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Description	P77538
Date	Thu Jan 5 12:30:21 GMT 2012
Unique Job ID	3ae6baf2b4348895

	1	10	20	30	40	50	60
Sequence	MALP V N K R V P K I L F I L F V V A F C V Y L V P R V A I N F F Y Y P D D K I Y G P D P W S A E S V E F T A K D G T						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	R L Q G W F I P S S T G P A D N A I A T I I H A H G N A G N M S A H W P L V S V L P E R N F N V F M F D Y R G F G K S K						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	G T P S Q A G L L D D T Q S A I N V V R H R S D V N P Q R L V L F G Q S I G G A N I L D V I G R G D R E G I R A V I L D						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	190	200	210	220	230	240	
Sequence	S T F A S Y A T I A N Q M I P G S G Y L L D E S Y S G E N Y I A S V S P I P L L L I H G K A D H V I P W Q H S E K L Y S						
Secondary structure							
SS confidence							
Disorder							
Disorder confidence							
	250	260	270	280			
Sequence	L A K E P K R L I L I P D G E H I D A F S D R H G D V Y R E Q M V D F I L S A L N P Q N						
Secondary structure							
SS confidence							
Disorder	? - ? ? ? ?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand