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Description	P0ADD7
Date	Thu Jan 5 11:20:44 GMT 2012
Unique Job ID	b4f6b28f793b2f8d

	1	10	20	30	40	50	60
Sequence	MLPGCKNGI V I SKI PVMQAGLKEVMRTHFPEYEI I SSASAEDLTLLQLRRSGLVIADLA						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	GESEDPRSVC EHYYSLI SQYREI HWVFMVSR SWSQAVELLMCPTATLLS DVEPI ENLVK						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	130	140	150	160	170	180	
Sequence	TVRSGNTHAERI SAML TSPAMTETHDFS YRSVIL T LSEKVLRL LGKGVWGINQIASLLKK						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	SNKTI SAQKNSAMRRLAI HSNAEMYAWI NSAQGARELNLPSVYGDAAEVNTAELRRMSH						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	S						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand