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Description	P0AC02
Date	Thu Jan 5 11:16:49 GMT 2012
Unique Job ID	6b53a978223eac03

	1	10	20	30	40	50	60
Sequence	MTRMKYLVA AATLSLFLAGCSGSKEEVPDNPPNEIYATAQQKLQDGNWRQAI TQLEALDN						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	RYPFGPY SQVQLDIYAYYKNADLPLAQAAIDRFIRLNPTHPNIDYVMYMRGLTNMALD						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	DSALQGFFGVDRSDRDPQHARA AFSDFSKLVRGYPNSQYTTDATKRLVFLKDRLAKYEYS						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	VAEYYTERGA VVAVVNRVEGMLRDY PDTQATRDALPLMENAYRQMQMNAQA EKVAKII AA						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							
Sequence	NSSNT						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand