

Supplementary Online Content

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eTable. Sample-size matched inter-biomarker comparisons of goodness of fit in generalized estimating equations models.

This supplementary material has been provided by the authors to give readers additional information about their work.

eTable. Sample-size matched inter-biomarker comparisons of goodness of fit in generalized estimating equations models

CSF vs. PET

	NC	MCI	AD
	(n = 17)	(n = 28)	(n = 8)
CSF model R ²	0.23	0.02	0.28
Age	0.09	0.01	-0.07
Time	-0.06 [*]	0.04	0.49 ^{***}
Baseline	-0.03 [*]	-0.01	0.02
Change	-0.02	-0.03	-0.01
PET model R ²	0.30	0.14	0.66
Age	-0.02	0.01	0.19
Time	-0.07 ^{**}	0.01	0.29 ^{***}
Baseline	-15.67 ^{**}	-11.38 ^{**}	-39.57 [*]
Change	-17.35 [*]	-22.80 [*]	-35.27 ^{**}

CSF vs. MRI

	NC	MCI	AD
	(n = 36)	(n = 53)	(n = 16)
CSF model R ²	0.13	0.10	0.25
Age	0.03	-0.01	-0.07
Time	-0.02	0.06	0.37 ^{**}
Baseline	-0.018 ^{**}	-0.02 [*]	0.08
Change	-0.01	-0.09 [*]	-0.26
MRI model R ²	0.02	0.10	0.18
Age	-0.03	-0.05	0.07
Time	-0.02	0.02	0.31

	Baseline	-5.3×10^{-4}	-2.7×10^{-3}	-3.3×10^{-3}
	Change	-3.1×10^{-3}	-0.01	-0.01
PET vs. MRI				
		NC	MCI	AD
		(n = 102)	(n = 198)	(n = 94)
	PET model R ²	0.08	0.16	0.38
	Age	0.01	0.02	0.15
	Time	-0.03	0.07 ^{***}	0.18 ^{***}
	Baseline	-6.47 ^{***}	-14.80 ^{***}	-31.53 ^{***}
	Change	-4.46	-11.20 ^{**}	-38.79 ^{***}
	MRI model R ²	0.05	0.19	0.21
	Age	0.07	-0.06	-0.25 ^{**}
	Time	-0.05 ^{**}	0.04	0.22 ^{**}
	Baseline	1.8×10^{-4}	-4.3×10^{-3} ^{***}	-5.4×10^{-3} ^{**}
	Change	-0.01	-0.01 ^{**}	-0.01 [*]

Time-varying ADAS-cog scores were modeled by baseline age, time and baseline value and change of biomarker. R-square was the percentage of the outcome variance explained by the model. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.