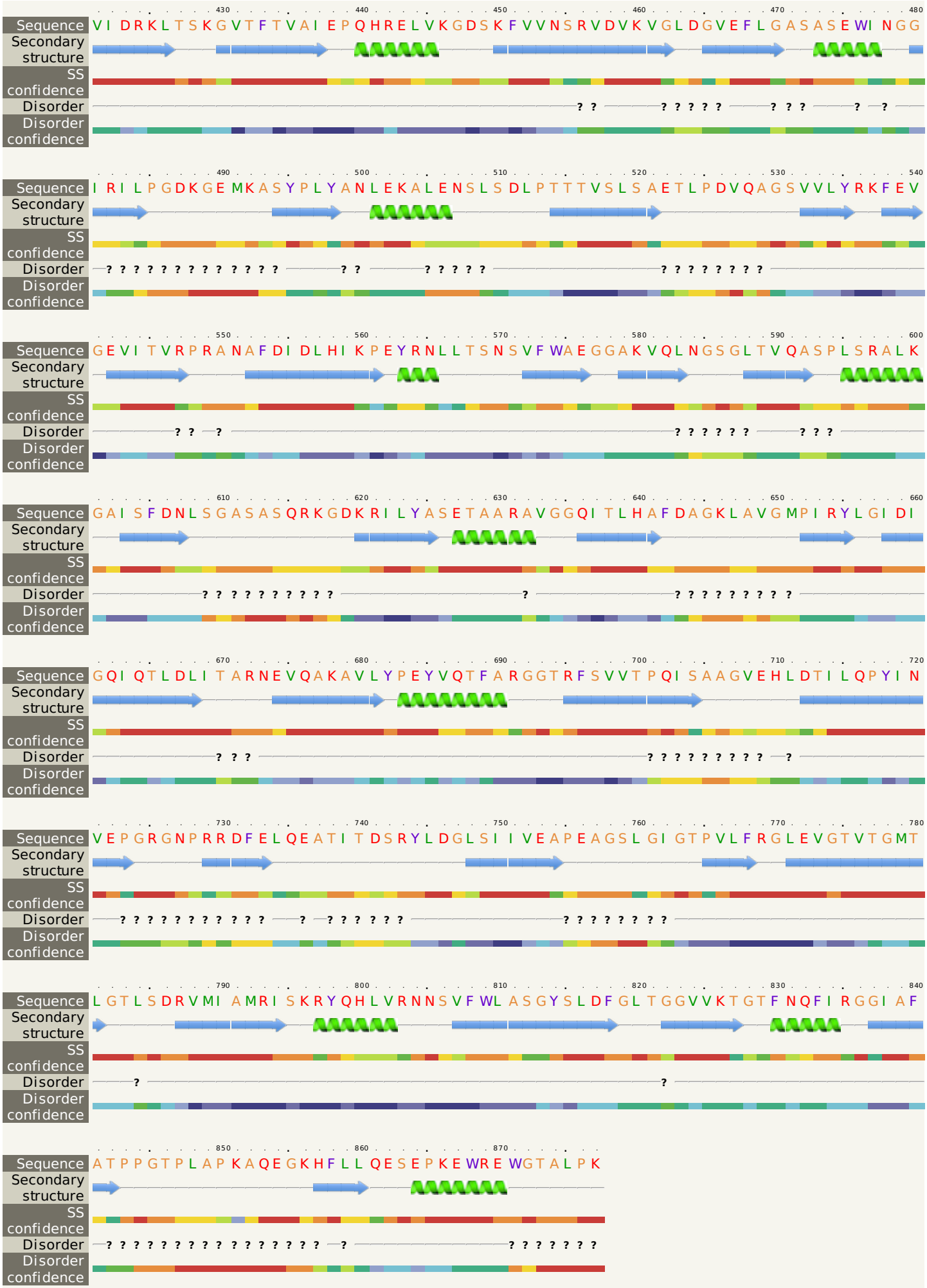
Email	l.a.kelley@imperial.ac.uk
Description	P76272
Date	Thu Jan 5 12:21:28 GMT 2012
Unique Job ID	a224dc802c7002a2

Secondary structure and disorder prediction


Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
VI DRKLTSKGVTFTVAI EPQHRELVKGDSKFVVNSRVDVKVGLDGVEFLGASASEWI NGG
490 500 510 520 530 540
I RI LPGDKGE MKASYPLYANLEKALENSLSDLPTTTVSLSAETLPDVQAGSVVLYRKFEV
550 560 570 580 590 600
GEVI TVRPRANAFDI DLHI KPEYRNLLTSNSVFWAEGGAKVQLNGSGLTVQASPLSRALK
610 620 630 640 650 660
GAI SFDNLSGASASQRKGDKRI LYASETAARAVGGQI TLHAFDAGKLAVGMPI RYLGI DI
670 680 690 700 710 720
GQI QTLDLI TARNEVQAKAVLYPEYVQTFARGGTRFSVVTPQI SAAGVEHLDTI LQPYI N
730 740 750 760 770 780
VEPGRGNPRRDFELQEATI TDSRYLDGLSI I VEAPEAGSLGI GTPVLFRGLEVGTVTGMT
790 800 810 820 830 840
LGTLSDRVMI AMRI SKRYQHLVRNNSVFWL ASGYSLDFGLTGGVVKTGTF NQFI RGGI AF
850 860 870
ATPPGTPLAP KAQEGKHFLL QESEPKEWRE WGTALPK
Confidence Key

High(9) Low (0)

? Disordered

Alpha helix

Beta strand