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Description	P77569
Date	Thu Jan 5 12:30:40 GMT 2012
Unique Job ID	01da6b95b5f6bf43

	1	10	20	30	40	50	60
Sequence	MQNN	EQT EYKTV	RGLTRGLMLLN	NMLNKL	DGGASV	GLLAELSGLH	RRTTVRRLL
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ?						
Disorder confidence							

	70	80	90	100	110	120
Sequence	VRRSP	SDDSFRLTI	KVRQLSEGF	RDEQWISAL	AAPLLGDL	LREVVWPTD
Secondary structure						
SS confidence						
Disorder	? ? ?					
Disorder confidence						

	130	140	150	160	170	180
Sequence	RETTHRF	SRLSFRAMV	GRRLPL	LKTASGLT	WLAFCEP	EQDRKELI
Secondary structure						
SS confidence						
Disorder	? ? ? ? ? ?					
Disorder confidence						

	190	200	210	220	230	240
Sequence	EPLKLEAI	LARARKEGY	GQNYRGWD	QE EKIASIA	VPLRSEQRV	I GCLNLVY
Secondary structure						
SS confidence						
Disorder	?					
Disorder confidence						

	250	260	270
Sequence	AAEKHL	PALQRVAKQI	EEGVESQA
Secondary structure			
SS confidence			
Disorder	? ? ? ? ? ? ? ? ? ?		
Disorder confidence			

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand