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Description	P37636
Date	Thu Jan 5 11:56:08 GMT 2012
Unique Job ID	c423edb132c55891

Protein: HNF1A (Human Nuclear Factor 1A)

Residues: 1 to 380

Sequence: M N R R R K L L I P L L F C G A M L T A C D D K S A E N A A A M T P E V G V V T L S P G S V N V L S E L P G R T V P Y E

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

Sequence: V A E I R P Q V G G I I K R N F I E G D K V N Q G D S L Y Q I D P A P L Q A E L N S A K G S L A K A L S T A S N A R I

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

Sequence: T F N R Q A S L L K T N Y V S R Q D Y D T A R T Q L N E A E A N V T V A K A A V E Q A T I N L Q Y A N V T S P I T G V S

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

Sequence: G K S S V T V G A L V T A N Q A D S L V T V Q R L D P I Y V D L T Q S V Q D F L R M K E E V A S G Q I K Q V Q G S T P V

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

Sequence: Q L N L E N G K R Y S Q T G T L K F S D P T V D E T T G S V T L R A I F P N P N G D L L P G M Y V T A L V D E G S R Q N

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

Sequence: V L L V P Q E G V T H N A Q G K A T A L I L D K D D V V Q L R E I E A S K A I G D Q V V V T S G L Q A G D R V I V S G L

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

Sequence: Q R I R P G I K A R A I S S S Q E N A S T E S K Q

Secondary structure: [Alpha-helices and Beta-strands]

SS confidence: [Confidence scale]

Disorder: [Disorder scale]

Disorder confidence: [Confidence scale]

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