

Supplementary material – CSF assay methodology

Total tau (T-tau), phosphorylated tau (P-tau) and β -amyloid 1-42 ($A\beta$ 1-42) were analyzed using INNOTEST enzyme-linked immunosorbent assays (ELISAs (Fujirebio Europe N.V., Gent, Belgium). Other markers of amyloid processing were measured using the MSD $A\beta$ Triplex assay (Meso Scale Discovery, Rockville, MD), a multiplexed method in which C-terminally specific antibodies are used selectively to capture $A\beta$ forms ending at amino acids 38, 40 and 42, respectively, which are then quantified using the 6E10 detector antibody. This assay is thus not specific to the 1st amino acid of the $A\beta$ peptides (the epitope of 6E10 lies within amino acids 3 to 8 in the $A\beta$ sequence) and the measured $A\beta$ isoforms are therefore called $A\beta$ X-38, $A\beta$ X-40 and $A\beta$ X-42. Neurofilament light chain (NFL) concentrations were determined using the NF-light method (UmanDiagnostics, Umeå Sweden); YKL-40, also known as chitinase-3-like protein 1 (CHI3L1), was measured using the Human Chitinase 3-like 1 Quantikine ELISA Kit (R&D systems, Minneapolis, MN). Amyloid precursor protein soluble metabolites α and β (sAPP α , sAPP β) were measured using a commercial duplex immunoassay with electrochemiluminescence detection (Meso Scale Discovery, Rockville, MD). Inter-plate co-efficients of variation for internal standards (pooled AD CSF) were: YKL-40: 9.59 %; NFL: 7.72%; sAPP α : 23.03%; sAPP β : 28.56%; $A\beta$ X-38: 5.52%; $A\beta$ X-40: 7.57%; $A\beta$ X-42: 10.17%.

Supplementary Table S1 Diagnostic accuracy of A β 1-42, T-tau, T-tau/A β 1-42 ratio, P-tau and A β X-42/X-40 ratio in test and validation cohorts based on pre-LP diagnostic classification and diagnostic accuracy in the pathologically or genetically defined sub-cohort. AD: Alzheimer's disease; DLB: dementia with Lewy bodies; bvFTD: behavioural variant frontotemporal dementia; PNFA: progressive non-fluent aphasia; SD: Semantic dementia; HC: healthy control.

Diagn ostic Groups compared	Biom arker	Test cohort (used to estimate the cut- point)(data from tables 3 and 4) (n=275)		Validation cohort (using test cohort cut-point) (n=143)		Pathologically or genetically confirmed sub-cohort (n=26)	
		Optim al cut- point at 85% sensitivity	Specif icity	Sensi tivity	Speci ficity	Sensit ivity	Specif icity
AD vs HC	A β X- 42/X-40	<0.06 0	93%	82%	80%		
	A β 1- 42 (pg/mL)	<529. 0	90%	71%	80%		
	T- tau/A β 1-42	>0.64	83%	88%	89%		
	T-tau (pg/mL)	>312. 0	53%	87%	78%		
	P-tau (pg/L)	>48.9	54%	83%	78%		
AD vs non- AD dementia	T- tau/A β 1-42	>0.64	56%	88%	76%	100%	60%
	A β X- 42/X-40	<0.06 0	68%	82%	74%	92%	100%
	T-tau (pg/mL)	>312. 0	51%	87%	53%	94%	60%
	P-tau (pg/L)	>48.9	41%	83%	70%	83%	43%
	A β 1- 42 (pg/mL)	<529. 0	48%	71%	77%	88%	60%
AD vs all HC)	T- tau/A β 1-42	>0.64	63%	88%	76%		
	A β X- 42/X-40	<0.06 0	76%	83%	74%		
	T-tau (pg/mL)	>312. 0	51%	87%	50%		
	A β 1- 42 (pg/mL)	<529. 0	59%	71%	77%		

	P-tau (pg/L)	>48.9	45%	83%	70%		
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