Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD0384c_(clpB)_459457_462003
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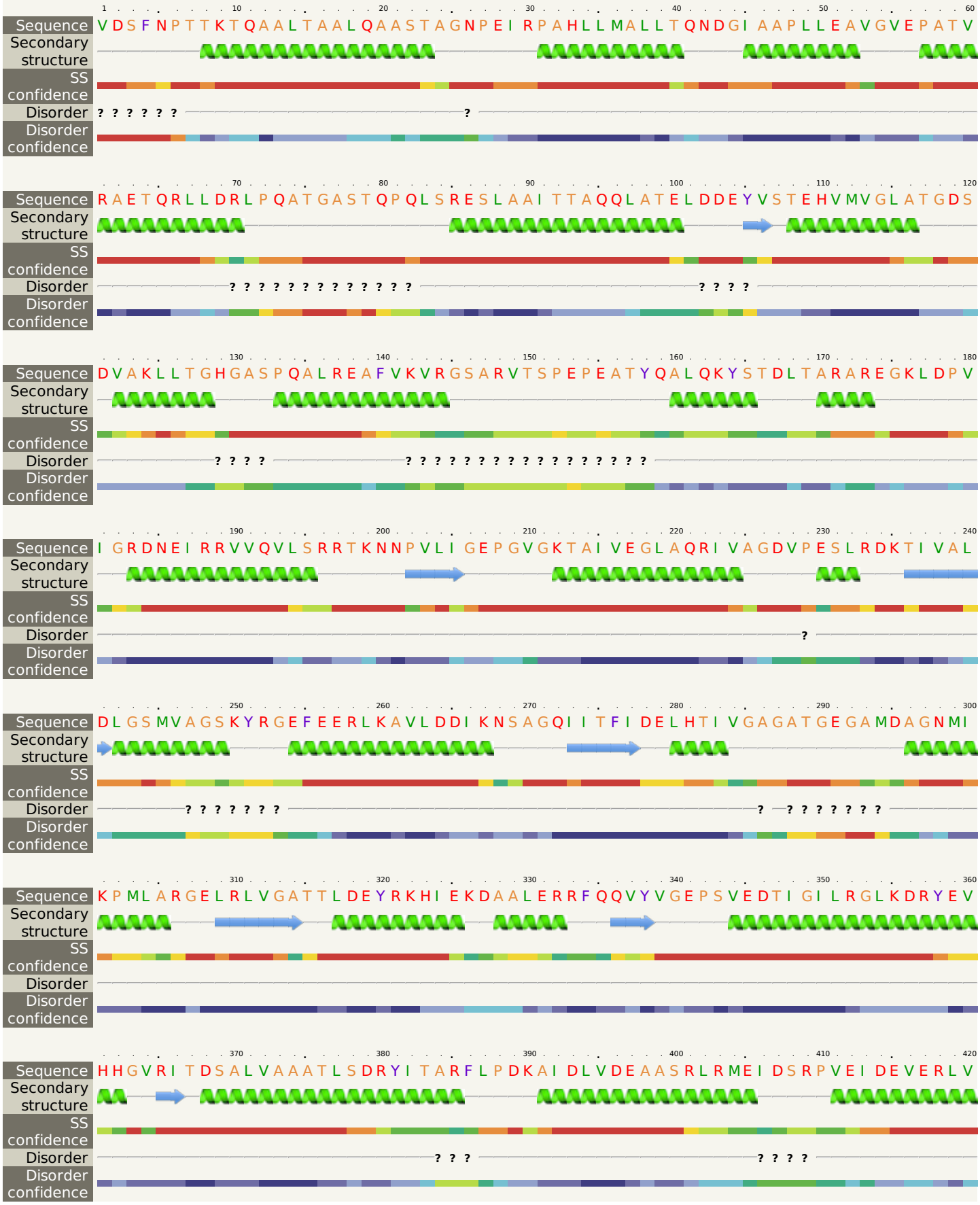
Secondary structure and disorder prediction



Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
RRLEI EEMALSKEEDEASAERLAKLRSELADQKEKLAELTTRWQNEKNAI EI VRDLKEQL
EALRGESERAERDGDLAKAAELRYGRI PEVEKKLDAALPQAQAREQVMLKEEVGPDDI AD
VVSAWTGI PAGRLLEGETAKLLRMEDELGKRVI GQKAAVTAVSDAVRRSRAGVSDPNRPT
GAFMFLGPTGVGKTELAKALADFLFDDERAMVRI DMSEYGEKHTVARLI GAPPGYVGYEA
GGQLTEAVRRRPYTVVLFDEI EKAHPDVFDVLLQVLDEGRLTDGHGRTVDFRNTI LI LTS
NLGSGGSAEQVLAAVRATFKPEFI NRLDDVLI FEGLNPEELVRI VDI QLAQLGKRLAQRR
LQLQVSLPAKRWLAQRGFDPVYGARPLRRLVQQAI GDQLAKMLLAGQVHDGDTVPVNVSP
DADSLI LG

Confidence Key

High(9) Low (0)

? Disordered (15%)

Alpha helix (59%)

Beta strand (9%)