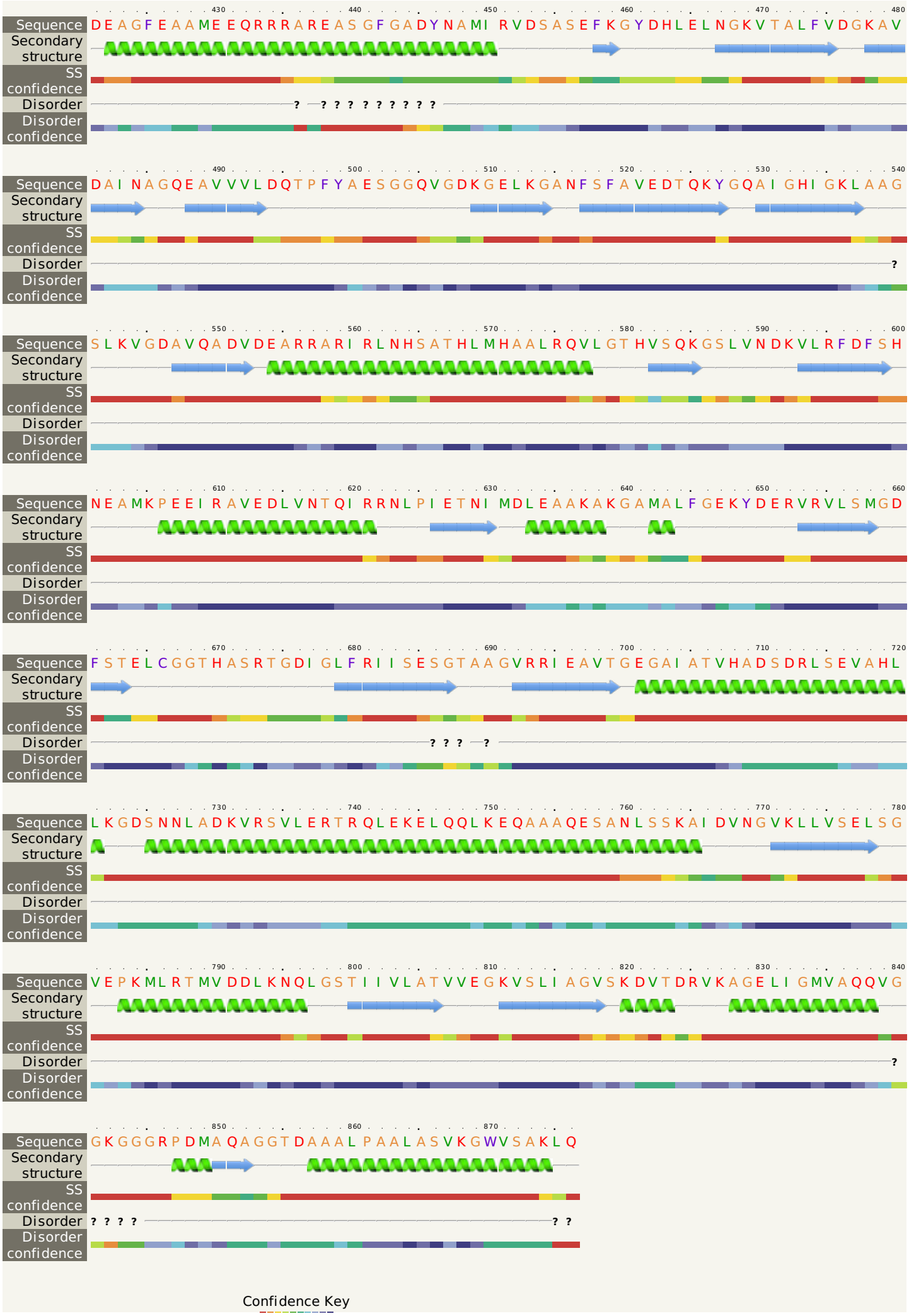
Email	l.a.kelley@imperial.ac.uk
Description	P00957
Date	Thu Jan 5 10:57:20 GMT 2012
Unique Job ID	be922b2a1da9a3c5

Secondary structure and disorder prediction


Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
DEAGFEAAMEEQRRRAREASGFGADYNAMI RVDSASEFKGYDHLELNGKVTALFVDGKAV
490 500 510 520 530 540
DAI NAGQEAVVVLDQTPFYAESGGQVGDKGELKGANFSFAVEDTQKYGQAI GHI GKLAAG
550 560 570 580 590 600
SLKVGDAVQADVDEARRARI RLNHSATHLMHAALRQVLGT HVSQKGSLVNDKVLRFDFSH
610 620 630 640 650 660
NEAMKPEEI RAVEDLVNTQI RRNLPI ETNI MDLEAAKAKGAMALFGEKYDERVRVLSMGD
670 680 690 700 710 720
FSTELCGGTHASRTGDI GLFRII SESGTAAGVRRI EAVTGEGAI ATVHADSDRLSEVAHL
730 740 750 760 770 780
LKGDSNNLADKVRSVLERTRQLEKELQQLKEQAAAQESANLSSKAI DVNGVKLLVSELSG
790 800 810 820 830 840
VEPKMLRTMV DDLKNQLGSTII VLATVVEGKVSLI AGVSKDVTDRVKAGELI GMVAQQVG
850 860 870
GKGGGRPDMA QAGGTDAAAL PAALASVKGWVSAKLQ
Confidence Key

High(9) Low (0)

? Disordered

Alpha helix

Beta strand