Email	l.a.kelley@imperial.ac.uk
Description	Q62415
Date	Tue Jul 30 13:16:45 BST 2013
Unique Job ID	eb04dc0bf7b652f9

Secondary structure and disorder prediction

Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
AI SSQPLPLS ALGATEKLGI EI GKGPPPI P GVGKPLPPSY GTYPSSGPLG PGSTSSLERR
490 500 510 520 530 540
KEGSLPRPGA GPPSRQKPAP LPPASNAPQP GSSQQI QQRI SVPPSPTYPP AGPPAFPTGD
550 560 570 580 590 600
GKPELPLTVA I RPFLADKGS RPQSPRKGPQ TVNSSSI YSMYLQQATPPKN YQPPAHGTLN
610 620 630 640 650 660
KSVKAVYGKP VLPSGSASPS PLPFLHGSLG TGTAQPQPPS DSAEKEPEQE GPSVPGEGST
670 680 690 700 710 720
VESLPRPLSP TKLTPI VHSP LRYQSDADLE ALRRKLANAP RPLKKRSSI T EPEGPGGPNI
730 740 750 760 770 780
QKLLYQRFNT LAGGMEGTPF YQPSPSQDFV GTLADMDNGN TNANGNLDEP FPPRPTAPLP
790 800 810 820 830 840
EELAPSSDAN DNELPSPEPE ELI CPQTTHQTAE PTEDNNN NVAPVPSTEQI PSPVAEAPS
850 860 870 880 890 900
EEDQVPPAPL SPVI HPPAAS ASKRTNLKKP NSERTGHGLR VRFNPLALLL DASLEGEFDL
910 920 930 940 950 960
VQRI I YEVED PSKPNDEGI T PLHNAVCAGH HHI VKFLLDF GVNVNAADSD GWTPLHCAAS

SS confidence
Disorder
Disorder confidence
970 980 990 1000 1010 1020
Sequence C N S V H L C K Q L V E S G A A I F A S T I S D I E T A A D K C E E M E E G Y I Q C S Q F L Y G V Q E K L G V M N K G T
Secondary structure
SS confidence
Disorder
Disorder confidence
1030 1040 1050 1060 1070 1080
Sequence V Y A L W D Y E A Q N S D E L S F H E G D A I T I L R R K D E N E T E W W W A R L G D R E G Y V P K N L L G L Y P R I K
Secondary structure
SS confidence
Disorder
Disorder confidence
Sequence P R Q R T L A
Secondary structure
SS confidence
Disorder
Disorder confidence
Confidence Key
High(9)
Low (0)
Disordered
Alpha helix
Beta strand