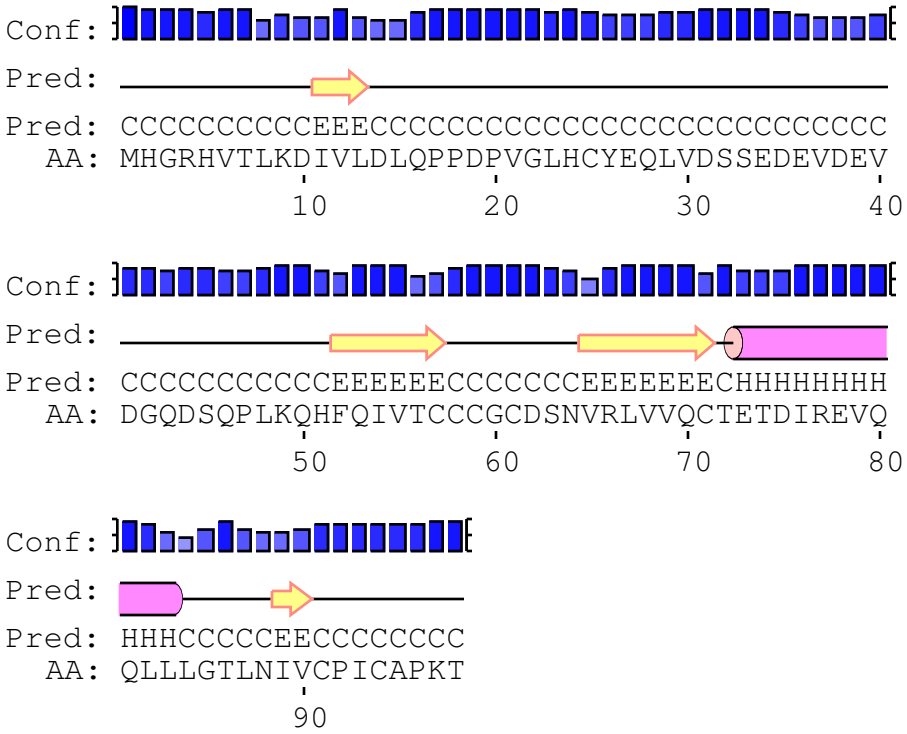

Legend:

= helix

= strand

= coil

Conf: = confidence of prediction

Pred: predicted secondary structure

AA: target sequence