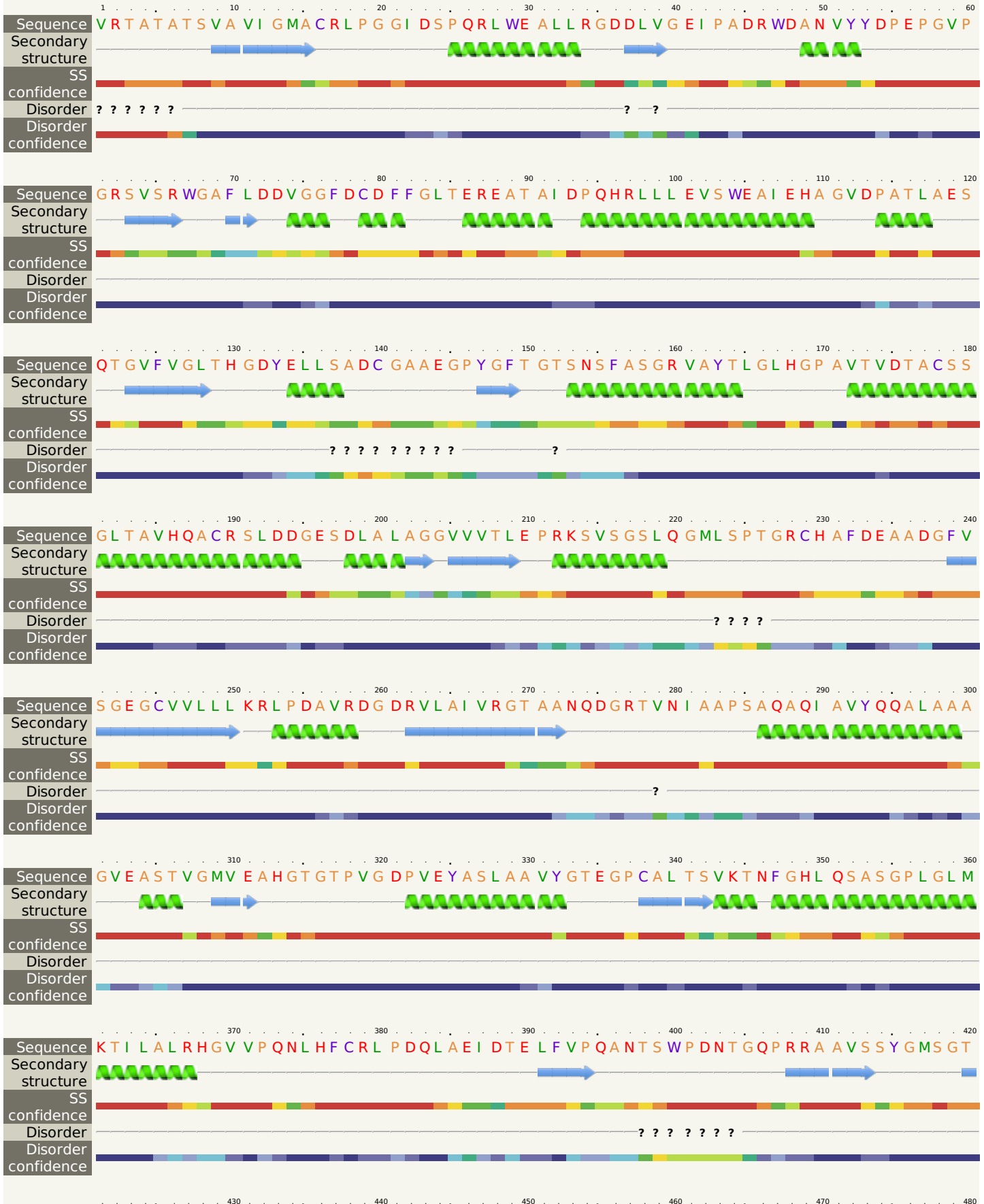
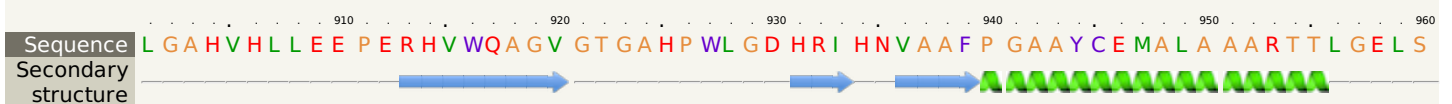
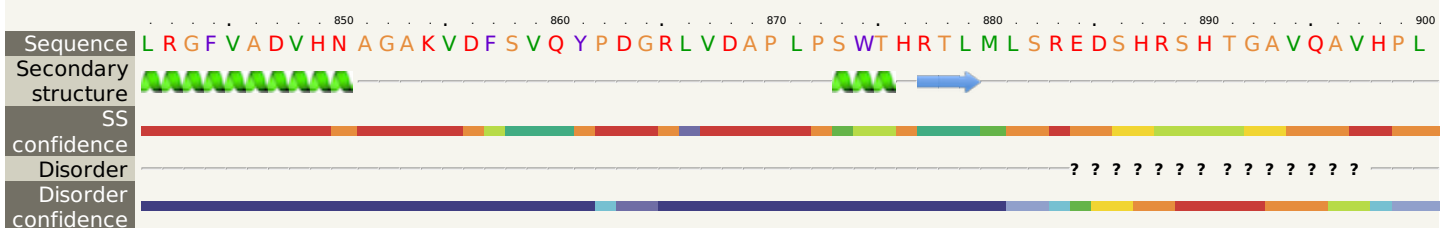
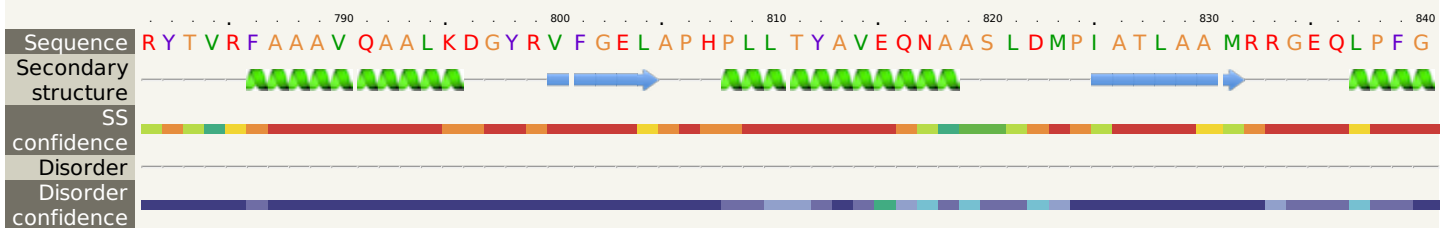
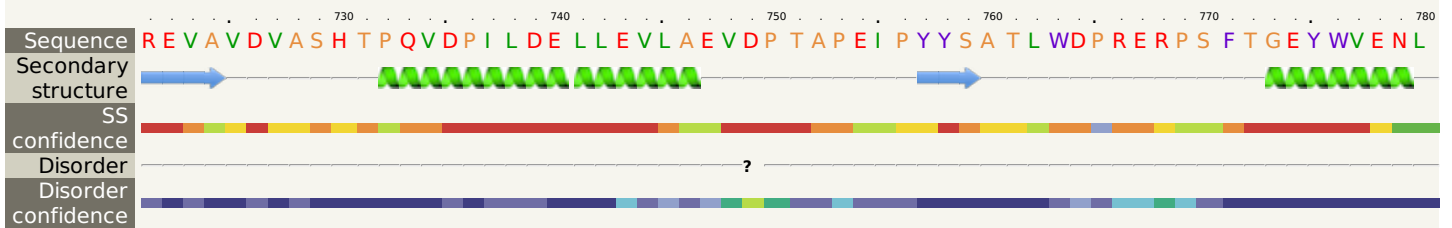
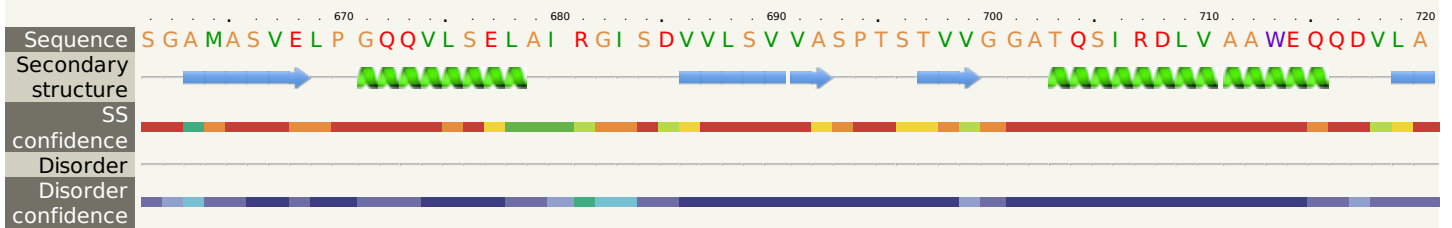
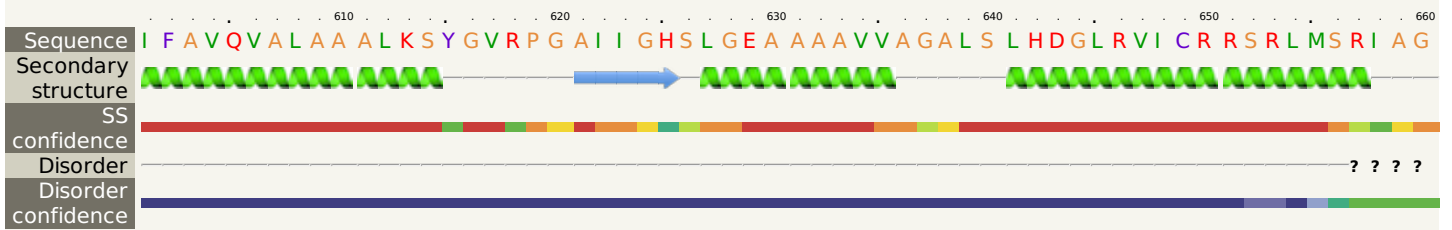
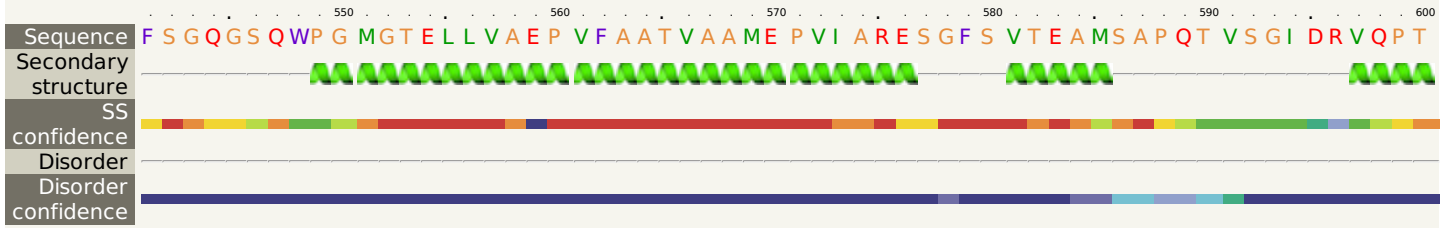
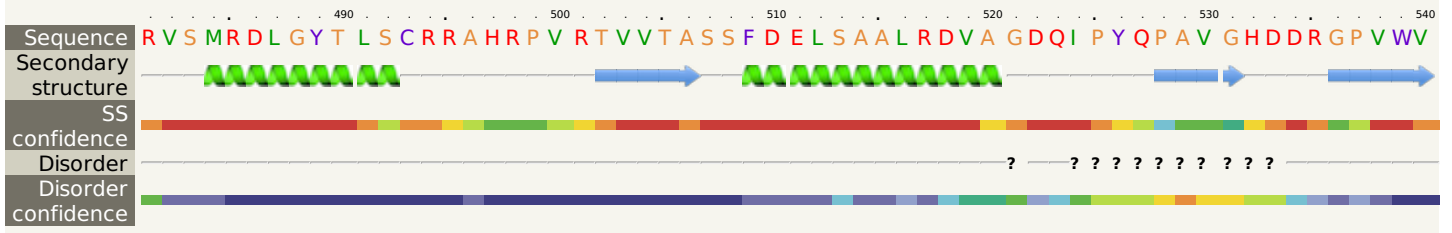
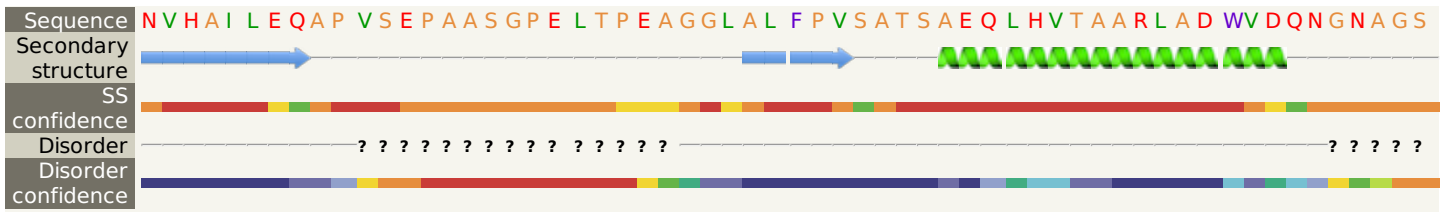


# Phyre2

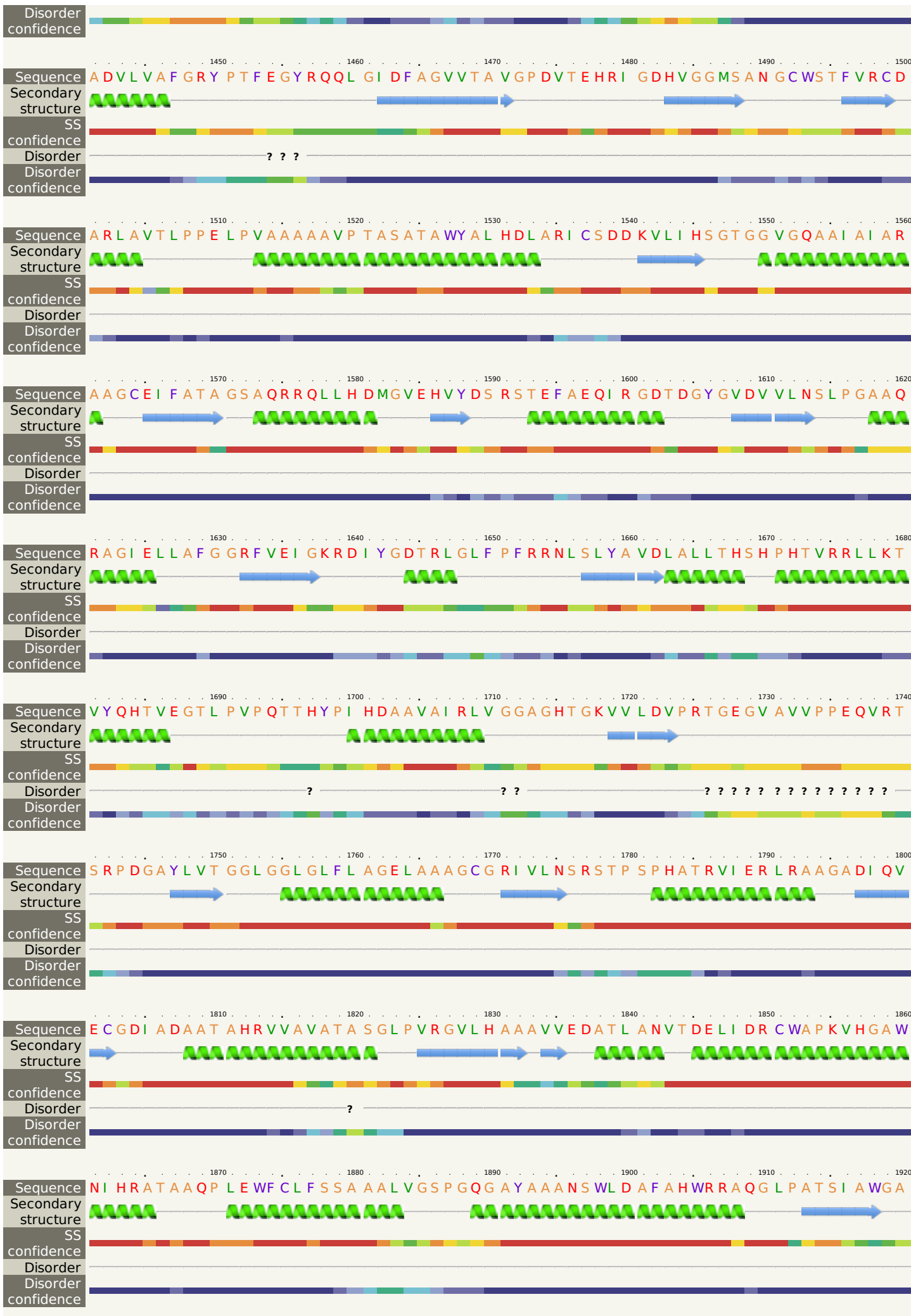
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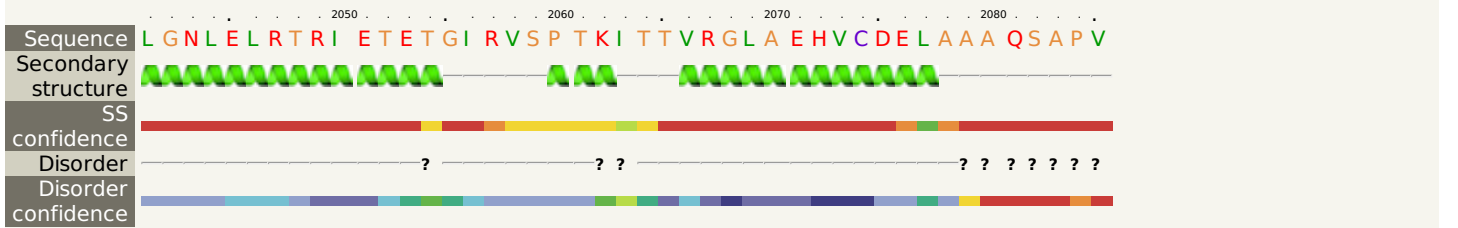
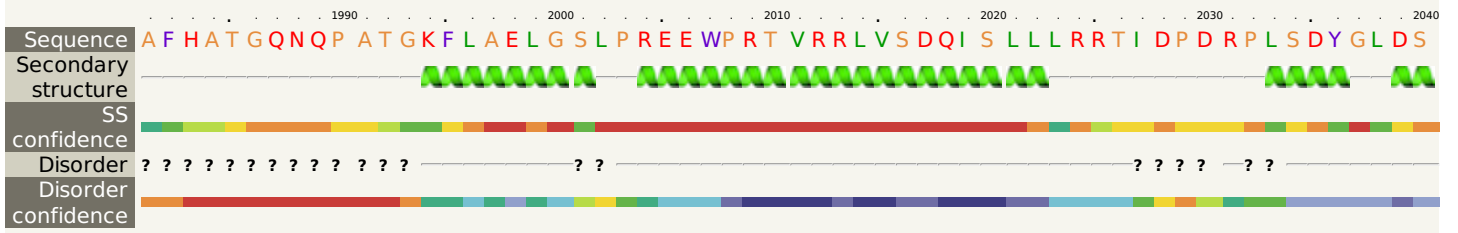
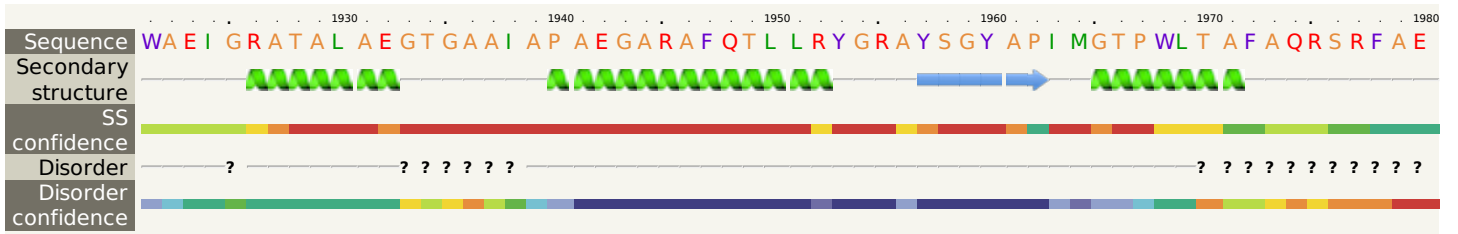
## Secondary structure and disorder prediction











Confidence Key

High(9) Low (0)

? Disordered ( 10%)

Alpha helix ( 36%)

Beta strand ( 19%)

TM helix ( 1%)