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|---------------|-----------------------------|
| Email | l.a.kelley@imperial.ac.uk |
| Description | P0A6D5 |
| Date | Thu Jan 5 11:02:53 GMT 2012 |
| Unique Job ID | 93862a7a82db52a6 |

| | 1 | 10 | 20 | 30 | 40 | 50 | 60 |
|---------------------|--|----|----|----|----|----|----|
| Sequence | MDVTAKEYELI GLMAYPI RHS LSP EMQNKAL EKAGLP FPTYMAFEVDNDSFP GAI EGLKALK | | | | | | |
| Secondary structure | | | | | | | |
| SS confidence | | | | | | | |
| Disorder | ? ? ? ? ? | | | | | | |
| Disorder confidence | | | | | | | |


| | 70 | 80 | 90 | 100 | 110 | 120 |
|---------------------|---|----|----|-----|-----|-----|
| Sequence | MRGTGV SMPNKQLACEYVDE LTPAAKLVGAI NTI VDDGYLRGYNTDGTGHI RAI KESGF | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | | | | | | |
| Disorder confidence | | | | | | |

| | 130 | 140 | 150 | 160 | 170 | 180 |
|---------------------|--|-----|-----|-----|-----|-----|
| Sequence | DI KGKTMVLL GAGGASTAI GAQGAI EGLKEI KLFNRRDEFFDKALAF AQRV NENTDCVV T | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | | | | | | |
| Disorder confidence | | | | | | |


| | 190 | 200 | 210 | 220 | 230 | 240 |
|---------------------|--|-----|-----|-----|-----|-----|
| Sequence | VTDLADQQQAF AEALASADI LTNGTKVGMKPLENESLVNDI SLLHPGLLVTECVYNPHMTK | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | ? ? ? ? | | | | | |
| Disorder confidence | | | | | | |


| | 250 | 260 | 270 | 280 |
|---------------------|--|-----|-----|-----|
| Sequence | LLQQAQQAGCKTI DGYGMLL WQGAEQFTLWTGKDFPLEYVKQVMGFGA | | | |
| Secondary structure | | | | |
| SS confidence | | | | |
| Disorder | ? ? ? | | | |
| Disorder confidence | | | | |

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand