

REPORT

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Table 1: Speed tests for swissprot_demo.fasta file (contains first 1k sequences from SwissProt)

Method	CPU	GPU
IUPred	41.44	NA
ProtBert	2015.47	42.91

Table 2: Percentages of disorder/order and secondary structure predictions

Data	Disorder	Helix	Strand	Coil	Total time (CPU)	Total time (GPU)
E. coli	6.3	42.8	16.4	40.8	3.1 min	3.3 min
H. sapiens	22.3	26.9	9.4	63.7	16.1 min	24.5 min
SwissProt	11.8	35.7	14.5	49.8	418.9 min	561.6 min

Table 3: MCC for short and long versions for human proteins

Version	MCC
Short	0.27
Long	0.29