

Email	l.a.kelley@imperial.ac.uk
Description	P0ADG4
Date	Thu Jan 5 11:20:58 GMT 2012
Unique Job ID	d2548836fe9ba9dd

1 10 20 30 40 50 60

Sequence MHPMLNI AVRAARKAGNLI AKNYETPDAVEASQKGSNDFVVTNVDKAAEAVII DTIRKSYIP

Secondary structure

SS confidence

Disorder

Disorder confidence

70 80 90 100 110 120

Sequence QHTII TEESGELEGT DQDVQWVI DPLDGTTFNFI KRLPHFAVSI AVRIRKGRTEVAVVYDPM

Secondary structure

SS confidence

Disorder

Disorder confidence

130 140 150 160 170 180

Sequence RNELFTATRGGQAQLNGYRLRGSTARDLDGTILATGFPFKAKQYATTYINIVGKLFNECA

Secondary structure

SS confidence

Disorder

Disorder confidence

190 200 210 220 230 240

Sequence DFRRTGSAALDLAYVAAGRV DGF FEIGLRP WDF AAGELLVREAGGI VSDFTGGHNYMLTG

Secondary structure

SS confidence

Disorder

Disorder confidence

250 260

Sequence NIVAGNPRVVKAMLANMRDELS DALKR


Secondary structure

SS confidence


Disorder


Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand