







Conf:

Pred:

Pred: CCCC  
 AA: FERC

Legend:

= helix      Conf: = confidence of prediction

= strand      Pred: predicted secondary structure

= coil      AA: target sequence

Analysis performed with the use of PSIPRED Protein Structure Prediction Server

Citation: Bryson K, McGuffin LJ, Marsden RL, Ward JJ, Sodhi JS. & Jones DT. (2005)  
 Protein structure prediction servers at University College London.  
 Nucl. Acids Res. 33(Web Server issue):W36-38