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Description	P0ACD4
Date	Thu Jan 5 11:17:57 GMT 2012
Unique Job ID	080ad46ef068e9dd

**Panel 1: Residues 1-60**

Residue	Sequence	Secondary structure	SS	Disorder	Disorder confidence
1-10	MAYSEKVI	Alpha-helix	High	Low	High
11-20	DHYENPRNV	Alpha-helix	High	Low	High
21-30	GSFDNNDE	Alpha-helix	High	Low	High
31-40	VGSGMVGAP	Alpha-helix	High	Low	High
41-50	ACGDV MKLQI	Beta-strand	Low	High	Low
51-60	KVNDEGI IEDARFKT	Beta-strand	Low	High	Low


**Panel 2: Residues 70-120**

Residue	Sequence	Secondary structure	SS	Disorder	Disorder confidence
70-80	YGC GSAIASSSLVTE	Alpha-helix	High	Low	High
81-90	WVKGKSLDEAQAI	Alpha-helix	High	Low	High
91-100	KNTDI AEELLP	Alpha-helix	High	Low	High
101-110	PVKIHC S I LAEDA I	Beta-strand	Low	High	Low
111-120	KAAIAD	Beta-strand	Low	High	Low


**Panel 3: Residues 1-10**


Residue	Sequence	Secondary structure	SS	Disorder	Disorder confidence
1-10	YKSKREAK	Alpha-helix	High	Low	High

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand