve</i>	66.99 \pm 4.990	3.99 \pm .061	0.02
Temporal cortex			
<i>Protoplasmic</i>	68.81 \pm 5.644	24.98 \pm 1.952	0.02
<i>Fibrous</i>	62.49 \pm 5.248	29.75 \pm 1.744	0.02
<i>IL-As</i>	65.75 \pm 1.771	35.24 \pm 1.384	0.02
<i>Vp-As</i>	26 \pm 1.145	73.49 \pm 6.068	0.02
<i>Thorn-shaped</i>	17.49 \pm 7.603	18 \pm 2.215	0.68
<i>Ramified</i>	3.75 \pm 0.76	55.75 \pm 4.71	0.02
<i>Granular/fuzzy</i>	2.5 \pm 0.28	26 \pm 5.10	0.02
<i>AT8+ve</i>	4.24 \pm 0.72	75 \pm 3.80	0.02
<i>AT8+ve (subtype I)</i>		65.50 \pm 4.39	
<i>AT8+ve (subtype II)</i>	0.75 \pm 0.32	7.50 \pm 0.72	0.02
<i>AT8-ve</i>	43.25 \pm 7.29	43.25 \pm 7.29	0.05
<i>As Plaque</i>	8.25 \pm 7.029	63.49 \pm 7.28	0.02
Cerebellum			
<i>Fibrous</i>	71.18 \pm 1.39	10.12 \pm 19.270	0.02
<i>Valate</i>	100.42 \pm 3.13	183.12 \pm 16.93	0.02
<i>Bergmann</i>	144.43 \pm 4.26	201.13 \pm 3.98	0.02
<i>Thorn-shaped</i>	9.74 \pm 0.37	11.75 \pm 1.20	0.85
<i>Ramified</i>	17.24 \pm 0.54	27.76 \pm 0.94	0.02
<i>Astrocytic plaque</i>	36.32 \pm 1.28	50.50 \pm 1.18	0.02
<i>AT8+ve</i>	2.25 \pm 0.72	54.64 \pm 4.94	0.02
<i>AT8+ve (subtype II)</i>		34.55 \pm 4.094	
<i>AT8-ve</i>	80.63 \pm 6.25	5.49 \pm 1.19	0.02
Caudate nucleus			
<i>Protoplasmic</i>	58.77 \pm 2.46	45.48 \pm 2.22	0.11
<i>Fibrous</i>	22.50 \pm 2.25	43.71 \pm 3.13	0.11
<i>Vp-As</i>	0.49 \pm 0.022	3.99 \pm 0.68	0.14
<i>Tufted</i>	1.99 \pm 0.31	33.37 \pm 2.43	0.02
<i>Thorn-shaped</i>	6.76 \pm 0.60	1.38 \pm 0.60	0.34
<i>Ramified</i>	3.98 \pm 0.56	9.5 \pm 0.41	0.02
<i>Granular/fuzzy</i>	2.25 \pm 0.57	4.25 \pm 0.57	0.60
<i>AT8+ve</i>	3.25 \pm 0.54	41.24 \pm 3.02	0.02
<i>AT8+ve (subtype I)</i>	0.99 \pm 0.043	24 \pm 2.54	0.02
<i>AT8+ve (subtype II)</i>		16.49 \pm 1.86	

<i>AT8-ve</i>	29.75±2.73	3.5±1.24	0.02
Putamen			
<i>Protoplasmic</i>	16.51±1.28	24.24±3.34	0.51
<i>Fibrous</i>	18.25±2.09	18±2.57	0.88
<i>Tufted</i>	1.25±0.32	8.99±0.88	0.02
<i>Thorn-shaped</i>	7.75±0.72	14.74±1.84	0.34
<i>Ramified</i>	5.99±1.27	5.99±0.40	>0.9
<i>Granular/fuzzy</i>		9.25±0.60	
<i>AT8+ve</i>		66.75±10.6	
<i>AT8+ve (subtype I)</i>		36.75±7.78	
<i>AT8+ve (subtype II)</i>		27.24±3.38	
<i>AT8-ve</i>	83.25±4.77	1.25±0.54	0.02
<i>As Plaque</i>	0.75±0.21	12.5±2.99	0.02
Medulla Oblongata			
<i>Protoplasmic</i>	77.38±4.58	16±1.31	0.02
<i>Fibrous</i>	67.63±4.58	51.39±5.84	0.48
<i>Tufted</i>	3.75±0.65	17.75±1.87	0.02
<i>Thorn-shaped</i>	15±1.02	6.49±1.53	0.11
<i>Ramified</i>	2.74±0.21	7.23±0.48	0.02
<i>Granular/fuzzy</i>	1.24±0.42	13.49±1.10	0.02
<i>AT8+ve</i>	0.25±0.011	8.74±0.94	0.02
<i>AT8+ve (subtype I)</i>		2.99±0.75	
<i>AT8+ve (subtype II)</i>		3.75±0.89	
<i>AT8-ve</i>	52±2.05		
<i>As Plaque</i>		3.75±0.60	

Note: Counts of astrocytes were expressed as numbers per unit area and exact p-values are reported. IL-As, interlaminar astrocytes; Vp-As, varicose projection astrocytes; AT8+Ve, tau positive astrocytes; AT8+ve (subtype I), Tau infiltrated astrocytic subtype I; AT8+ve (subtype II), Tau infiltrated astrocytic subtype II; AT8-ve, tau negative astrocytes.

Supplementary Table 3: Detailed information regarding antibodies, chemicals and software used in the study

Reagent/ resources	Source	Identifier/ details
Antibodies		
rabbit anti-GFAP	Abcam	Cat # ab194324, RRID: AB_3662092
mouse anti-ICAM-1	Proteintech	Cat# CL594-60299, RRID:AB_2883457
mouse anti-S100B	EMD Milipore	Cat# MAB079-1, RRID:AB_571112
rabbit anti-GSK3B	Cell Signaling Technology	Cat # 9315S, RRID: AB_490890
mouse anti-AT8	Invitrogen	Cat# MN1020, RRID:AB_223647
rabbit anti-IBA-1	Invitrogen	Cat# PA5-27436, RRID:AB_2544912
NeuroTrace	ThermoFisher	Cat # N21479, RRID: AB_2341101
rabbit anti-Claudin 5	Abcam	Cat# ab131259, RRID:AB_11157940
DAPI	Milipore	
Goat anti-rabbit TRITC	Santa Cruz Biotechnology	Cat# sc-2780, RRID: AB_649006
Goat anti-mouse AlexaFluor 555	Thermo Fisher Scientific	Cat# A-21422, RRID: AB_2535844
Goat anti-mouse AlexaFluor 647	Abcam	Cat# ab150115, RRID: AB_2687948
Goat anti-mouse AlexaFluor 488	Thermo Fisher Scientific	Cat# A-11001, RRID: AB_2534069
Donkey anti-rabbit AlexaFluor 488	Thermo Fisher Scientific	Cat# A-21206, RRID: AB_2535792
Chemicals		
Mercury(II) chloride	Milipore	Cat # M1136, CAS: 7487-94-7
Potassium chromate	Milipore	Cat # 12249, CAS: 7789-00-6
Potassium di-chromate	Milipore	Cat # 207802, CAS: 7778-50-9
Ethyl alcohol	Milipore	Cat # 493511, CAS: 64-17-5
Sodium Thiosulfate	Milipore	Cat # 72049, CAS: 7772-98-7
Sodium chloride	Milipore	Cat # S9888, CAS: 7647-14-5
di-sodium hydrogen phosphate	Milipore	Cat # 1.06585, CAS: 7558-79-4
Triton X-100	Milipore	Cat # X100, CAS: 9036-19-5
Tween 20	Milipore	Cat # P1379, CAS: 9005-64-5
Software and algorithms		
Fiji 1.54p	<i>Schneider et al., (1)</i>	https://imagej.net/
Image J 1.53m	<i>Schneider et al., (1)</i>	https://imagej.net/
Imaris 10	Oxford Instrument	https://imaris.oxinst.com/
Inkscape 1.4.2	Inkscape project	https://inkscape.org/release/inkscape-1.4.2/windows/64-bit/msi/dl/
Wind rose plot	Jupiter notebook	https://pypi.org/project/windrose/1.6/
Simple Neurite Tracer	<i>Longhair et al.,(2)</i>	https://imagej.net/plugins/snt/
Neuron Studio	<i>Wearne et al., (3)</i>	https://biii.eu/neuronstudio

NeuroMorphoVis	<i>Abdellah et al., (4)</i>	https://github.com/BlueBrain/NeuroMorphoVis
Blender 3.1.0	Blender development team	https://www.blender.org/
Prism 10.1.1	GraphPad	https://www.graphpad.com/features

Note: Secondary antibodies were selected based on experimental design and spectral compatibility.

References:

1. J. Schindelin *et al.*, Fiji: an open-source platform for biological-image analysis. *Nat Methods* **9**, 676-682 (2012).
2. M. H. Longair, D. A. Baker, J. D. Armstrong, Simple Neurite Tracer: open source software for reconstruction, visualization and analysis of neuronal processes. *Bioinformatics* **27**, 2453-2454 (2011).
3. S. L. Wearne *et al.*, New techniques for imaging, digitization and analysis of three-dimensional neural morphology on multiple scales. *Neuroscience* **136**, 661-680 (2005).
4. M. Abdellah *et al.*, NeuroMorphoVis: a collaborative framework for analysis and visualization of neuronal morphology skeletons reconstructed from microscopy stacks. *Bioinformatics* **34**, i574-i582 (2018).