

Email	l.a.kelley@imperial.ac.uk
Description	A4UR77
Date	Thu Jan 5 10:55:05 GMT 2012
Unique Job ID	48dfafd19ed8ae98

Sequence: MMQPVI NAPEI ATAREQQLF NGKNFHVFI YNKTESI SGLHQHDYEFETLVLTGRYFQEI N

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [Color scale]

Disorder: [Color scale]

Disorder confidence: [Color scale]

Sequence: GKRVL LERGD FVFI PLGSHHQS FYEFGATR ILNVGI SKRFFEQHLYLPLL PYCFVASQVYR

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [Color scale]

Disorder: [Color scale]

Disorder confidence: [Color scale]

Sequence: TNNALFTYVETVI SSLNVRETGLEEFVEMVTFYVINRLRHYYREEQVI DDVPQWLKSTVEK

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [Color scale]

Disorder: [Color scale]

Disorder confidence: [Color scale]

Sequence: MHDKEQFS ESAL ENMV ALSAKSQEYLTRATQRYYGKTPMQI INEIRINFAKKQLEMTNYS

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [Color scale]

Disorder: [Color scale]

Disorder confidence: [Color scale]

Sequence: VTDIAFEAGYSSPSLFI KTFKKLTSFTP KSYRKKLTEFNQ

Secondary structure: [Alpha-helices and beta-strands]

SS confidence: [Color scale]

Disorder: [Color scale]

Disorder confidence: [Color scale]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand