

This file contains the agreement between disorder predictors on a pairwise basis of the per-residue binary scores.

	dynamine-esp_xray	esp_xray-iup_short	iup_short-dynamine
conserved	0.840734012	0.835773983	0.839444041
contained	0.841516292	0.847389175	0.835576098
conflict	0.88944439	0.915477004	0.880929597
overlap	0.837850456	0.849032379	0.842729416