Phyre²

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Description	RhoE_____
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Secondary structure and disorder prediction

Sequence (1–60)
MKERRASQKLSSKSIMDPNQNVKCKIVVVGDSQCGKTALLHVFAKDCFPENYVPTVFENY

Secondary structure

SS confidence

Disorder: ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ? ?

Disorder confidence

Sequence (61–120)
TASFEIDTQRIELSLWDTSGSPYYDNVRPLSYPDSDAVLICFDISRPETLDSVLKKWKGE

Secondary structure

SS confidence

Disorder

Disorder confidence

Sequence (121–180)
IQEFCPNTKMLLVGCKSDLRTDVSTLVELSNHRQTPVSYDQGANMAKQIGAATYIECSAL

Secondary structure

SS confidence

Disorder: ? ? ? ? ? ? ? ? ? ? ? ? ?

Disorder confidence

Sequence (181–240)
QSENSVRDIFHVATLACVNKTNKNVKRNKSQRATKRISHMPSRPELSAVATDLRKDKAKS

Secondary structure

SS confidence

Disorder: ?

Disorder confidence

Sequence (241–244)
CTVM

Secondary structure

SS confidence

Disorder: ? ? ? ?

Disorder confidence

Confidence Key

High(9) ▮▮▮▮▮▮▮▮▮ Low (0)

? Disordered (32%)

Alpha helix (33%)

Beta strand (20%)