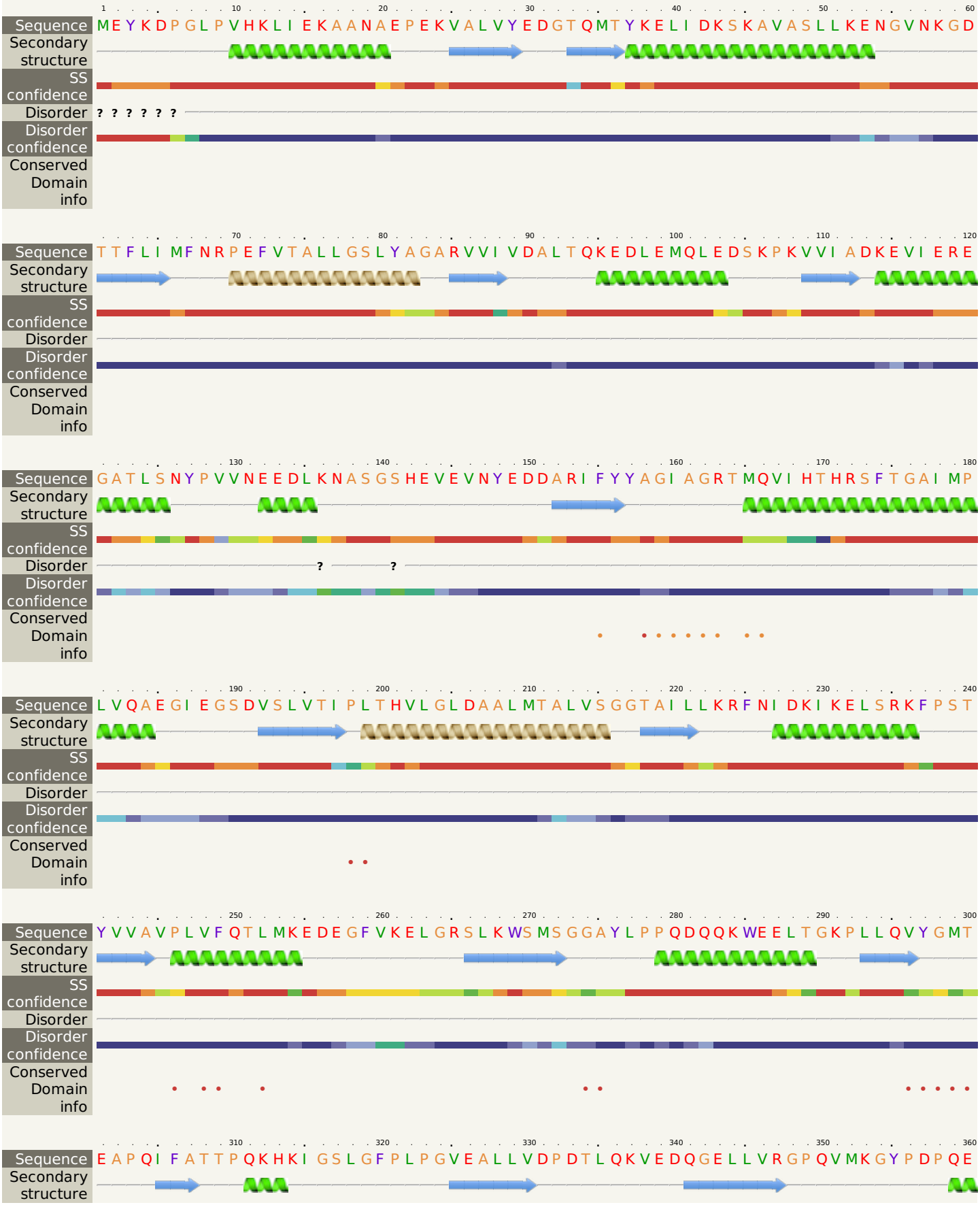
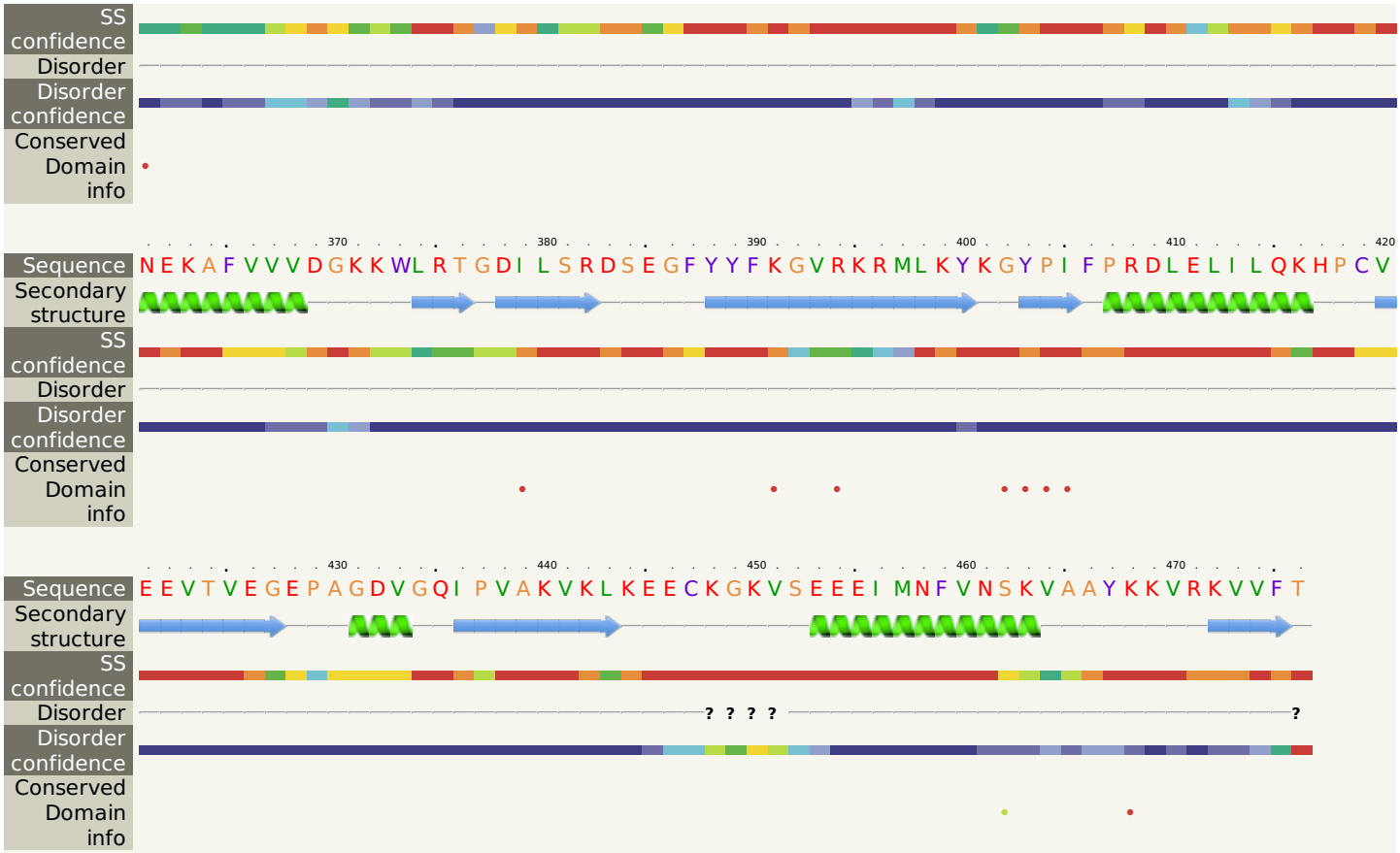
Phyre2

Email	Jennifer.Flechsler@ur.de
Description	4hbcoa_islandicus_00941_0379
Date	Fri Feb 27 14:13:32 GMT 2015
Unique Job ID	0a05811bb2f434b2

Secondary structure and disorder prediction





Confidence Key

High(9) Low (0)

CDD sites

- AMP binding site
- active site
- CoA binding site
- acyl-activating enzyme (AAE) consensus motif

? Disordered (3%)

Alpha helix (36%)

Beta strand (24%)

TM helix (11%)