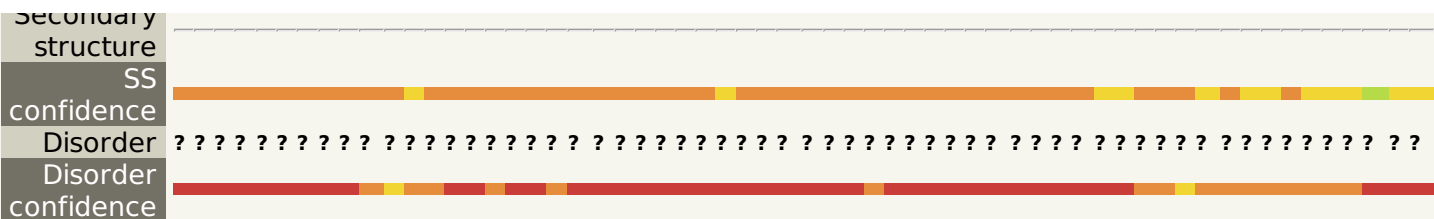


Phyre2

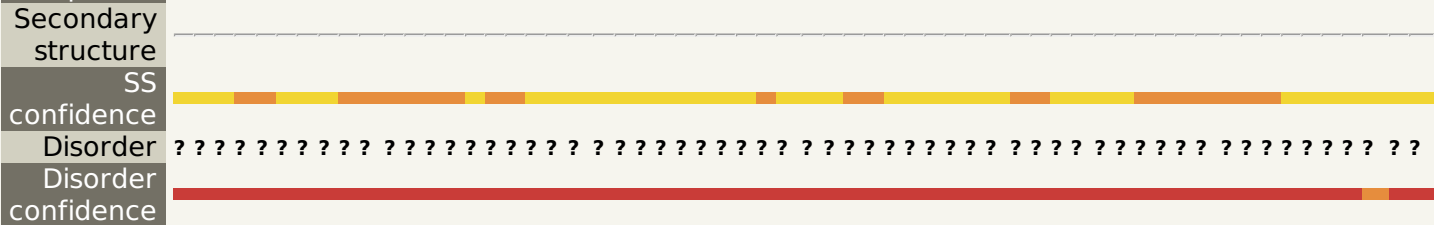
| | |
|---------------|------------------------------------|
| Email | mdejesus@rockefeller.edu |
| Description | RVBD0833_(PE_PGRS13)_925364_927613 |
| Date | Fri Jul 26 01:50:42 BST 2019 |
| Unique Job ID | 0c1277b4a26c86a7 |

Secondary structure and disorder prediction

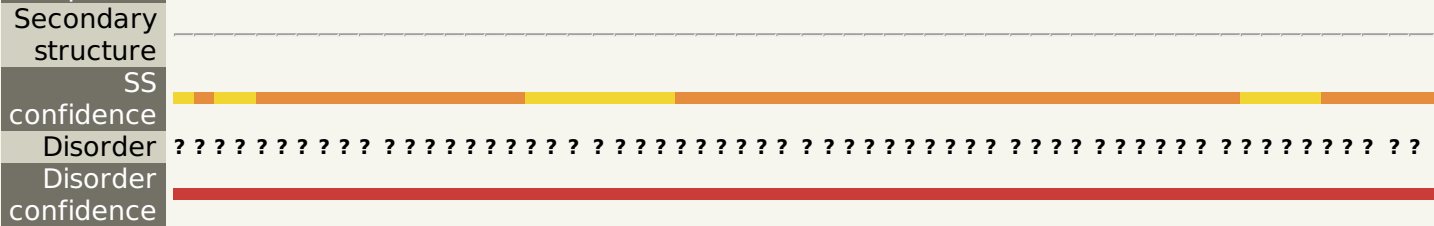




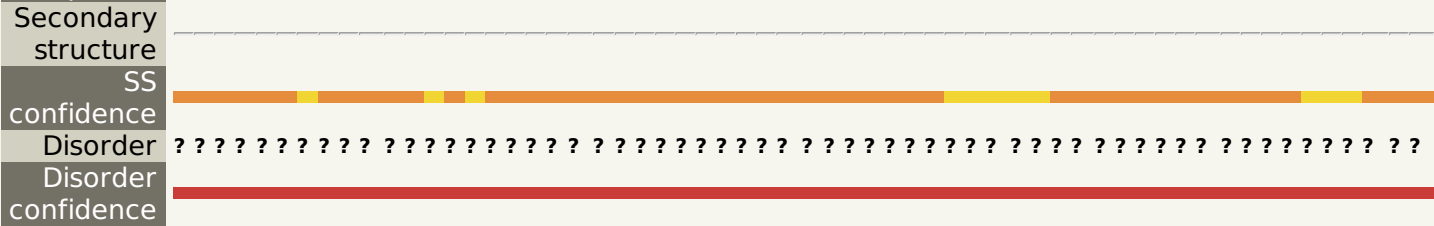
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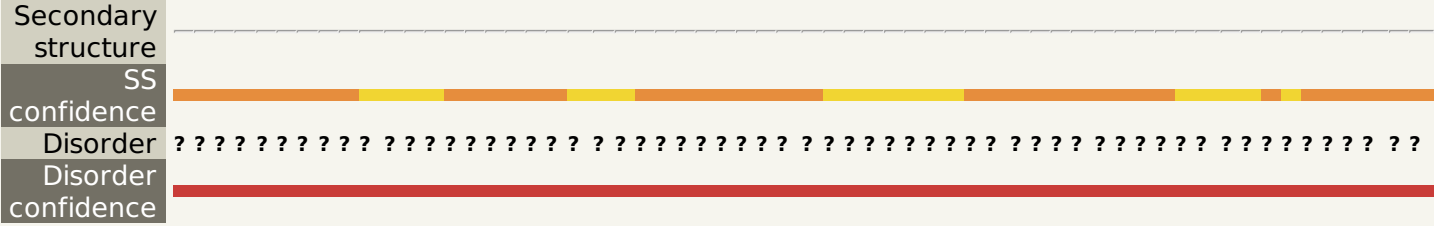
Sequence GGGGTGSGV SGGAGGDGGA GGI L F GAGGA GGAGGAVTGT GATGSGSGAG GGALLFGA GG



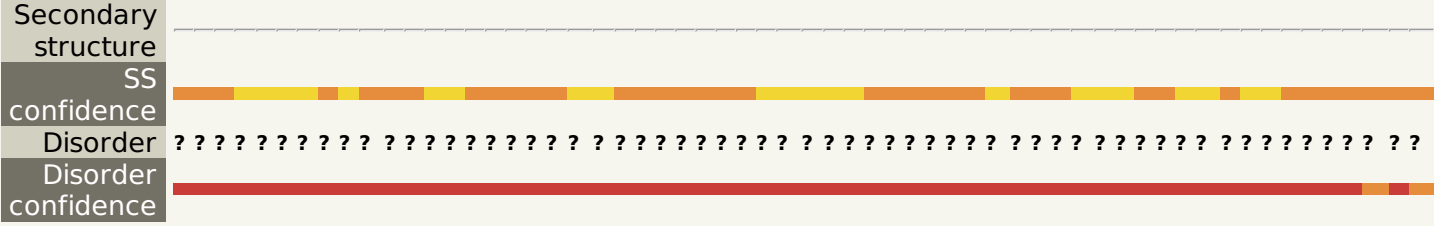
Sequence AGGAGGSSGI GGF AAGGAGG P GGAGGLF NG GGAGGAGGSGV SGGAGGEGG AGGAGGLF AG



Sequence GGAGGAGGSG NNVGGAGGAG GVGGGLF GAGGAGGSGGGGSGV AGDGGAGGNA GLLAPGLA GG

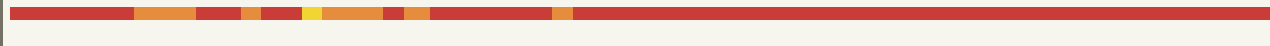


Sequence AGGGGGQGF DTGGAGGPGGDAGLLVGS GG V GGAGGFGLTTGGPGAAGGDA GLLFGSGGAG



Sequence GAGGSGRTDL GGAGGAGGKA GLI GNNGGAGGAGGNGGG DGGP GGAAFGLGNNGG NG

Disorder
confidence



..... 730 740

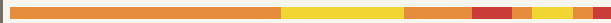
Sequence

GTGT SAGSPGAGGAGGSLI GA EGPLPGLLP

Secondary
structure



SS
confidence



Disorder

????????????????????????????????

Disorder
confidence



Confidence Key
 High(9) Low (0)
 ? Disordered (100%)
 Alpha helix (0%)
 Beta strand (0%)