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Job GOR4 (ID: c1c56a433320) is running on NPS@ server (started on 20250711-143525).
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-----

GOR4 result for : UNK_309500

Abstract (/NPSA/npsa_references.html#gor4) GOR secondary structure prediction method version IV, J. Garnier, J.-F. Gibrat, B. Robson, Methods in Enzymology, R.F. Doolittle Ed., vol 266, 540-553, (1996)

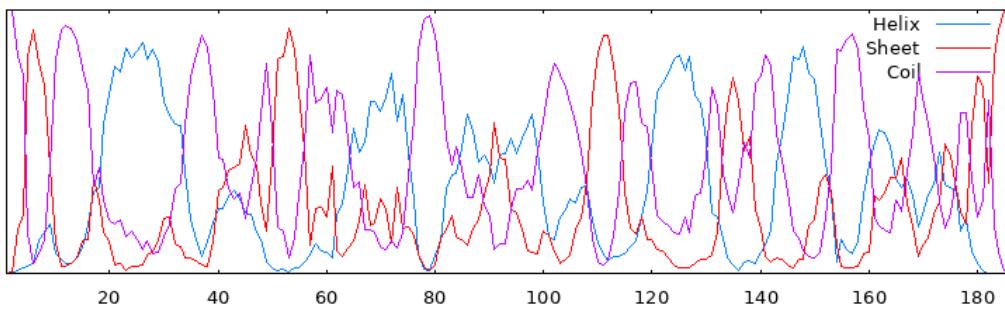
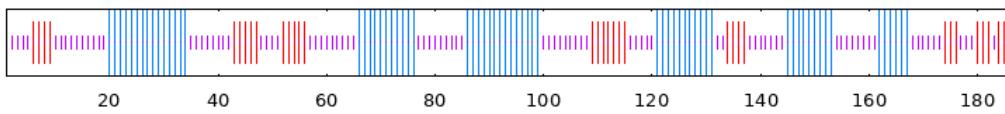
View GOR4 in: [AnTheProt (PC) (http://npsa-pbil.ibcp.fr/cgi-bin/npsa_mpsaview.pl?fn=/tmp/c1c56a433320.gor4.mpsa.mpsa&gui=antheprot), Download... (<http://antheprot-pbil.ibcp.fr>)] [HELP (http://npsa-pbil.ibcp.fr/cgi-bin/npsa_automat.pl?page=/NPSAHELP/npsahlp_locsoft.html)]

10	20	30	40	50	60	70
CSQPLDVILLDGSSFPASYFDEMKSFAKAFISKANIGPRLTQVSVLQYGSITTIDVPWNVPEKAHLL						
ccccccceeecccccccccuhhhhhhhhhhhccccccceeeeeccccceeeeecccccccccuhhh						
SLVDVMQREGGPSQIGDALGFAVRYLTSSEMHGARPAGASKAVVILVTDVSDSVDAADAARSNRVTVFPI						
hhhhhhcccccccccuhhhhhhhhhccccccceeeeeccccchhhhhhhhhccccccc						
GIGDRYDAAQLRILLAGPAGDSNVKLQRIEDLPTMVLGNNSFLHKLC						
ccccchhhhhhhcccccccccuhhhccccccceeeccccceeeccc						

Sequence length : 187

GOR4 :

Alpha helix (Hh) :	66 is 35.29%
β_10 helix (Gg) :	0 is 0.00%
Pi helix (Ii) :	0 is 0.00%
Beta bridge (Bb) :	0 is 0.00%
Extended strand (Ee) :	33 is 17.65%
Beta turn (Tt) :	0 is 0.00%
Bend region (Ss) :	0 is 0.00%
Random coil (Cc) :	88 is 47.06%
Ambiguous states (?) :	0 is 0.00%
Other states :	0 is 0.00%



Prediction result file (text): [GOR4 (/tmp/c1c56a433320.gor4.mpsa)]

User : public Last modification time : Fri Jul 11 14:36:57 2025. Current time : Fri Jul 11 14:36:57 2025



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