





Phyre2

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Description: RVBD1966_(mce3A)_2209335_2210612
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Secondary structure and disorder prediction



Sequence	A G E Q P
Secondary structure	_____
SS confidence	██████████
Disorder	? ? ? ? ?
Disorder confidence	██████████

Confidence Key
 High(9)  Low (0)
 ? Disordered (26%)
 Alpha helix (44%)
 Beta strand (14%)
 TM helix (5%)