

Phyre2

Email: mdejesus@rockefeller.edu
 Description: RVBD3464_(rmlB)_3882016_3883011
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 Unique Job ID: 35dbc7705bcf2d3f

Secondary structure and disorder prediction



Confidence Key
 High(9) Low (0)
 ? Disordered (4%)
 Alpha helix (47%)
 Beta strand (16%)
 TM helix (5%)

