


Email	l.a.kelley@imperial.ac.uk
Description	P32719
Date	Thu Jan 5 11:50:42 GMT 2012
Unique Job ID	0f1f6edfc6bf637b


The figure displays three segments of protein analysis tracks:


- Segment 1 (Residues 1-60):** Sequence MKI SPS L M C M D L L K F K E Q I E F I D S H A D Y F H I D I M D G H F V P N L T L S P F F V S Q V K K L A T K P L. Secondary structure shows alpha-helices (green) and loops/coils (blue). SS confidence is high (red/yellow). Disorder is indicated by question marks.
- Segment 2 (Residues 70-120):** Sequence D C H L M V T R P Q D Y I A Q L A R A G A D F I T L H P E T I N G Q A F R L I D E I R R H D M K V G L I L N P E T P V E. Similar structural and confidence annotations.
- Segment 3 (Residues 130-180):** Sequence A M K Y Y I H K A D K I T V M T V D P G F A G Q P F I P E M L D K L A E L K A W R E R E G L E Y E I E V D G S C N Q A T. Includes disorder confidence track.
- Partial Segment 4 (Residues 190-230):** Sequence Y E K L M A A G A D V F I V G T S G L F N H A E N I D E A W R I M T A Q I L A A K S E V Q P H A K T A. Shows disorder confidence with multiple question marks.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand