

Email	l.a.kelley@imperial.ac.uk
Description	P0ABP8
Date	Thu Jan 5 11:16:02 GMT 2012
Unique Job ID	421ea58865332278

1 10 20 30 40 50 60

Sequence MATPHI NAEMGDFADVVLMPGDPLRAKYIAETFL EDAREVNNVRGMLGFTGT YKGRKISV

Secondary structure

SS

confidence

Disorder

Disorder confidence

70 80 90 100 110 120

Sequence MGHGMGIPSCSIYTKELITDFGVKKIIRVGSCGAVLPHVKLRD VVIGMGACTDSKVNRI R

Secondary structure

SS

confidence

Disorder

Disorder confidence

130 140 150 160 170 180

Sequence FKDHDFAAIADFDMVRNAVDAAKALGIDARVGNLFSADLFYSPDGEMFDVMEKYGILGVE

Secondary structure

SS

confidence

Disorder

Disorder confidence

190 200 210 220 230

Sequence MEAAGIYGVAAEFGAKAL TICTVSDHIRT HEQT TAAERQTTFNDMIKIALESVLLGDKE

Secondary structure

SS

confidence

Disorder

Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand