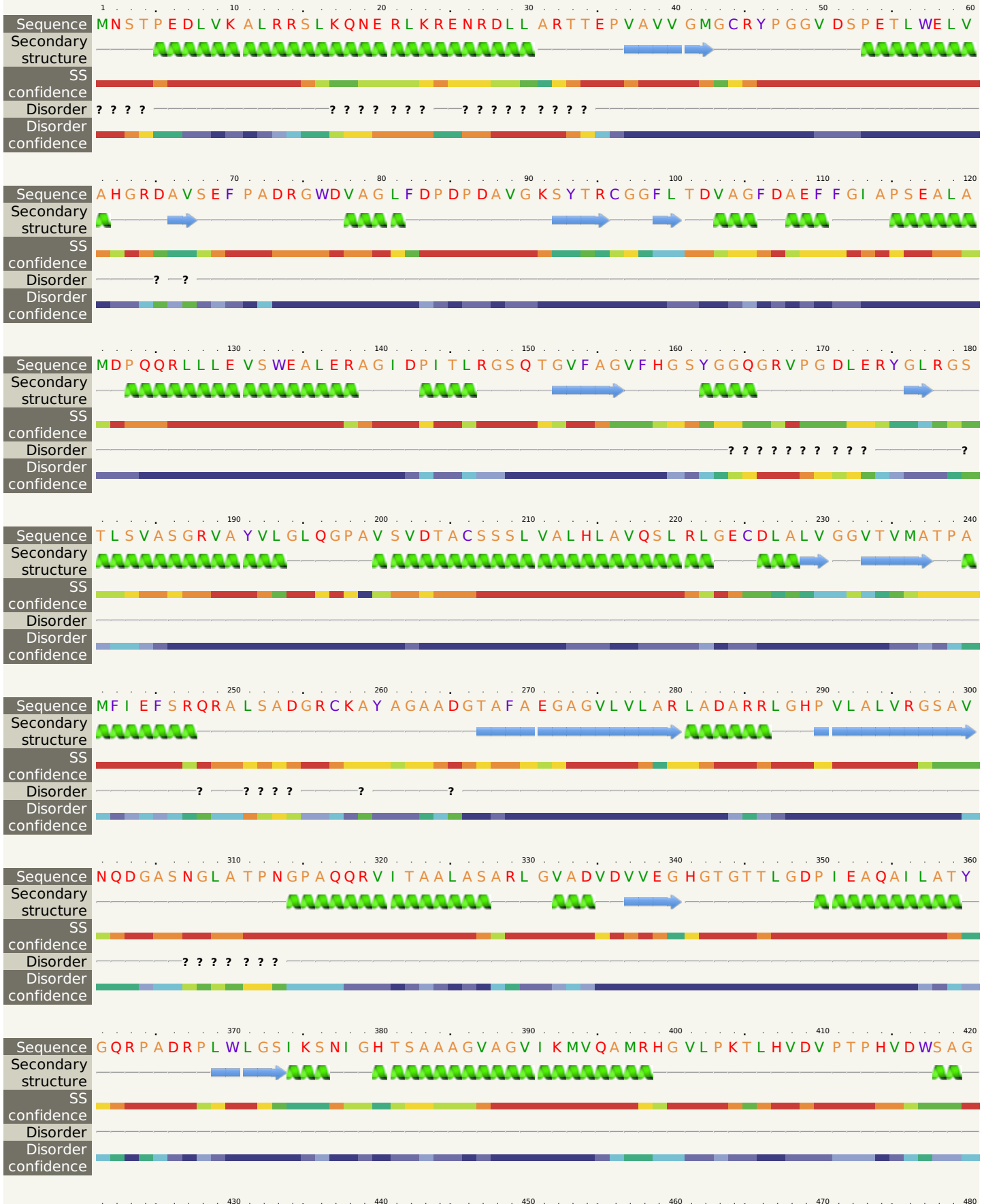
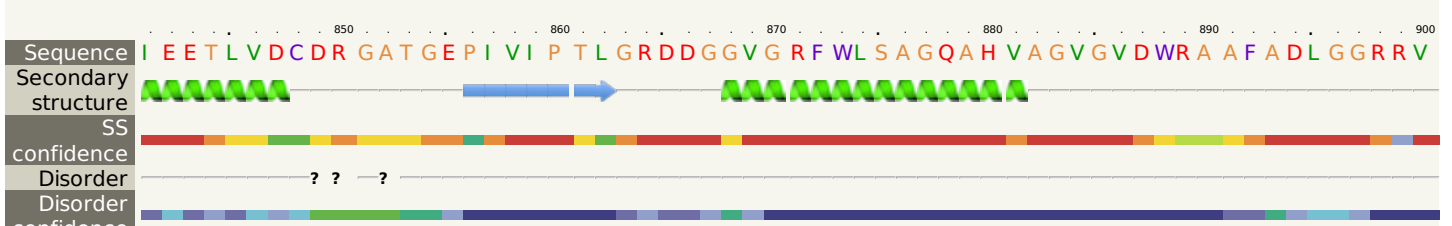
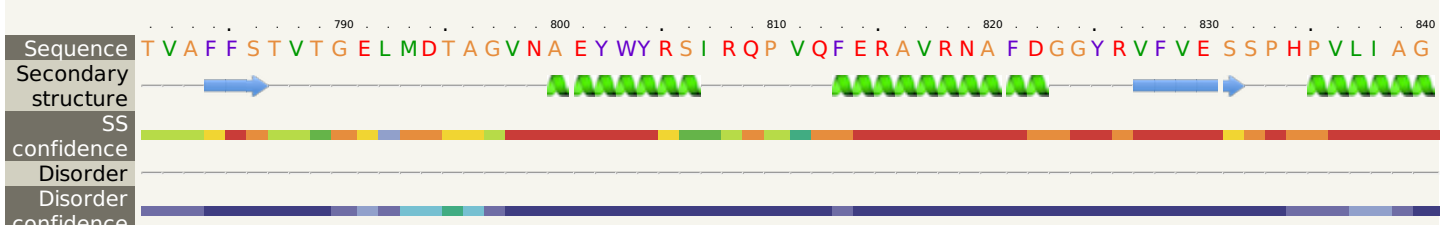
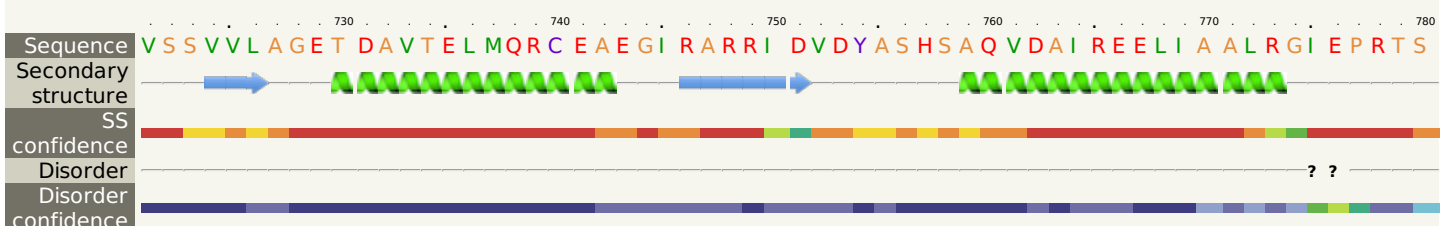
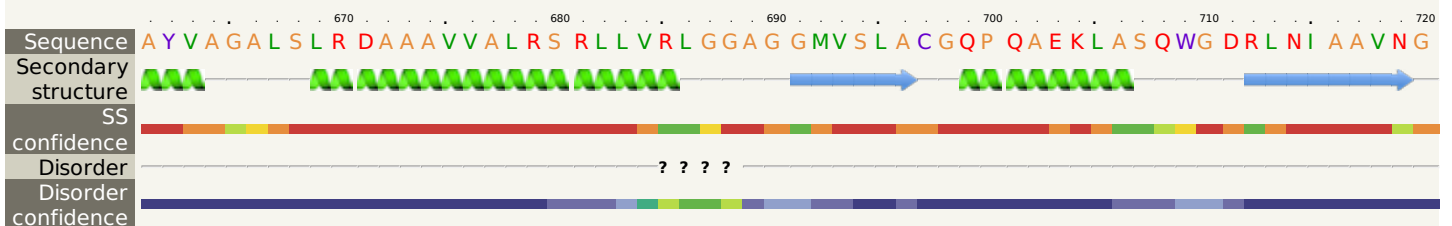
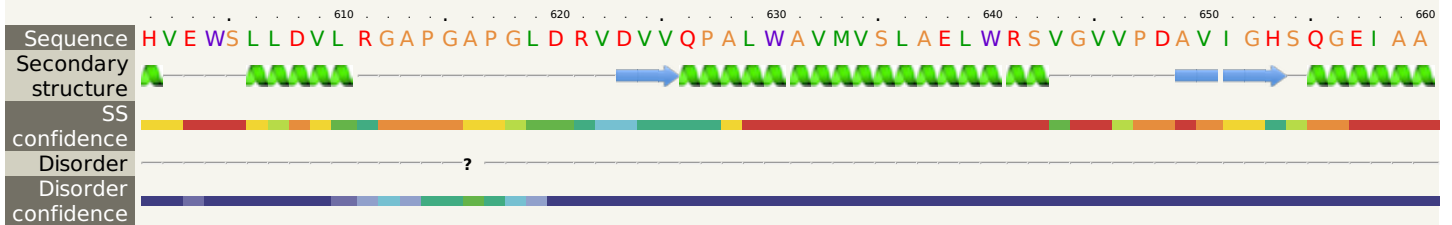
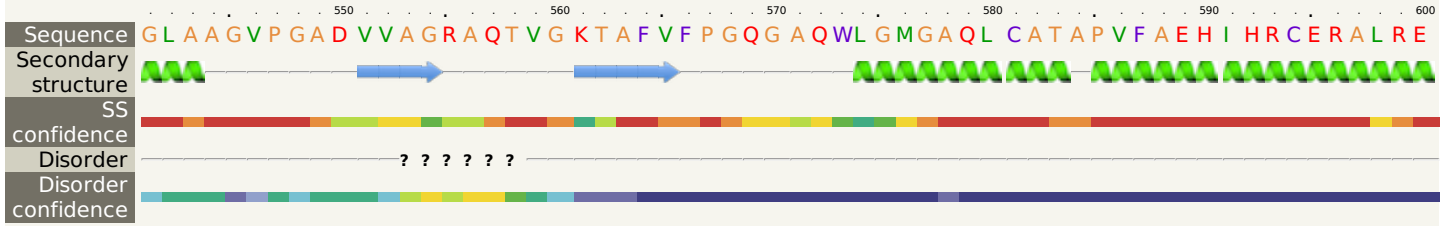
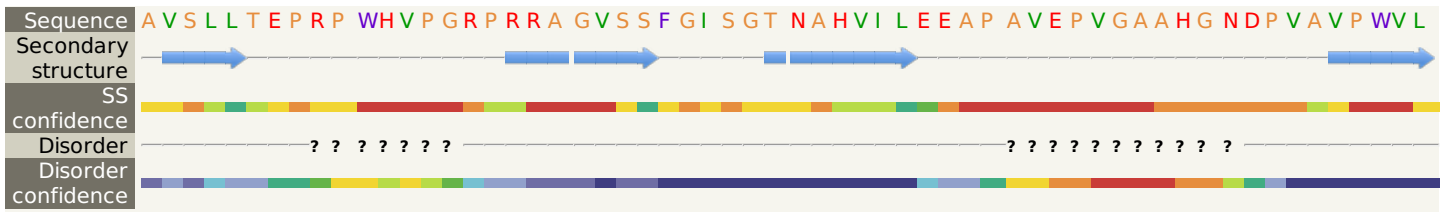


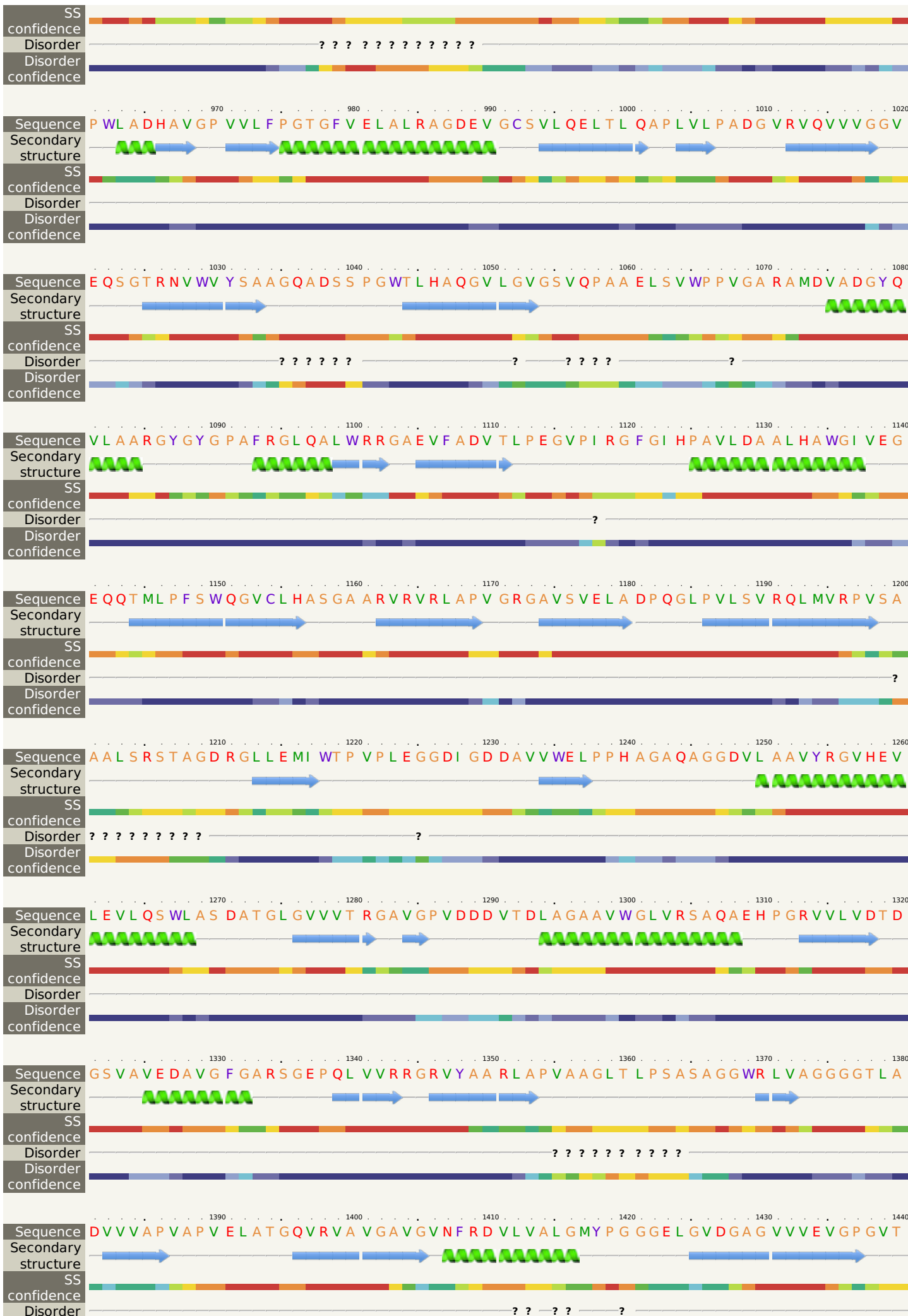
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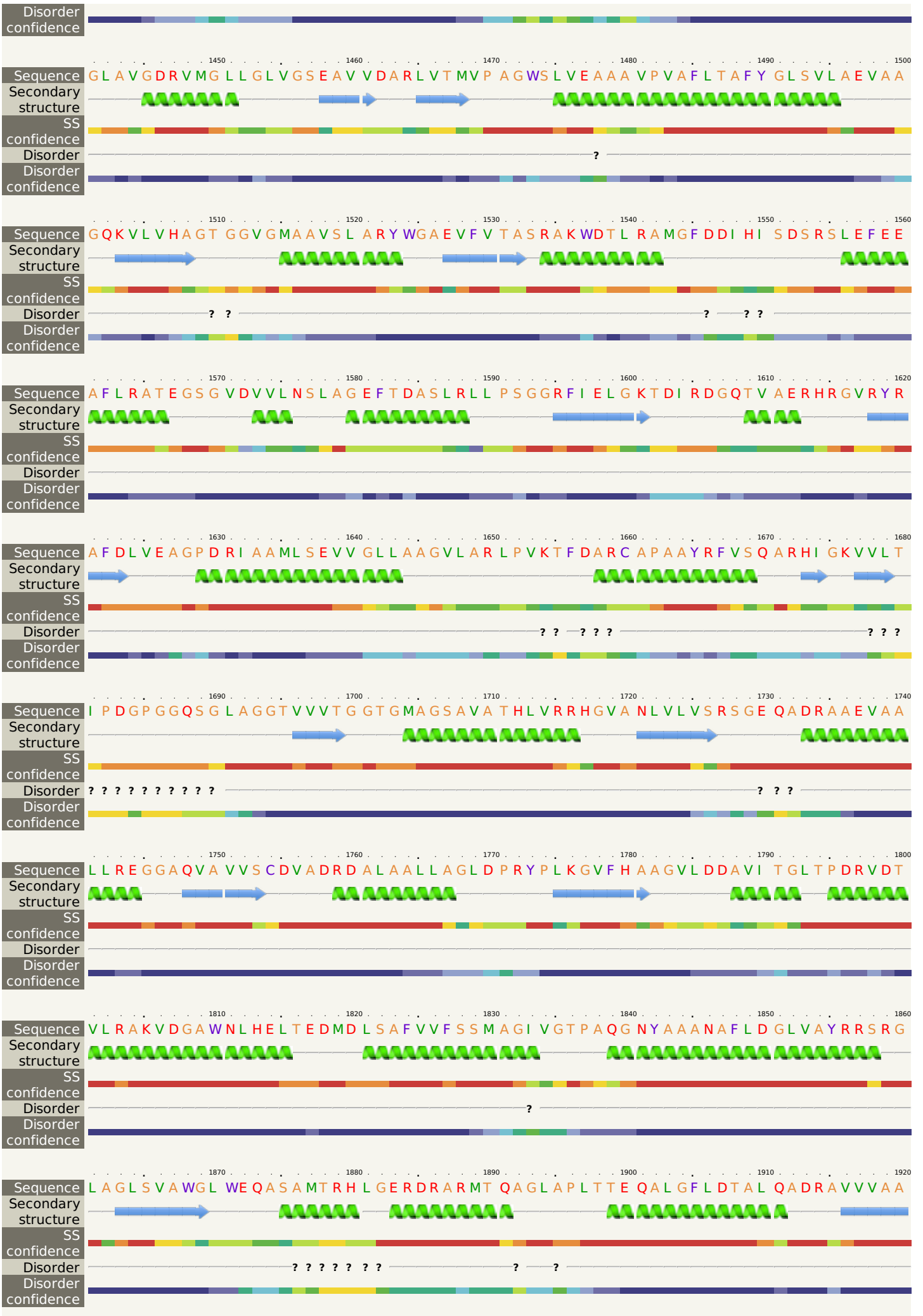
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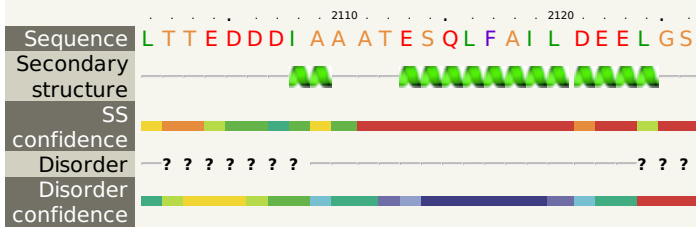
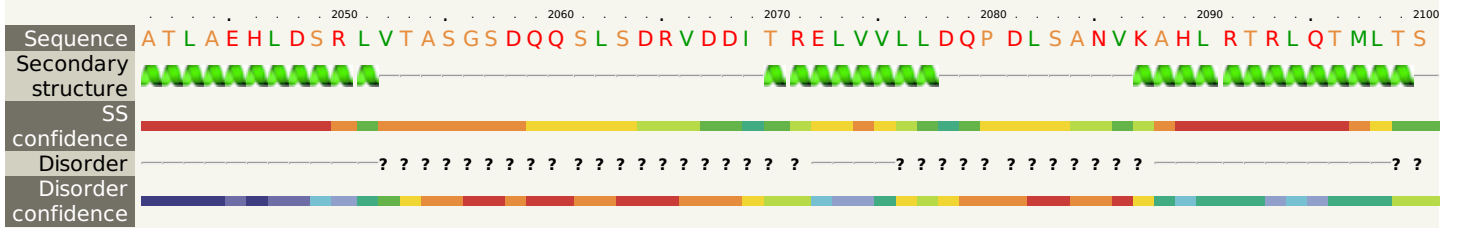
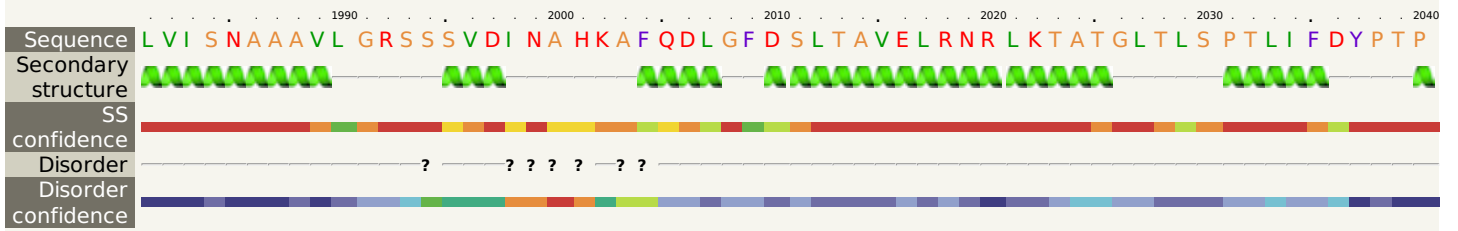
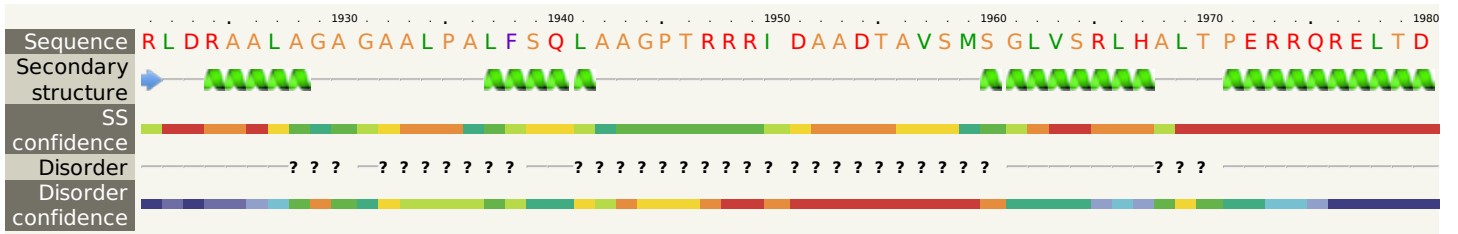
Secondary structure and disorder prediction














Confidence Key
 High(9)  Low (0)
 ? Disordered (12%)
 Alpha helix (38%)
 Beta strand (18%)