

Table C. Aggregated results of cross-validation, using separate clinical features, i.e. no TICF; without the relational network. NA means that the model didn't converge.

ML Model	Train R2		Explained Variance		Negative Mean Absolute Error		Negative Median Absolute Error	
	Mean	StD	Mean	StD	Mean	StD	Mean	StD
SVR-RBF	NA	NA	NA	NA	NA	NA	NA	NA
SVR-LINEAR	0.866	0.025	0.877	0.029	-14.485	2.359	-12.594	2.025
DTR	0.985	0.013	0.987	0.011	-4.526	2.133	-4.088	1.684
SVR-POLY	0.764	0.073	0.805	0.033	-17.165	1.376	-14.516	1.963