

All Projects

2QAC mutated.fasta Created: today at 08:34

Summary Templates 50 Models 1

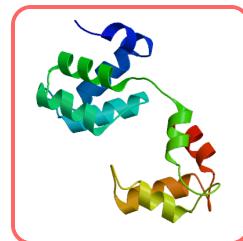
Project Data ▾

Order by: GMQE ▾

Model Results ⓘ



Model 01



Structure Assessment

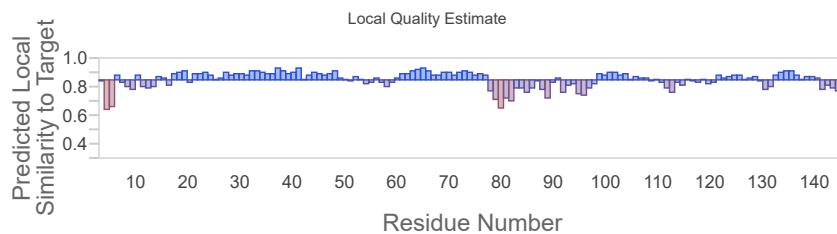
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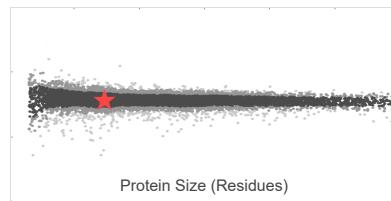
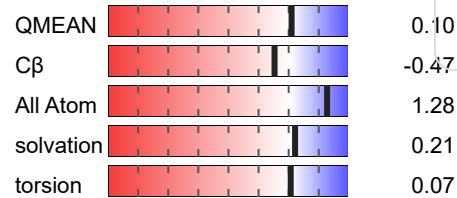
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Oligo-State
MonomerGMQE
0.91QMEANDisCo Global:
0.85 ± 0.07

QMEANDisCo Local



QMEAN Z-Scores



Template

2qac.1.A Myosin A tail domain interacting protein MTIP

The closed MTIP-MyosinA-tail complex from the malaria parasite invasion machinery

Seq Identity

99.31%

Coverage



Model-Template Alignment

Model_01	M E SVADIQQLEEEVDES DVRIYFNEKSSGGKISID	35
2qac.1.A	- - SVADIQQLEEKVD ESDVRIYFNEKSSGGKISID	35
Model_01	NASYNARKLGLAPS SIDEKKIKELYGDNLTYEQYL	70
2qac.1.A	NASYNARKLGLAPS SIDEKKIKELYGDNLTYEQYL	70
Model_01	EYLSTICVHDKDNEEELIKMFAHFDNNCTGYLTKSQ	105
2qac.1.A	EYLSTICVHDKDNEEELIKMFAHFDNNCTGYLTKSQ	105
Model_01	MKNILTTWGDALT DQE AIDALNAFSSEDNIDYKLF	140
2qac.1.A	MKNILTTWGDALT DQE AIDALNAFSSEDNIDYKLF	140
Model_01	CEDILQ	146
2qac.1.A	CEDILQ	146