

Email	l.a.kelley@imperial.ac.uk
Description	P37647
Date	Thu Jan 5 11:56:19 GMT 2012
Unique Job ID	91d84f335408d2f4

1 10 20 30 40 50 60

Sequence MSKKI AVI GECMI ELSEKGA DVKRG FGGDT LNTSVYI ARQV DPAALT VHY VTAL GTDS FS

Secondary structure

SS

confidence

Disorder ? ? ?

Disorder confidence

70 80 90 100 110 120

Sequence QQML DA WHGE NVDTSL TQR MENRL PGLYYI ETDST GERT FYY WRNE AAK FVLE SEQSA A

Secondary structure

SS

confidence

Disorder

Disorder confidence

130 140 150 160 170 180

Sequence ICEEL ANFDY LLSGI SLAI LSPTS REKLL SLLRECR ANGKVI FDN NYR PRL WASKE ET

Secondary structure

SS

confidence

Disorder ?

Disorder confidence

190 200 210 220 230 240

Sequence QQVY QQMLE CTDI AFLTL DDE DALWGQQ PVEDVI ARTHNA GVK E VVKRG ADSCL VSI AG

Secondary structure

SS

confidence

Disorder

Disorder confidence

250 260 270 280 290 300

Sequence EGLVD VPAVK LPEKVI DTTAAGDS FSAGY LAVRL TGGS AEDA AKRGHL TASTVI QYRGA

Secondary structure

SS

confidence

Disorder

Disorder confidence

Sequence I I P R E A M P A

Secondary structure


SS

confidence


Disorder ? ? ? ? ? ? ?


Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand