

eTable 1 Follow-up frequency and genetic variants of the study cohort						
	Total (n = 314)	bvFTD (n = 152)	nvPPA (n = 61)	lvPPA (n = 32)	svPPA (n = 31)	ncfPPA (n = 38)
Follow-up 1	154/314	75/152	32/61	13/32	16/31	18/38
Follow-up 2	58/314	34/152	10/61	4/32	3/31	7/38
Follow-up 3	14/314	7/152	3/61	2/32	0/31	2/38
Follow-up 4	1/314	1/152	0/61	0/32	0/31	0/38
C9orf72, n (%)	11/314 (3.5%)	10/152 (6.6%)	0/61 (0.0%)	0/32 (0.0%)	0/31 (0.0%)	1/38 (2.6%)
GRN, n (%)	15/314 (4.8%)	3/152 (2.0%)	5/61 (8.2%)	2/32 (6.3%)	1/31 (3.2%)	4/38 (10.5%)
MAPT, n (%)	2/314 (0.6%)	2/152 (1.3%)	0/61 (0.0%)	0/32 (0.0%)	0/31 (0.0%)	0/38 (0.0%)
Other, n (%)	6/314 (1.9%)	4/152 (2.6%)	0/61 (0.0%)	1/32 (0.0%)	1/31 (3.2%)	0/38 (0.0%)