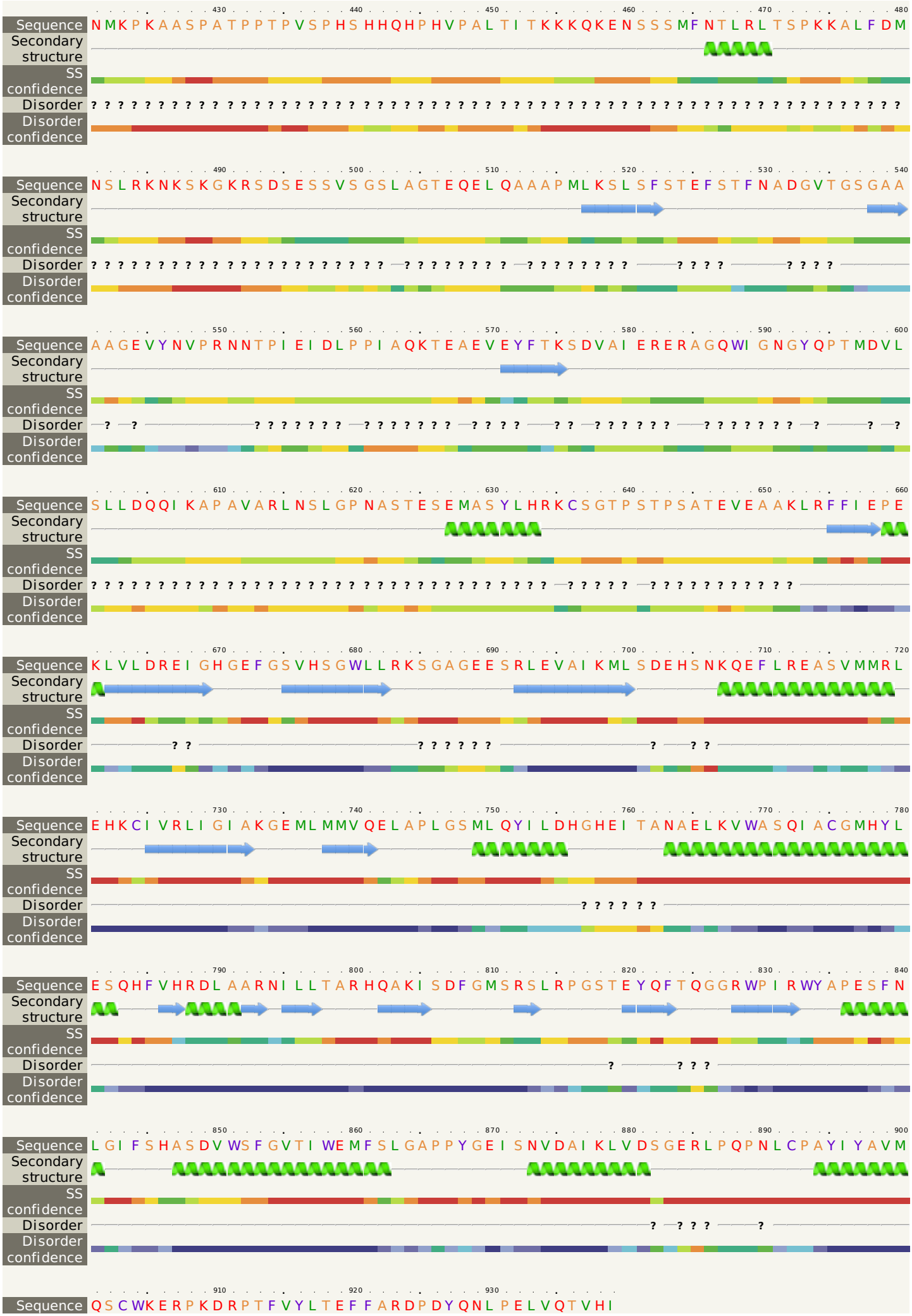
Email	l.a.kelley@imperial.ac.uk
Description	Q24145
Date	Tue Jan 22 11:35:34 GMT 2013
Unique Job ID	8157dca3e4a2af7d

Secondary structure and disorder prediction


Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
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490 500 510 520 530 540
NSLRKNKSKGKRSDSESSVSGSLAGTEQELQAAAPMLKSLSFSTEFSTFNADGVTGSGAA
550 560 570 580 590 600
AAGEVYNVPRNNTPIEIDLPPIAQKTEAEVEYFTKSDVAIERERAGQWIGNGYQPTMDVL
610 620 630 640 650 660
SLLDQQIKAPAVARLNSLGPNASTESEMASYLHRKCSGTPSTPSATEVEAAKLRFFIEPE
670 680 690 700 710 720
KLVLDREIGHGEFGSVHSGWLLRKSGAGEESRLEVAIKMLSDEHSNKQEFLREASVMMRL
730 740 750 760 770 780
EHKCIVRLIGIAKGEMLMMVQELAPLGSMLQYILDHGHEITANAELKVWASQIACGMHYL
790 800 810 820 830 840
ESQHFVHRDLAARNILLTARHQAKISDFGMSRSLRPGSTEYQFTQGGRWPIRWYAPESFN
850 860 870 880 890 900
LGIFSHASDVWSFGVTIWEMFSLGAPPYGEISNVDAIKLVDSGERLPQPNLCPAYIYAVM
910 920 930
Sequence
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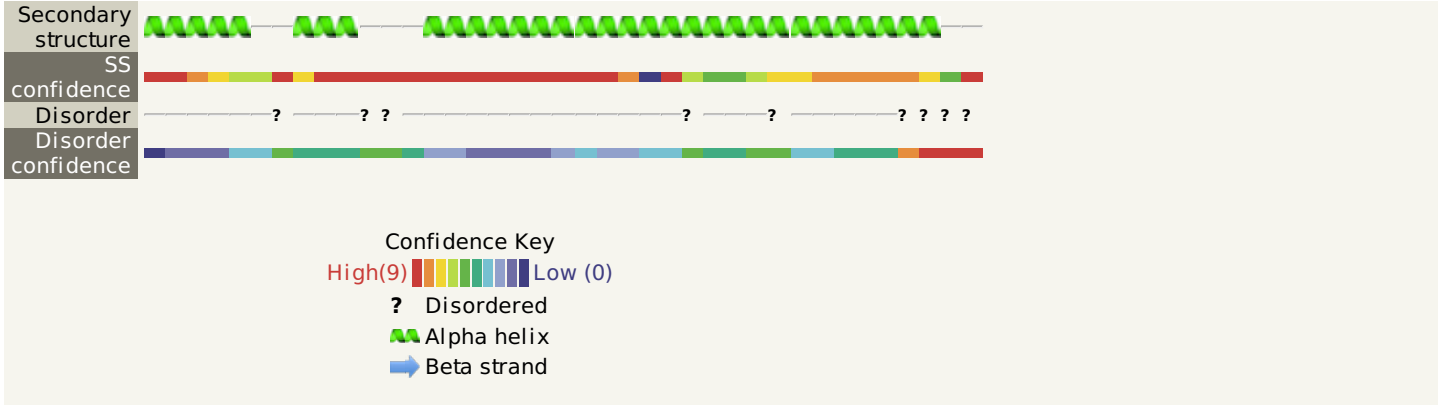
Secondary structure
SS confidence
Disorder
Disorder confidence
Confidence Key
High(9)
Low (0)
? Disordered
Alpha helix
Beta strand