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Description	P43319
Date	Thu Jan 5 12:02:15 GMT 2012
Unique Job ID	69bb761763fa5c2a

	1	10	20	30	40	50	60
Sequence	MKRAPLI	TGLLLI	STSCAYASSGGCGADSTSGATNYS	SSVVDDVTVNQT	DNV	TGREFTSAT	
Secondary structure							
SS confidence							
Disorder	?	?	?	?			
Disorder confidence							
	70	80	90	100	110	120	
Sequence	LSS	TNWQYAC	SCSAGKAVKL	VYMVSPVLT	TTHGHQTGY	KLNDSLDI	KTTLQANDIPGLTT
Secondary structure							
SS confidence							
Disorder	?	?	?	?	?	?	?
Disorder confidence							
	130	140	150	160	170	180	
Sequence	DQVVSVNTRF	TQIKNNTVYS	AATQTGVCQGDTSR	YGPNIGANTTFTLY	GTKPFLGSMTI		
Secondary structure							
SS confidence							
Disorder	?	?	?	?	?	?	?
Disorder confidence							
	190	200	210	220	230	240	
Sequence	PKT	DAVKGA	WVDGMGSPSTGDFHDLV	KLSIQGNLTAPQSCKI	NQGDVI	KVNFGFIN	NGQ
Secondary structure							
SS confidence							
Disorder	?	?	?	?	?	?	?
Disorder confidence							
	250	260	270	280	290	300	
Sequence	KFT	TRNAMPD	GFTPVD	FIDITDCGDTSKI	KNSLQMRI	DGTTGVVDQYNLVARRR	SSDNP
Secondary structure							
SS confidence							
Disorder	?	?	?	?	?	?	?
Disorder confidence							
	310	320	330	340	350	360	
Sequence	DVGIRIENLG	GGVANIP	FQNGILPVDP	SGHGTVMNR	AWPVNLVGGGELETG	KFQGTATI	TV
Secondary structure							
SS confidence							
Disorder	?	?	?	?	?	?	?
Disorder confidence							

Sequence I V R

Secondary structure

SS

confidence

Disorder

Disorder confidence

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand