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Description	P12996
Date	Thu Jan 5 11:33:15 GMT 2012
Unique Job ID	0b6ce921cfb45f8a

	1	10	20	30	40	50	60
Sequence	MAHRPRWTL S QVTELF EKPLLDLLFEAQQVHRQHFDP RQVQVSTLLSIKTGACPEDCKYC						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	PQSSRYKTGL EAERLMEVEQVLESARKAKAAGSTRFCMGA A WKNPHERDMPYLEQMVQGV						
Secondary structure							
SS confidence							
Disorder	- ? - ? ? ? ? - ? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	KAMGLEACMTL GTLSE SQAQRLANAGLDYYNHNLDTSPEFYGNII TTRTYQERLDTLEKV						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	190	200	210	220	230	240	
Sequence	RDAGI KVC SGGI VGLGETVKDRAGLLQLANLPTPPESVPI NMLVKVKGTPLADNDDVDA						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	250	260	270	280	290	300	
Sequence	FDFIRTI AVARI MMPTSYVRLSAGREQMNEQTQAMCFMAGANSI FYGCKLLTTPNP EEDK						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	310	320	330	340			
Sequence	DLQLFRKLG LNPQQTAVLAGDNEQQQRLEQALMTPD TDEYYNAAAL						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand