

**Table A. Aggregated results of cross-validation, using separate clinical features, i.e. no TICF; and relational network.**

ML Model	Train R2		Explained Variance		Negative Mean Absolute Error		Negative Median Absolute Error	
	Mean	StD	Mean	StD	Mean	StD	Mean	StD
SVR-RBF	0.206	0.044	0.230	0.023	-54.923	5.267	-49.310	8.098
SVR-LINEAR	0.972	0.009	0.973	0.008	-5.494	1.009	-5.312	1.515
DTR	0.991	0.004	0.991	0.004	-5.258	1.125	-4.443	1.235
SVR-POLY	0.883	0.020	0.886	0.021	-18.900	1.112	-14.272	3.289