Supplementary materials for ”Assessment of protein assembly prediction in CASP13”

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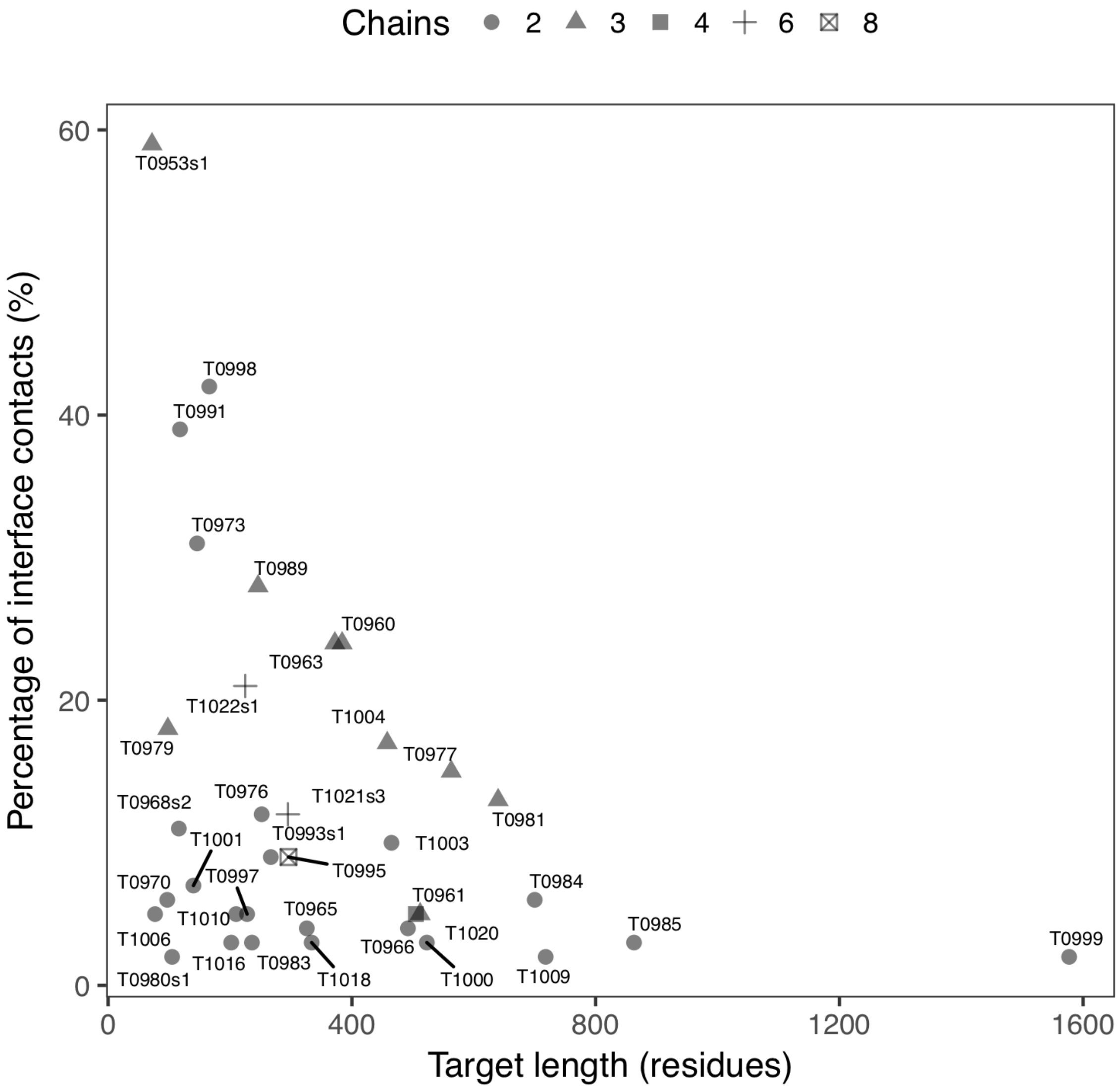
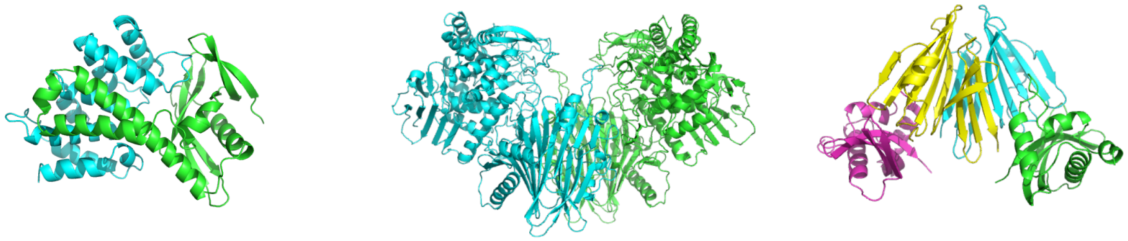


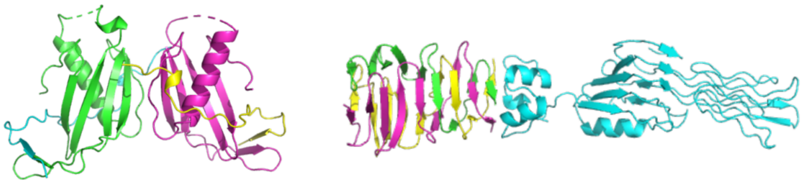
Figure S1: Percentage of homomeric interface contacts in CASP13 targets.



0957: SAXS, X-links, x-links

0985: SAXS (A1 in regular)

0968: SAXS, X-links, x-links



0980: NMR, SAXS

0953: SAXS,SANS, X-links

Images not available:
0981: SAXS, X-links
0999: SAXS, X-links

Figure S2: Data-assisted targets.

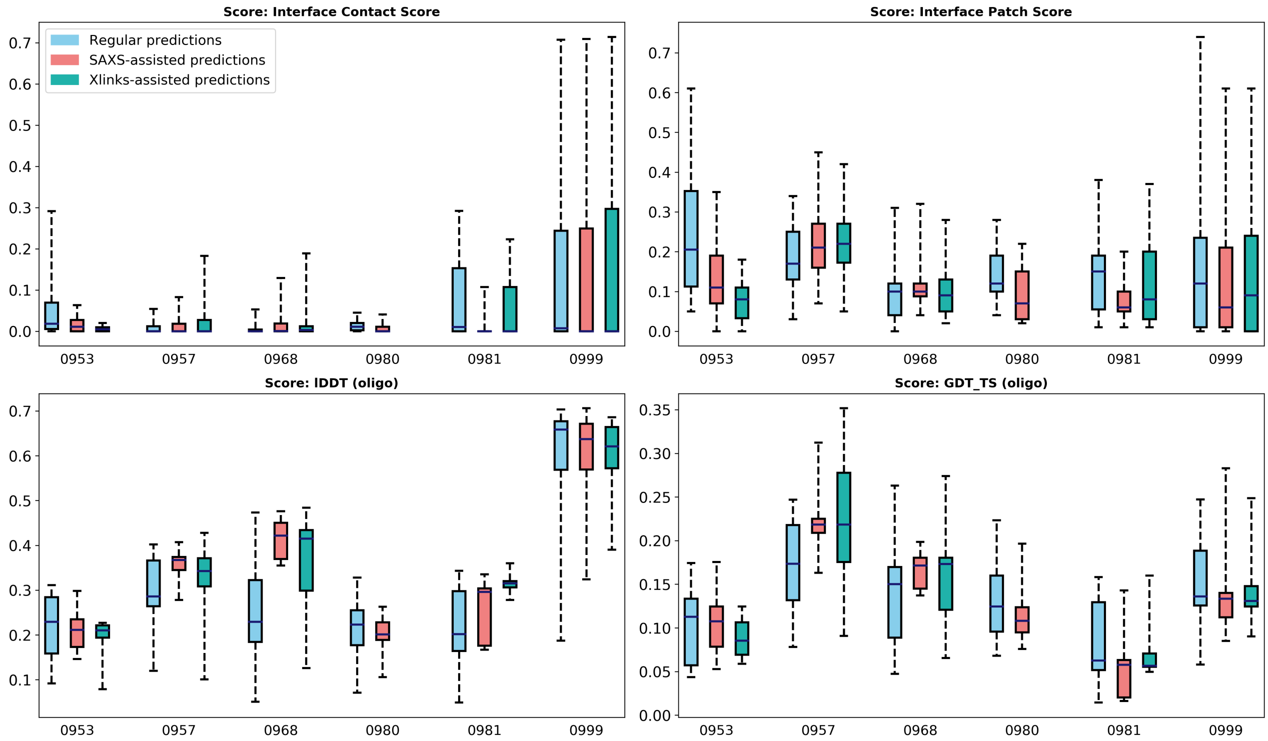


Figure S3: Score distributions for all predictions of data-assisted and the corresponding non-assisted targets. Two types of crosslinks and two types of scattering datasets are merged for the purpose of this figure.

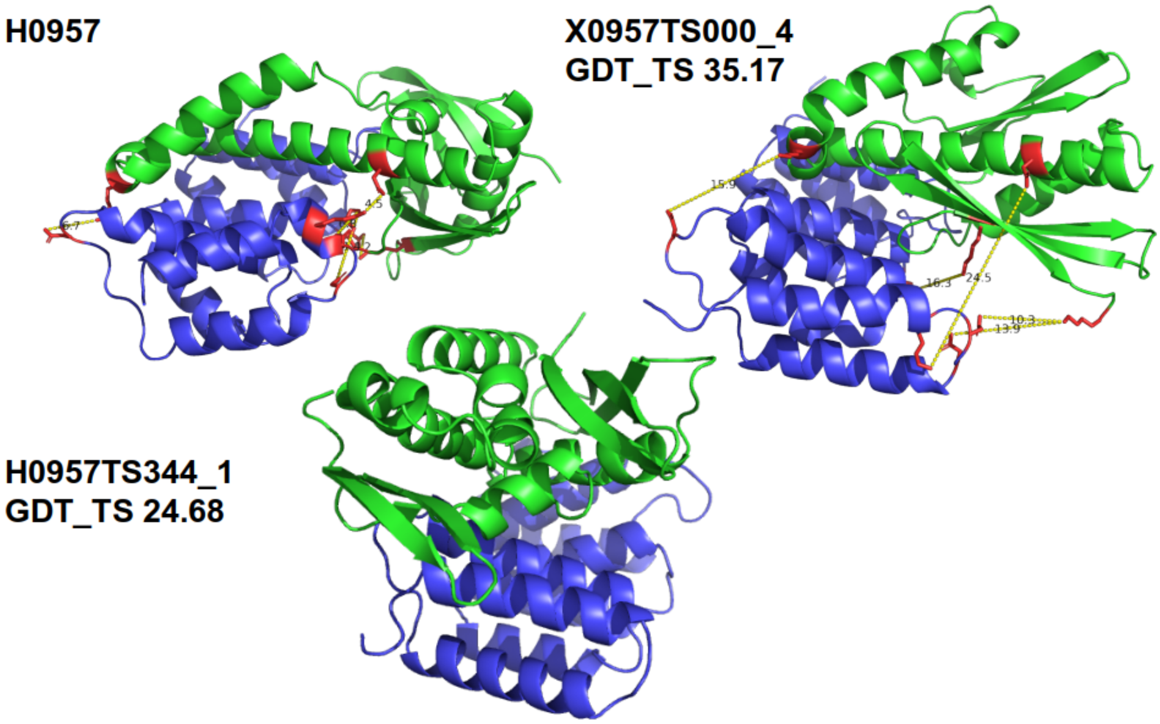


Figure S4: Target X0957 shows improvement across all scores considered due to several fortunate intermolecular crosslinks. Crosslinked residues in the target and the assisted prediction are highlighted in red and connected with a dashed yellow line. Crosslinks between missing residues are not shown. Best regular prediction (bottom) has a significantly lower GDT.