





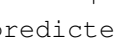
Legend:


 = helix

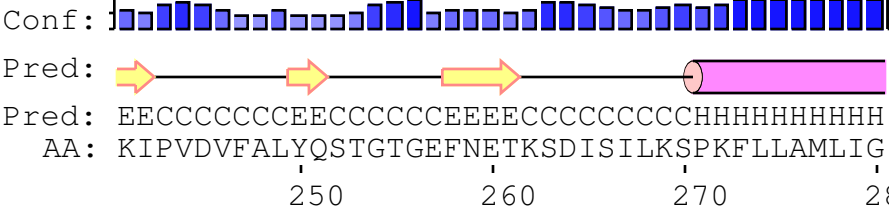
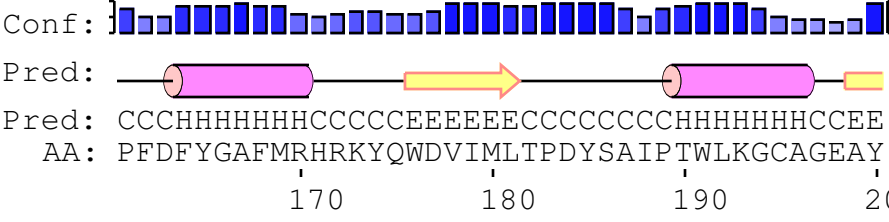
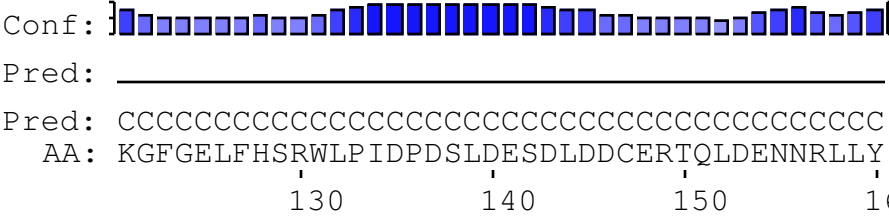
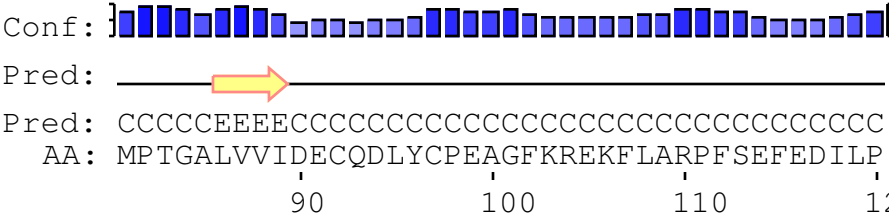
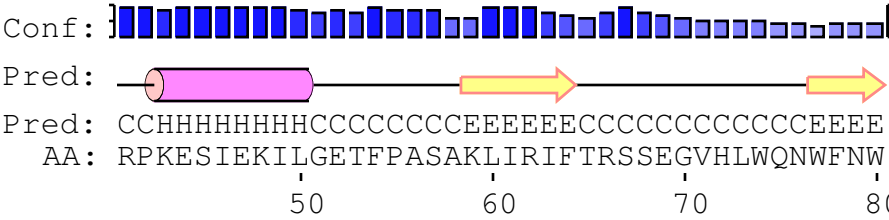
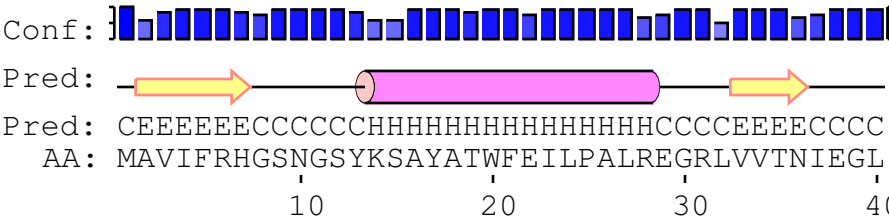
 = strand

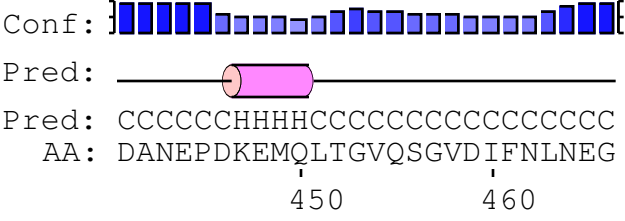
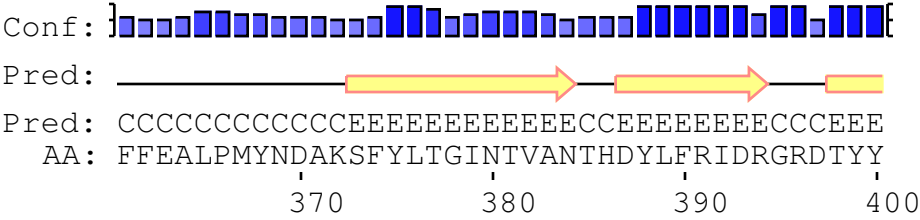
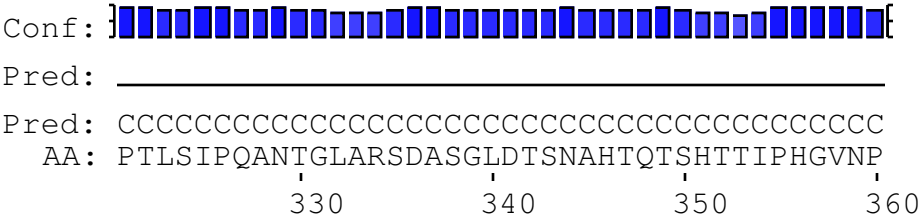
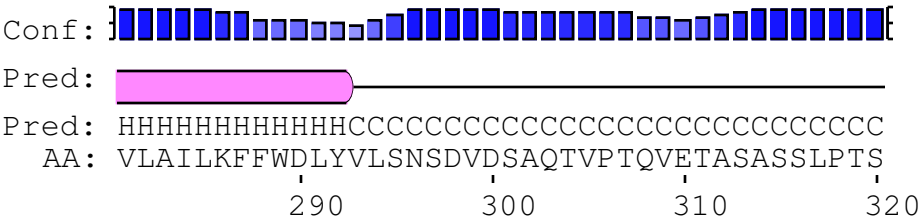
 = coil

Conf:  = confidence of prediction


Pred:  = predicted secondary structure

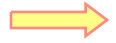
AA:  = target sequence







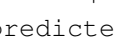
Legend:


 = helix

 = strand

 = coil

Conf:  = confidence of prediction

Pred:  = predicted secondary structure

AA:  = target sequence

