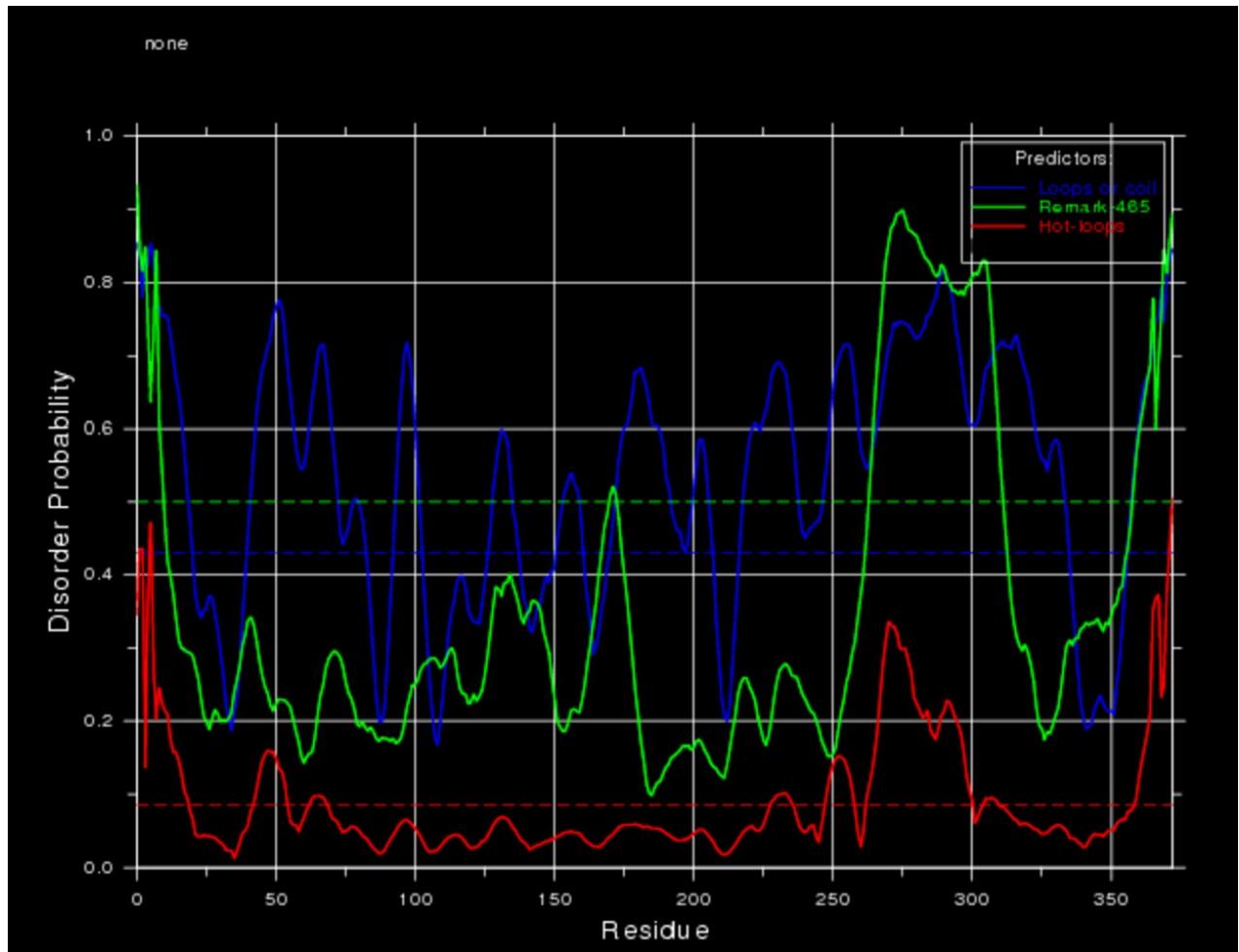


DisEMBL™

Prediction results for Q8I6E4_LEIIN



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Disordered by Loops/coils definition

```
> Q8I6E4_LEIIN_LOOPS 1-20, 41-83, 94-103, 128-138, 152-161, 171-207, 218-335, 357-373
MTASPRAPHQ EHVLGEPTLE glahyirekn vrrilvlvga GASVAAGIPD FRSPDTGIYA NLGKYNLEDP TDAFSLTLLR
EKPeifysia relNLWPGHF QPTavhhfir llqdegrllr cctqnidGLE KAAGVSPEll veahgsfaaa aCIECHTPFS
Ieqnyleams GTVSRCSTCG GIVKPNVVFF GENLPDAFFD ALHHDAPIae lviiigtSMQ VHPFALLPCV VPKSVPVVM
NRERVGGLLF RFPDDPLNTV HEDAVAKEGR SSSSQSRSPS ASPRREEGGT EDSPPSPNEE VEEASTSSSS DGYGQYGDYH
AHPDVCRDVL FRGDCqenvv tlaeylglse alakrmrlsd AAPATAQRAP NET
```

Disordered by Hot-loops definition

```
> Q8I6E4_LEIIN_HOTLOOPS 1-19, 44-55, 249-258, 263-301, 360-373
MTASPRAPHQ EHVLGEPTLe glahyirekn vrrilvlvga gasVAAGIPD FRSPDTgiya nlgkynledp tdafsltllr
ekpeifysia relnlwpghf qptavhhfir llqdegrllr cctqnidgle kaagvspell veahgsfaaa aciechtpfs
ieqnyleams gtvsrcstcg givkpnvvff genlpdaffd alhhdaapiae lviiigtsmq vhpfallpcv vpksvpvvm
nrervgglLF RFPDDPLNtv heDAVAKEGR SSSSQSRSPS ASPRREEGGT EDSPPSPNEE Veeastssss dgygqygdyh
ahpdvcrdvl frgdcqenvv tlaeylglse alakrmrlsD AAPATAQRAP NET
```

Disordered by Remark-465 definition

> Q8I6E4_LEIIN_REM465 1-10, 264-312, 359-373

MTASPRAPHQ ehvlgeptle glahyirekn vrrilvlvga gasvaagipd frspdgtiya nlgkynledp tdafsltllr ekpeifysia relnlwpghf qptavhhfir llqdegrllr cctqnidgle kaagvspell veahgsfaaa aciechtpfs ieqnyleams gtvsr cstcg givkpnnvff genlpdaffd alhhdapiae lviiigtsmq vhpfallpcv vpksvprvvm nrervgllf rfpddplntv hed**AVAKEGR SSSSQSRSPS APRREEGGT EDSPSSPNEE VEEASTSSSS DGyqygdyh ahpdvcrdvl frgdcqenvv tlaeylglse alakrmrlSD AAPATAQRAP NET**

JOB-ID Q8I6E4_LEIIN_164640WgSGcAoLCBIAAH45I88AAAAB

Frames used smooth=8 peak=8 join=4

Thresholds used coils=0.516 rem465=0.6 hot loops=0.1204

Name Q8I6E4_LEIIN

Description none

Title/ID none

Sequence length 373

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