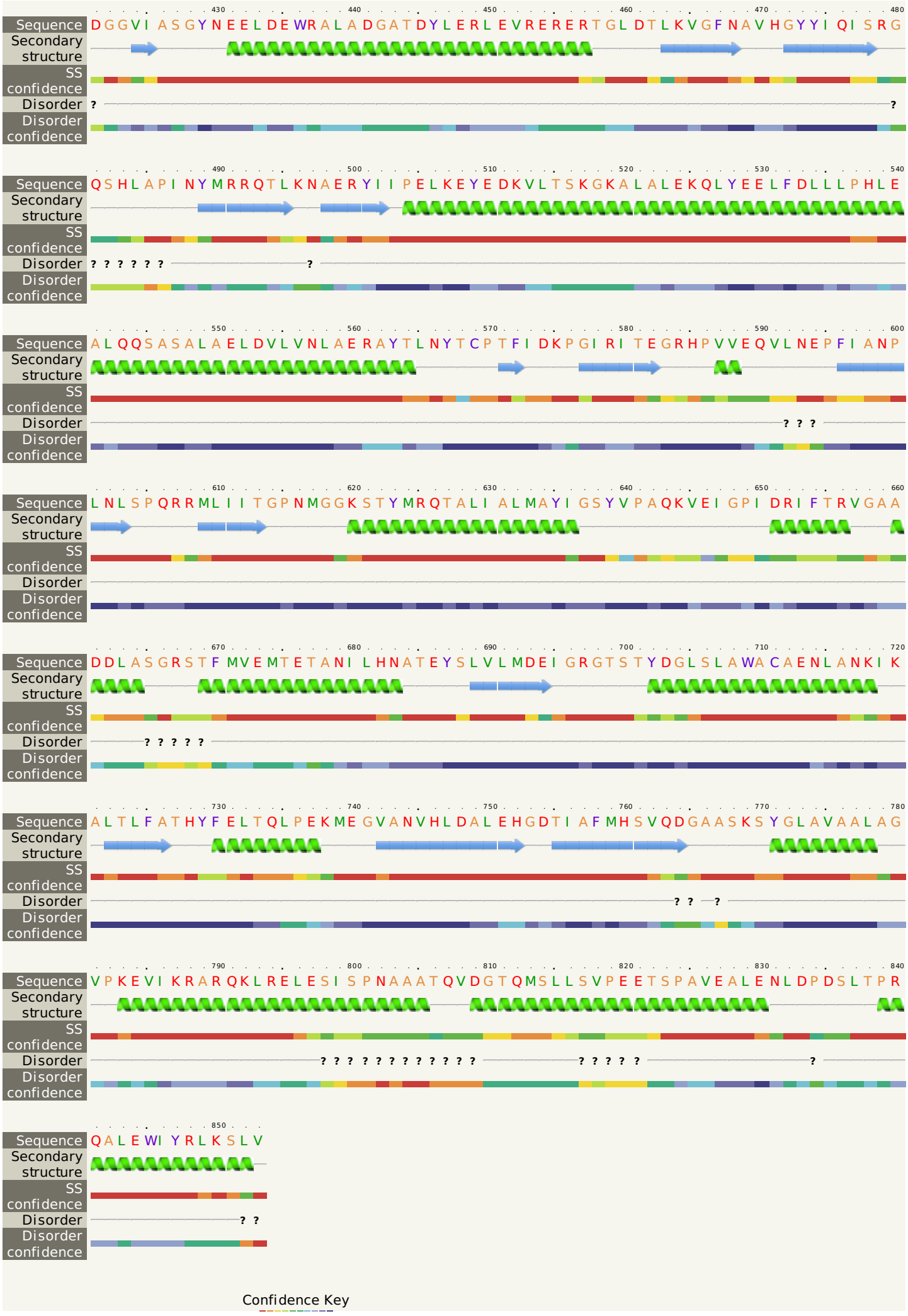
Email	l.a.kelley@imperial.ac.uk
Description	P23909
Date	Thu Jan 5 11:40:34 GMT 2012
Unique Job ID	cff3e7de5403a2df

Secondary structure and disorder prediction


Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
DGGVI ASGYNEELDEWRALADGATDYLERLEVRERERTGLDTLKVGFNAVHGYYI QI SRG
490 500 510 520 530 540
QSHLAPI NYMRRQTLKNAERYI I PELKEYEDKVLTSKGKALALEKQLYEEELFDLLLPHLE
550 560 570 580 590 600
ALQQSASALAELDVLVNLAERAYTLNYTCPTFI DKPGI RI TEGRHPVVEQVLNEPFI ANP
610 620 630 640 650 660
LNLSPQRRMLI I TGPNMGGKSTYMRQTALI ALMAYI GSYVPAQKVEI GPI DRI FTRVGAA
670 680 690 700 710 720
DDLASGRSTF MVEMTETANI LHNATEYSLVLMDEI GRGTSTYDGLSLAWACAENLANKI K
730 740 750 760 770 780
ALTLFATHYF ELTQLPEKME GVANVHLDALEHGDTI AFMHSVQDGAASKSYGLAVAALAG
790 800 810 820 830 840
VPKEVI KRARQKLRELESI SPNAAATQVDGTQMSLLSVPEETSPAVEALENLDPDSLTPR
850
QALEWI YRLKSLV
Confidence Key

High(9) Low (0)

? Disordered

Alpha helix

Beta strand