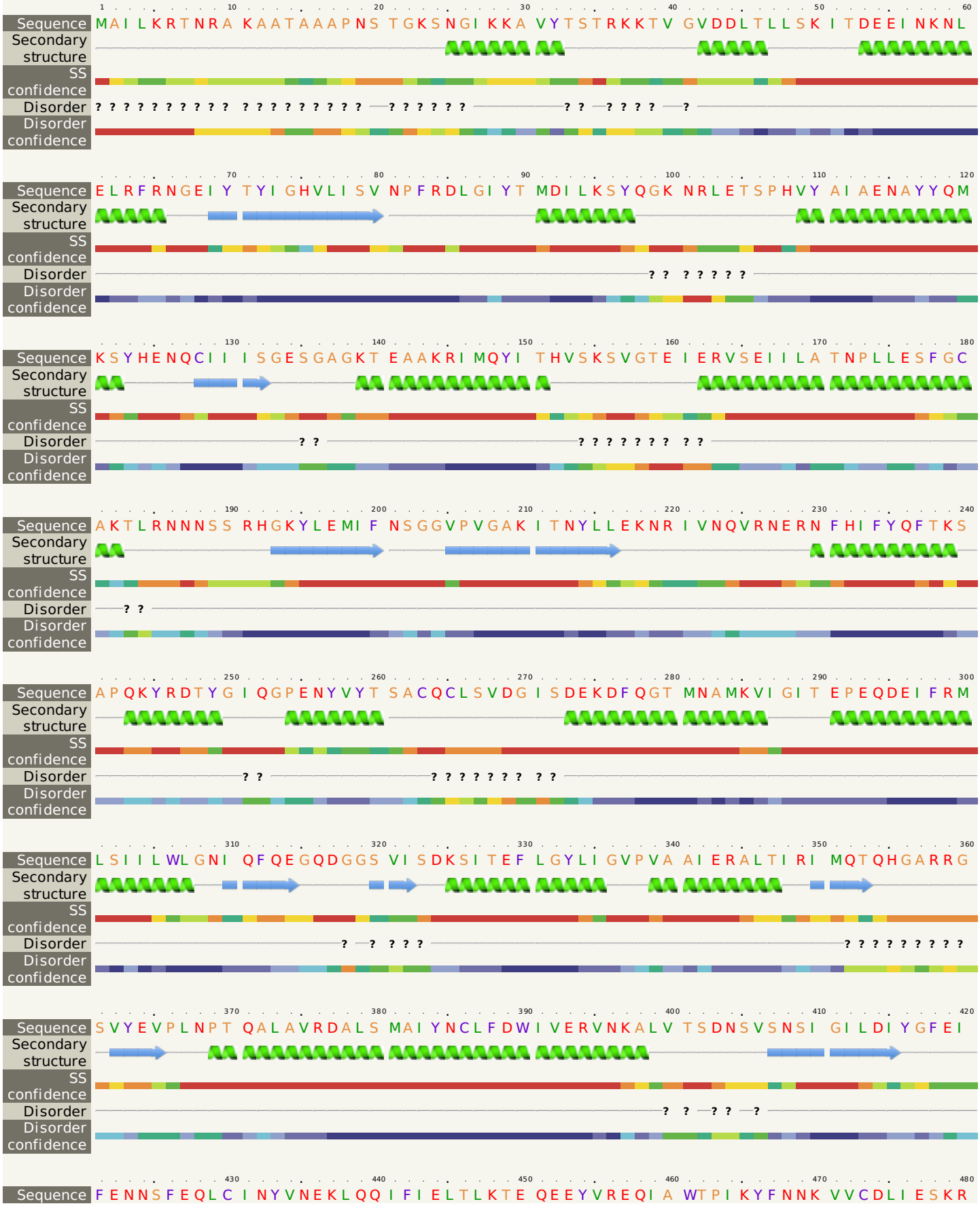
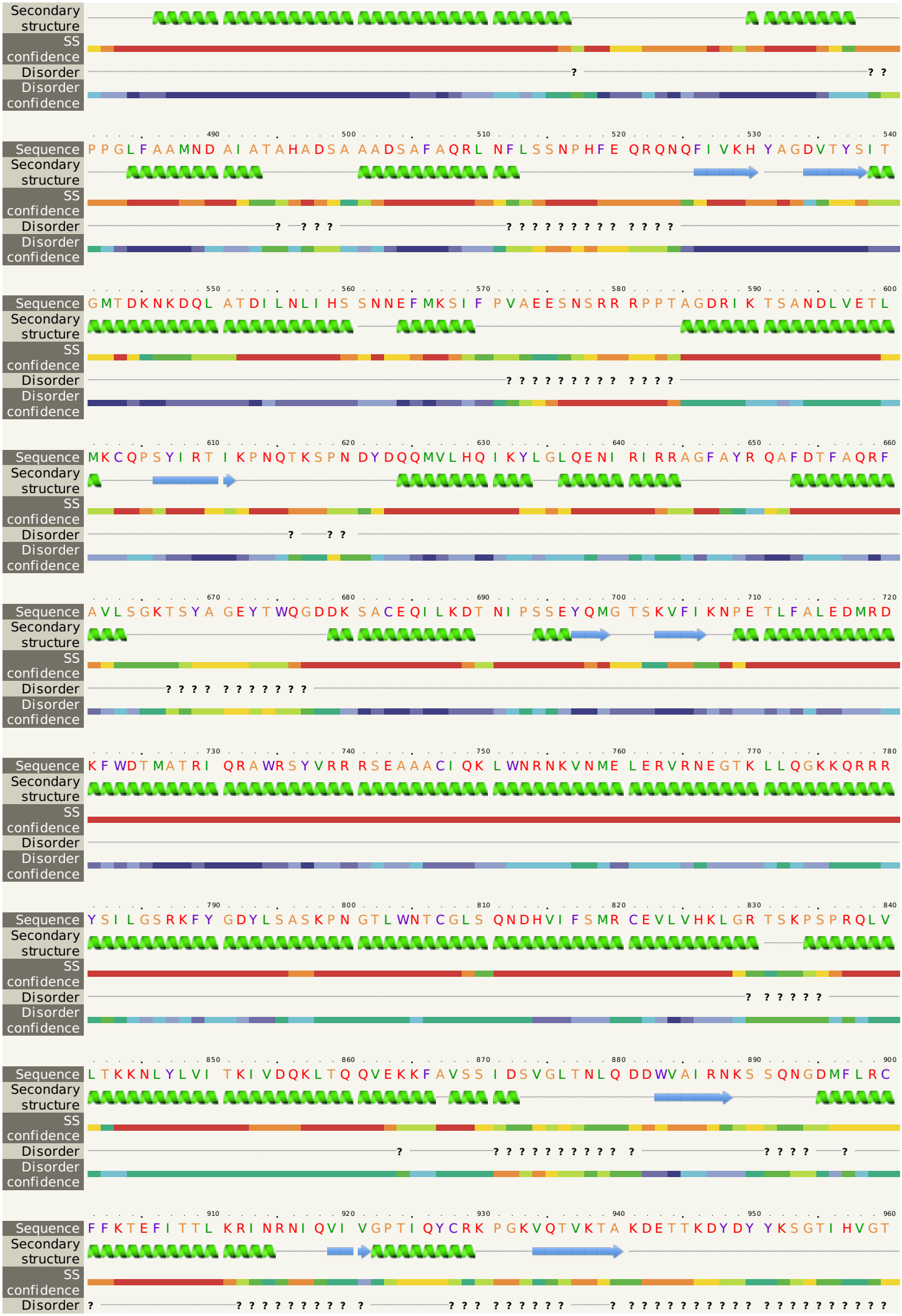
| | |
|---|---|
| Email | l.a.kelley@imperial.ac.uk |
| Description | Q9Y7Z8 |
| Date | Sun Jul 8 12:04:58 BST 2012 |
| Unique Job ID | 0d87b2125ba9b95e |

## Secondary structure and disorder prediction




Secondary structure
SS confidence
Disorder
Disorder confidence
Sequence
PPGLFAAMND AIATAHADSA AADSAFAQRL NFLSSNPHFE QRQNQFIVKH YAGDVTYSIT
GMTDKNKDQL ATDILNLIHS SNNEFMKSIF PVAEESNSRR RPPTAGDRIK TSANDLVETL
MKCQPSYIRT IKPNQTKSPN DYDQQMVLHQ IKYLGLQENI RIRRAGFAYR QAFDTFAQRF
AVLSGKTSYA GEYTWQGDDK SACEQILKDT NIPSSEYQMG TSKVFIKNPE TLFALEDMRD
KFWDTMATRI QRAWRSYVRR RSEAAACIQK LWNRNKVNME LERVRNEGTK LLQGKKQRRR
YSILGSRKFY GDYLSASKPN GTLWNTCGLS QNDHVIFSMR CEVLVHKLGR TSKPSPRQLV
LTKKNLYLVI TKIVDQKLTQ QVEKKFAVSS IDSVGLTNLQ DDWVAIRNKS SQNGDMFLRC
FFKTEFITTL KRINRNIQVI VGPTIQYCRK PGKVQTVKTA KDETTKDYDY YKSGTIHVGT

Disorder confidence
Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
970 980 990 1000 1010 1020
GLPPTSKSKP FPRLATGGST AAARGPRPVV QNKPAATKPV SMPAAKSKPA PMANPVSTAQ
1030 1040 1050 1060 1070 1080
QTQNRPPAPA MQARPNTTQA AAPVTSTTTT IKQATTVSAS KPAPSTVTSA ASSPSNISKP
1090 1100 1110 1120 1130 1140
SAPVANNVSK PSAVPPPPPP PPAEVEKKDL YLALYDFAGR SPNEMTIKKD EIIEIVQKEP
1150 1160 1170 1180 1190 1200
SGWWLALKNG AEGWVPATYV TEYKGSTPQT TASSTNVAAQ ANNNASPAEV NNLAGSLADA
1210
LRMRASAVRG SDEEEDW
Confidence Key
High(9)
Low (0)
? Disordered
Alpha helix
Beta strand