

Phyre2

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 Description: RVBD0527_(ccdA)_617496_618275
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Secondary structure and disorder prediction



Confidence Key
 High(9) [red bar] [orange bar] [yellow bar] [green bar] [blue bar] Low (0)
 ? Disordered (25%)
 Alpha helix (87%)
 Beta strand (0%)
 TM helix (60%)