


Email	l.a.kelley@imperial.ac.uk
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Date	Thu Jan 5 11:40:29 GMT 2012
Unique Job ID	b00ea95fe9a4b437


The figure displays two panels of protein analysis. The top panel shows the full-length protein (1-60) with its sequence, secondary structure (green alpha-helices), SS confidence (red bar), and disorder (blue bar with green segments). The bottom panel shows a protein fragment (70-110) with similar analysis. The disorder analysis indicates regions of low confidence (green) and high confidence (blue) for both proteins.


Panel	Sequence	Secondary structure	SS confidence	Disorder	Disorder confidence
Full-length (1-60)	MNPYIYLGGAILEVIGTTLMKFSEGFTRLWPSVGTIICYCASFWLLAQTLAYIPTGIA Y	Green alpha-helices	Red bar with green segments	Blue bar with green segments	Blue bar with green segments
Fragment (70-110)	AIWSGVGIVLISLLSWGFFGQRLDLPAILI GMMLICAGVLIINLLSRSTPH	Green alpha-helices	Red bar with green segments	Blue bar with green segments	Blue bar with green segments

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand