

Table 5.

The performance of the methods on 95 template-free proteins in the CASP dataset

Method	Top1	Top5	Top10
DeepSF	24.21%	51.58%	70.53%
HHSearch	11.58%	34.74%	44.21%
Cons_HH_DeepSF	23.16%	56.84%	70.53%
PSI-BLAST	8.42%	15.79%	32.63%