

Table 4.

The performance of the methods on 88 template-based proteins in the CASP dataset

Method	Top1	Top5	Top10
DeepSF	46.59%	73.86%	84.09%
HHSearch	43.20%	61.40%	67.00%
Cons_HH_DeepSF	59.10%	77.30%	85.20%
PSI-BLAST	15.90%	31.80%	47.70%