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Description	P0A870
Date	Thu Jan 5 11:07:02 GMT 2012
Unique Job ID	1606c35c02d4306e

	1	10	20	30	40	50	60
Sequence	M T D K L T S L R Q Y T T V V A D T G D I A A M K L Y Q P Q D A T T N P S L I L N A A Q I P E Y R K L I D D A V A W A K						
Secondary structure							
SS confidence							
Disorder	? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	Q Q S N D R A Q Q I V D A T D K L A V N I G L E I L K L V P G R I S T E V D A R L S Y D T E A S I A K A K R L I K L Y N						
Secondary structure							
SS confidence							
Disorder	? ? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	D A G I S N D R I L I K L A S T W Q G I R A A E Q L E K E G I N C N L T L L F S F A Q A R A C A E A G V F L I S P F V G						
Secondary structure							
SS confidence							
Disorder	?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	R I L D W Y K A N T D K K E Y A P A E D P G V V S V S E I Y Q Y Y K E H G Y E T V V M G A S F R N I G E I L E L A G C D						
Secondary structure							
SS confidence							
Disorder	? - ? ? ? ? ? ? ? ?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	R L T I A P A L L K E L A E S E G A I E R K L S Y T G E V K A R P A R I T E S E F L W Q H N Q D P M A V D K L A E G I R						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ?						
Disorder confidence							
	310						
Sequence	K F A I D Q E K L E K M I G D L L						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand