

Supplemental Table: Baseline interaction effects analysis results for primary model, including NART, gender, race.

	Old, nadir >200				Old, nadir <200			
	β	95% C.I.		p-value	β	95% C.I.		p-value
		Lower	Upper			Lower	Upper	
RAVLT trial 5	0.26	-0.77	1.29	0.62	-0.19	-1.14	0.76	0.69
RAVLT total score	0.41	-3.60	4.4	0.84	0.02	-3.68	3.7	0.99
RAVLTir	0.33	-1.06	1.71	0.64	0.39	-0.89	1.66	0.55
RAVLTdr	0.32	-1.17	1.80	0.68	0.17	-1.20	1.54	0.80
RCFcopy	-0.39	-1.89	1.11	0.61	0.29	-1.09	1.68	0.68
RCFir	-1.48	-4.33	1.37	0.31	1.78	-0.86	4.4	0.19
RCF-DR	-1.84	-4.63	0.95	0.20	2.2	-0.38	4.8	0.094
Digit Symbol	0.61	-4.38	5.6	0.81	0.26	-4.34	4.9	0.91
GPdom	2.3	-7.58	12.2	0.65	-2.95	-12.04	6.1	0.52
GPnon dom	6.4	-5.73	18.5	0.30	0.17	-10.96	11.3	0.98
FAS	-2.54	-7.76	2.7	0.34	-3.57	-8.39	1.25	0.15
Animals	1.11	-1.24	3.5	0.35	0.51	-1.66	2.7	0.65
Trails A	-0.26	-4.44	3.9	0.90	0.13	-3.71	4.0	0.95
Trails B	0.38	-11.82	12.6	0.95	6.6	-4.65	17.8	0.25
Digit Span-F	-0.52	-1.49	0.46	0.30	-0.55	-1.45	0.34	0.23
Digit Span-B	-0.96	-1.98	0.06	0.064	-1.20	-2.15	-0.26	0.012
BNT	-1.34	-3.34	0.65	0.19	-0.60	-2.44	1.24	0.52
Gait	0.79	0.07	1.51	0.031	0.55	-0.12	1.22	0.11
CalCap Choice	14.5	-7.32	36	0.19	5.3	-14.92	26	0.60
CalCap Seq.	10.8	-34.55	56	0.64	-0.89	-42.91	41	0.97