

# Phyre2

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 Description: RVBD3141\_(fadB4)\_3508092\_3509063  
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## Secondary structure and disorder prediction



Confidence Key  
 High(9) [Colorful bar] Low (0)  
 ? Disordered ( 4%)  
 AA Alpha helix ( 36%)  
 > Beta strand ( 27%)  
 TM TM helix ( 5%)

