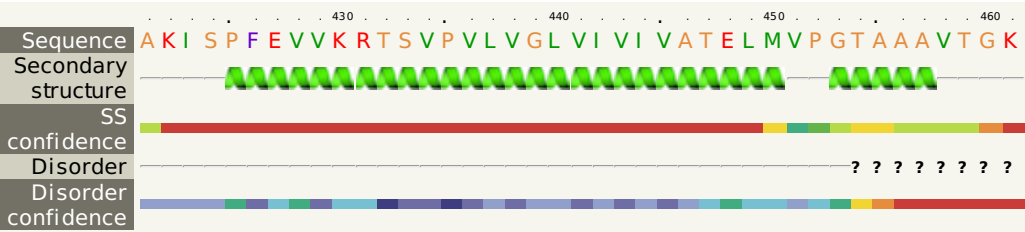


Email	l.a.kelley@imperial.ac.uk
Description	P0ABP3
Date	Thu Jan 5 11:15:59 GMT 2012
Unique Job ID	047988ec1e49e2f7

	1	10	20	30	40	50	60
Sequence	MLTFI ELLI GVVVI VGVARYI I KGY SATGVL FVGGLLLLI I SAI MGHKVL PSSQASTGY S						
Secondary structure							
SS confidence							
Disorder	? ? ?						
Disorder confidence							
	70	80	90	100	110	120	
Sequence	ATDI VEYVKI LLMSRGGDL GMMI MMLCGFAAYMTHI GANDMVVKLASKPL QYI NSPYLLM						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ?						
Disorder confidence							
	130	140	150	160	170	180	
Sequence	IAAYFVACLMSLAVSSATGL GVLLMATLFPVMVNVGI SRGAAAAI CASPAAI I LAPTSGD						
Secondary structure							
SS confidence							
Disorder	? ? ? ? ?						
Disorder confidence							
	190	200	210	220	230	240	
Sequence	VVLAAQASEMSLI DFAFKTTLPISI AAI I GMAI AHFFWQRYL DKKEHI SHEMLDVSEI TT						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							
	250	260	270	280	290	300	
Sequence	TAPAFYAI LPFTP I I GVLI F DGKWGPQLHI I TILVICMLI ASILEFLRSF NTQKVFSGLE						
Secondary structure							
SS confidence							
Disorder	? ? ? ?						
Disorder confidence							
	310	320	330	340	350	360	
Sequence	VAYRGMADAF ANVVMLLVAAGVFAQGLSTI GFI QSLI SIATSFGSASI I LMLVLVILTML						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							
	370	380	390	400	410	420	
Sequence	AAVTTGSGNAPFYAFVEMI PKLAHSSGI NPAYLTI PMLQASNLGRTLSPVSGVVVAVAGM						
Secondary structure							
SS confidence							
Disorder	? ?						
Disorder confidence							



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