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|---------------|------------------------------|
| Email | l.a.kelley@imperial.ac.uk |
| Description | Q95PX0 |
| Date | Sat Apr 13 13:19:58 BST 2013 |
| Unique Job ID | 957911be8e771a01 |

1 10 20 30 40 50 60

Sequence MS SKTKAVVKFTYEPRLDELGLTKGDFVYVVEKSTDGWWKGEAPNGGGVGVFP SNYVEEV

Secondary structure

SS

confidence

Disorder

Disorder confidence

70 80 90 100 110 120

Sequence EASTNGNQGS I ENRNPA AAVPAPI MMQAPPPKLQASRSSFEI QVVVALY SFDASSSEEL

Secondary structure

SS

confidence

Disorder

Disorder confidence

130 140 150 160 170 180

Sequence SF KKGERLEI VDHPEHDPD VWMARNASGTTGLVPRNYI EVVVNDSSSSKASHQDFAPQYS

Secondary structure

SS

confidence

Disorder

Disorder confidence

190 200 210 220 230 240

Sequence GNGEI P MEQQP WYFGRISRERAEDLLHGREGF LVRDSESNPGDLSI SMRGI ERNKHFK

Secondary structure

SS

confidence

Disorder

Disorder confidence

250 260 270 280

Sequence VQNV DGL LKI GNRTF VDMNALI NHYTTSP I FSSPTEK LFLTGPLPK

Secondary structure


SS

confidence


Disorder


Disorder confidence

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand