

Email	l.a.kelley@imperial.ac.uk
Description	P22586
Date	Thu Jan 5 11:39:00 GMT 2012
Unique Job ID	7a4a98cd69a2b3ca

Protein 1 (P1): Sequence: M N N H A T V Q S S A P V S A A P L L Q V S G A L I A I I A L I L A A A W L V K R L G F A P K R T G V N G L K I S A S A. Secondary structure: Coiled-coil. SS confidence: High. Disorder: Low. Disorder confidence: High.

Protein 2 (P2): Sequence: S L G A R E R V V V V D V E D A R L V L G V T A G Q I N L L H K L P P S A P T E E I P Q T D F Q S V M K N L L K R S G R. Secondary structure: Domain structure with alpha-helices and beta-strands. SS confidence: High. Disorder: Low. Disorder confidence: High.

Protein 3 (P3): Sequence: S. Secondary structure: None. SS confidence: Low. Disorder: High. Disorder confidence: Low.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand