

If you use this service, could you please send us a mail to npsang@ibcp.fr with details about your usage of the NPSA service (tools used, frequency, type of sequence, ..) ? Could you explain what makes this service unique for you ? Could you please add information about your country and your laboratory ? Thanks



Job **SOPMA** (ID: b9492f28380c) is running on **NPS@** server (started on 20250422-090427).

Results will be shown below. **Please wait and don't go back.**

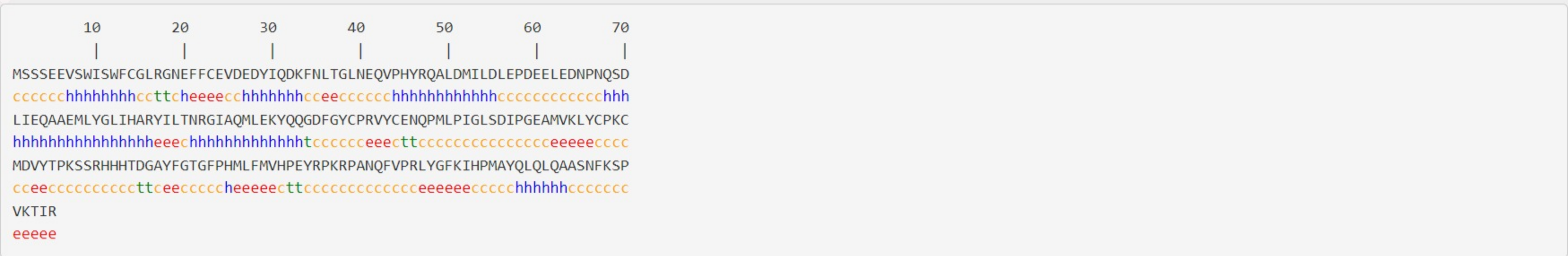
In your publication cite :
NPS@: Network Protein Sequence Analysis
TIBS 2000 March Vol. 25, No 3 [291]:147-150
Combet C., Blanchet C., Geourjon C. and Deléage G.

Running-----

SOPMA result for : CSNK2B_WT

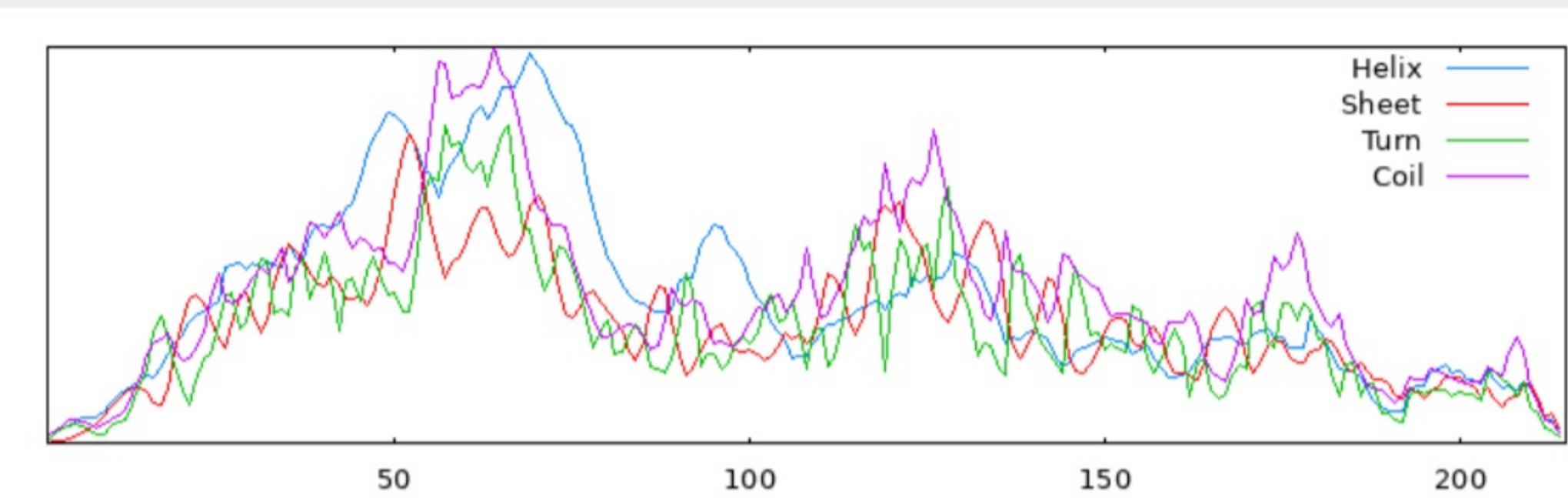
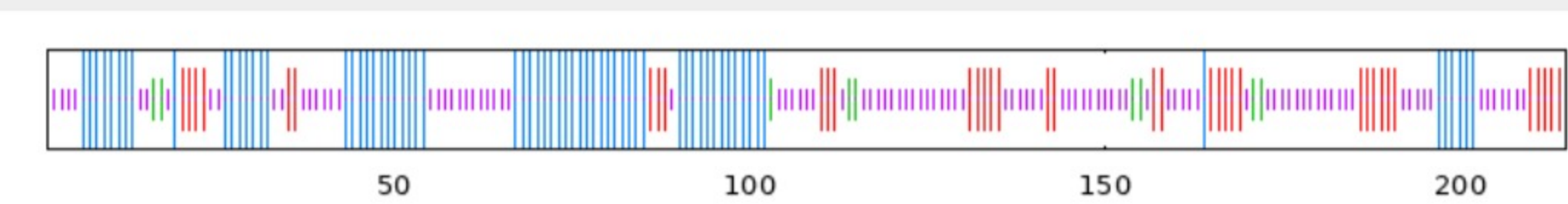
[Abstract](#) Geourjon, C. & Deléage, G., SOPMA: Significant improvement in protein secondary structure prediction by consensus prediction from multiple alignments., Cabios (1995) 11, 681-684

View SOPMA in: [\[AnTheProt \(PC\)](#) , [Download...](#)] [\[HELP\]](#)



Sequence length : 215

SOPMA :			
Alpha helix	(Hh) :	67 is	31.16%
3 ₁₀ helix	(Gg) :	0 is	0.00%
Pi helix	(Ii) :	0 is	0.00%
Beta bridge	(Bb) :	0 is	0.00%
Extended strand	(Ee) :	37 is	17.21%
Beta turn	(Tt) :	9 is	4.19%
Bend region	(Ss) :	0 is	0.00%
Random coil	(Cc) :	102 is	47.44%
Ambiguous states (?)	:	0 is	0.00%
Other states	:	0 is	0.00%



Parameters :
Window width : 17
Similarity threshold : 8
Number of states : 4

Prediction result file (text): [\[SOPMA\]](#)
Intermediate result file (text): [\[BLASTP on NRPROT\]](#) [\[CLUSTALW\]](#)

User : public Last modification time : Tue Apr 22 09:06:47 2025. Current time : Tue Apr 22 09:06:47 2025