

Email	l.a.kelley@imperial.ac.uk
Description	P0AD42
Date	Thu Jan 5 11:19:57 GMT 2012
Unique Job ID	4ca95936a2c68456

Sequence: MATHERRVVF¹FDL¹⁰DGTL²⁰HQQDM³⁰FGS⁴⁰FLRYLLRRQP⁵⁰LNALLVLP⁶⁰LLPI⁷⁰IAI⁸⁰ALLIKGRAAR⁹⁰

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bars]

Sequence: WPMS⁷⁰LLLVGCT⁸⁰FGHSEARL⁹⁰QTLQADFVRWF¹⁰⁰RDNVTA¹¹⁰FP¹²⁰LVQERLT¹³⁰TYLLSSDADI¹⁴⁰WLI¹⁵⁰TG

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bars]

Sequence: SPQPLVEAVY¹³⁰FDTP¹⁴⁰WLP¹⁵⁰RVNLI¹⁶⁰ASQIQRGY¹⁷⁰GGWVL¹⁸⁰TMRC¹⁹⁰LGHEKVAQLERKI²⁰⁰GTPLRLYS²¹⁰

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bars]

Sequence: GYSDSNQDN¹⁹⁰PLL²⁰⁰LYFCQHR²¹⁰WRVTPRGELQQL²²⁰E

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

SS confidence: [Confidence bars]

Disorder: [Disorder prediction]

Disorder confidence: [Disorder confidence bars]

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand