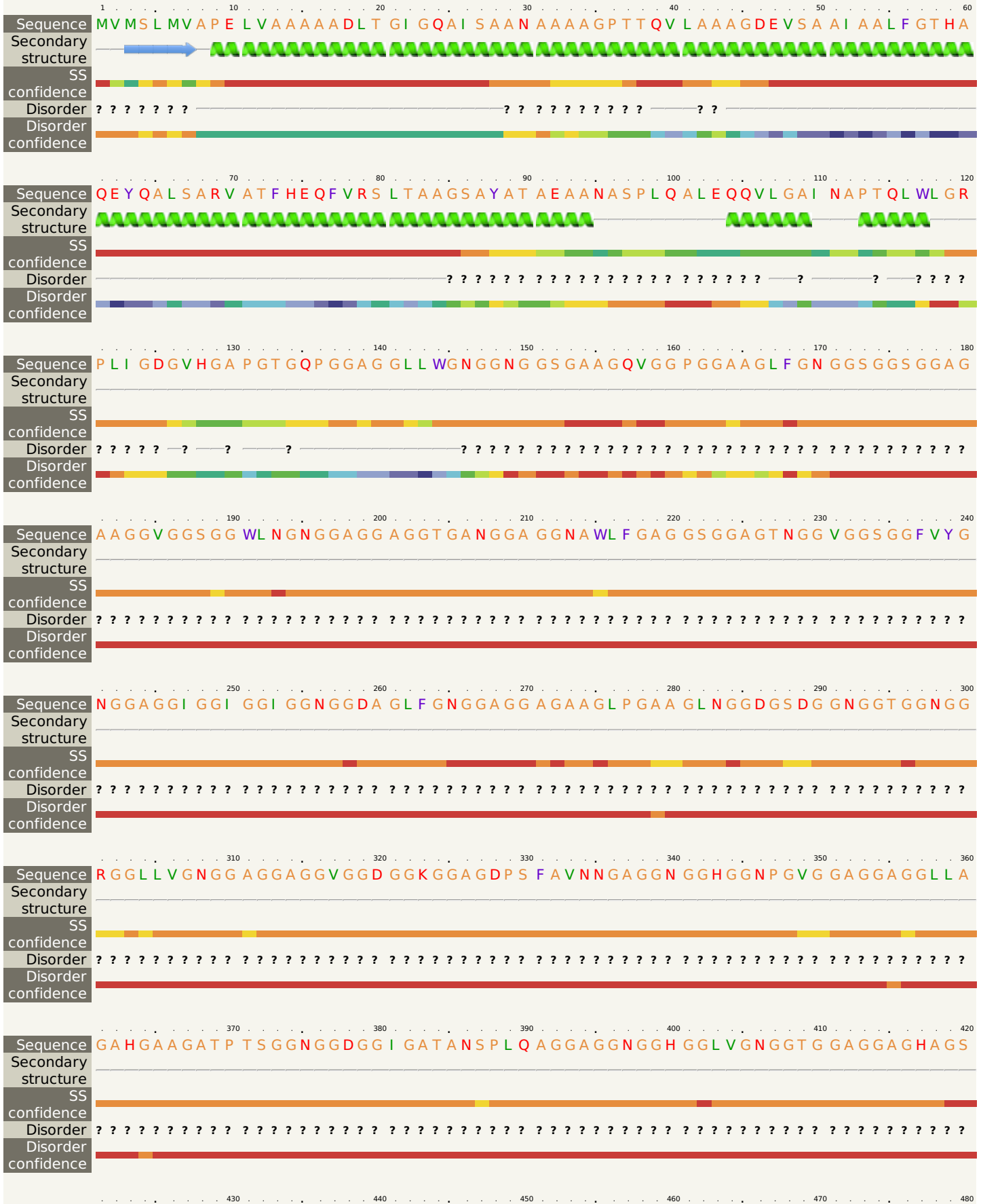


Phyre2

Email mdejesus@rockefeller.edu
Description RVBD3345c_(PE_PGRS50)_3738329_3742945
Date Thu Aug 8 16:20:55 BST 2019
Unique Job ID 9ea81385472b6237

Secondary structure and disorder prediction



Sequence T G A T G T A L Q P T G G N G T N G G A G G H G G N G G N G G A Q H G D G G V G G K G G A G G S G G A G G N G F D A A T
Secondary structure
SS confidence
Disorder
Disorder confidence

490 500 510 520 530 540
Sequence L G S P G A D G G M G G N G G K G G D G G K A G D G G A G A A G D V T L A V N Q G A G G D G G N G G E V G V G G K G G A
Secondary structure
SS confidence
Disorder
Disorder confidence

550 560 570 580 590 600
Sequence G G V S A N P A L N G S A G A N G T A P T S G G N G G N G G A G A T P T V A G E N G G A G G N G G H G G S V G N G G A G
Secondary structure
SS confidence
Disorder
Disorder confidence

610 620 630 640 650 660
Sequence G A G G N G V A G T G L A L N G G N G G N G G I G G N G G S A A G T G G D G G K G G N G G A G A N G Q D F S A S A N G A
Secondary structure
SS confidence
Disorder
Disorder confidence

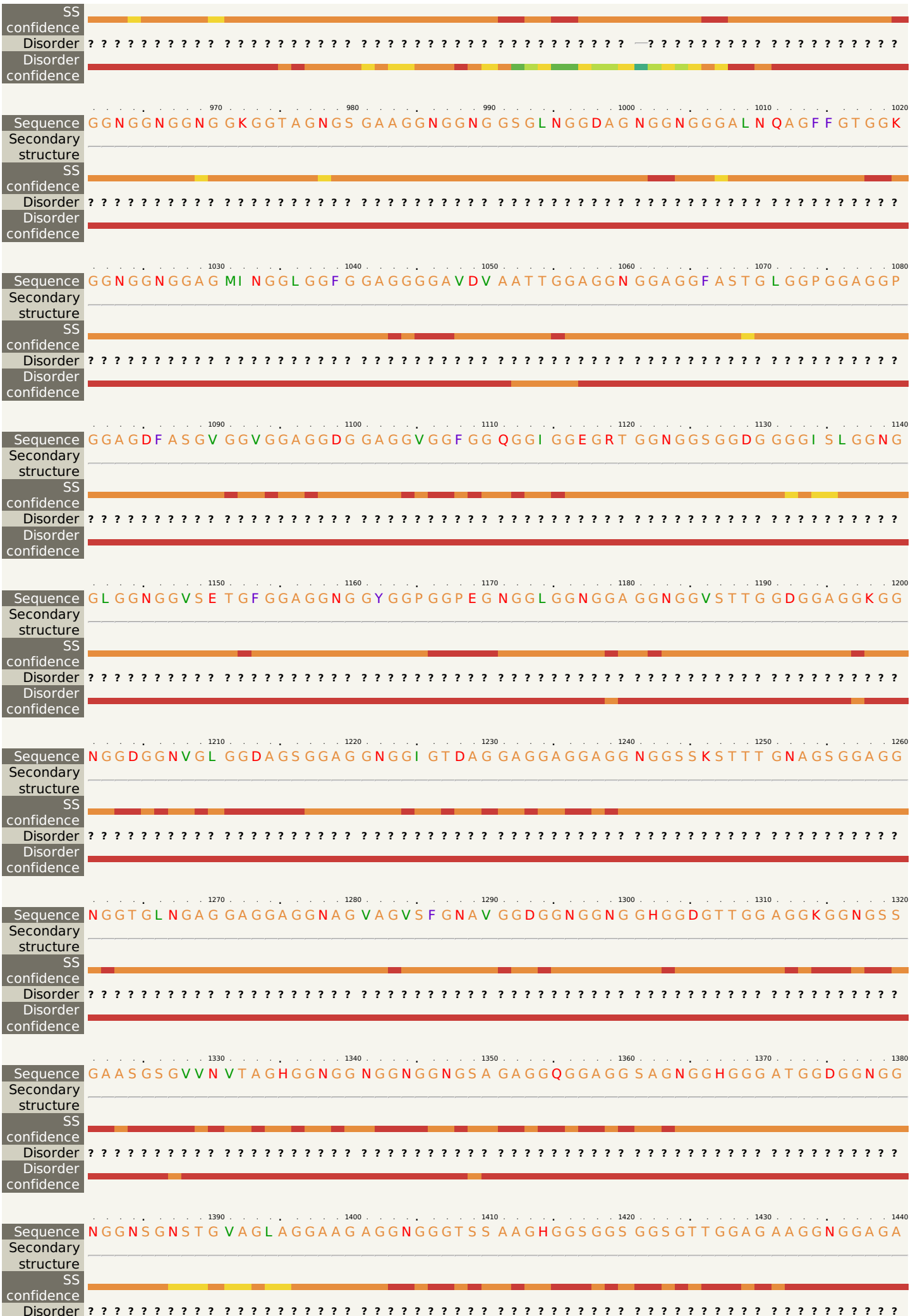
670 680 690 700 710 720
Sequence N G G Q G G N G G N G G I G G K G G D A F A T F A K A G N G G A G G N G G N V G V A G Q G G A G G K G A I P A M K G A T
Secondary structure
SS confidence
Disorder
Disorder confidence

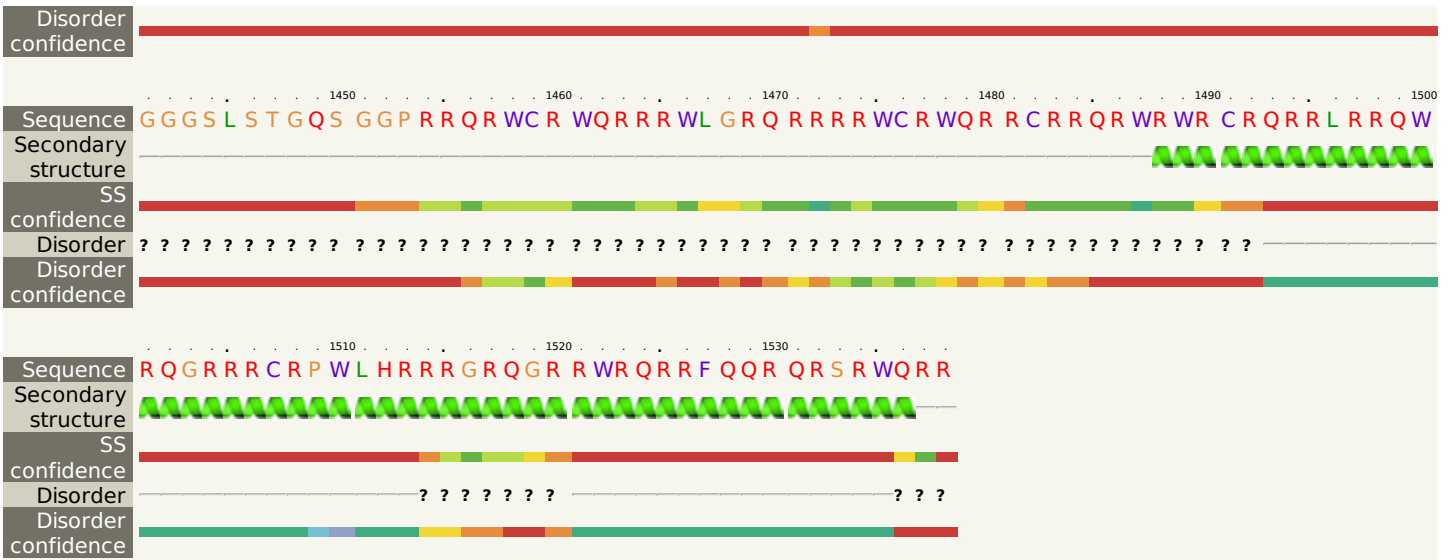
730 740 750 760 770 780
Sequence G A D G T A P T S G G D G G N G G N G A S P T V A G G N G G D G G K G G S G G N V G N G G N G G A G G N G A A G Q A G T
Secondary structure
SS confidence
Disorder
Disorder confidence

790 800 810 820 830 840
Sequence P G P T S G D S G T S G T D G G A G G N G G A G G A G G T L A G H G G N G G K G G N G G Q G G I G G A G E R G A D G A G
Secondary structure
SS confidence
Disorder
Disorder confidence

850 860 870 880 890 900
Sequence P N A N G A N G E N G G S G G N G G D G G A G G N G G A G G K A Q A A G Y T D G A T G T G G D G G N G G D G G K A G D G
Secondary structure
SS confidence
Disorder
Disorder confidence

910 920 930 940 950 960
Sequence G A G E N G L N S G A M L P G G G T V G N P G T G G N G G N G G N A G V G G T G G K A G T G S L T G L D G T D G I T P N
Secondary structure





Confidence Key
 High(9) [red bar] [orange bar] [yellow bar] [green bar] [light green bar] [light blue bar] [dark blue bar] Low (0)
 ? Disordered (92%)
 AA Alpha helix (9%)
 → Beta strand (0%)