

| | |
|---------------|-----------------------------|
| Email | l.a.kelley@imperial.ac.uk |
| Description | Q47138 |
| Date | Thu Jan 5 12:36:15 GMT 2012 |
| Unique Job ID | ac0f23ef389be2d5 |

| | 1 | 20 | 30 | 40 | 50 | 60 |
|---------------------|--|----|----|----|----|----|
| Sequence | MSKVFICAAIPDELATREEGAVAVATAIEAGDERRARRAKFHWQFLEHYPAAQDCAYKFI V | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | ? ? ? ? ? ? ? ? ? ? | | | | | |
| Disorder confidence | | | | | | |


| | 70 | 80 | 90 | 100 | 110 | 120 |
|---------------------|--|----|----|-----|-----|-----|
| Sequence | CEDKPGIPRPALDSWDAEYMQENRWDEESASFVPVETESDPMNVTFDKLAPENVQNAVMMVK | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | ? ? ? ? ? ? ? ? ? ? ? ? | | | | | |
| Disorder confidence | | | | | | |

| | 130 | 140 | 150 | 160 | 170 | 180 |
|---------------------|---|-----|-----|-----|-----|-----|
| Sequence | FDTCEINTVDMVISAQELLQEDMATFDGHI VEALMKMPEVNAMYPELKLHAI GVKHKCI | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | ? ? ? | | | | | |
| Disorder confidence | | | | | | |


| | 190 | 200 | 210 | 220 | 230 | 240 |
|---------------------|---|-----|-----|-----|-----|-----|
| Sequence | PGAKWPVEIQAE MRI WKKRREG ERKETGKYTSVVDLARARANQQYTENSTGKISPVI AAIH | | | | | |
| Secondary structure | | | | | | |
| SS confidence | | | | | | |
| Disorder | ? | | | | | |
| Disorder confidence | | | | | | |


| | 250 |
|---------------------|------------------|
| Sequence | REYKQTWKTL DDELA |
| Secondary structure | |
| SS confidence | |
| Disorder | ? ? ? ? |
| Disorder confidence | |

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand