

Email	l.a.kelley@imperial.ac.uk
Description	P31142
Date	Thu Jan 5 11:47:20 GMT 2012
Unique Job ID	ceb3777181587bc2

Sequence: MSTTWFVGADWLAELHIDDEPILQIIDARMASPGQEDRNVAQEYLNNGHIPGAVVFFDIEALSD

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

SS confidence: [Color-coded bar]

Disorder: [Disordered regions marked with question marks]

Disorder confidence: [Color-coded bar]

Sequence: HTSPLPHMLPRPETFAVAMRELGVNQDKHLIVYDEGNLFSAPRAWVWMLRTFGVEKVSILG

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

SS confidence: [Color-coded bar]

Disorder: [Disordered regions marked with question marks]

Disorder confidence: [Color-coded bar]

Sequence: GGLAGWQRDDLLEEGAVELPEGEFNAAFNPPEAVVKVTDVLLASHENTAQIIDARPAARF

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

SS confidence: [Color-coded bar]

Disorder: [Disordered regions marked with question marks]

Disorder confidence: [Color-coded bar]

Sequence: NAEVDEPRPGLRRGHI PGALNVPWTELVREGELKTTDELDAIFFGRGVSYDKPIIVSCGS

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

SS confidence: [Color-coded bar]

Disorder: [Disordered regions marked with question marks]

Disorder confidence: [Color-coded bar]

Sequence: GVTAAVLLALATLDVPPNVKLYDGAWSEWGARADLPVEPVK

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

SS confidence: [Color-coded bar]

Disorder: [Disordered regions marked with question marks]

Disorder confidence: [Color-coded bar]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand