

All Projects

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

Summary    Templates 50    Models 1

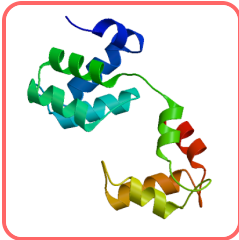
Project Data ▼

## Model Results ⓘ

Order by: GMQE ▼





### Model 01



Structure Assessment

Compare ☐

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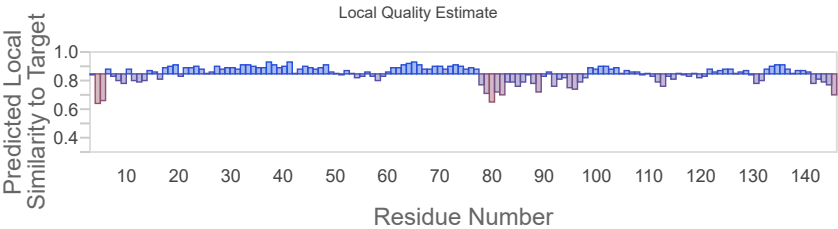
Display files 

Oligo-State  
Monomer

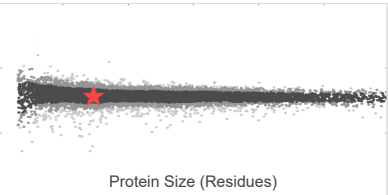
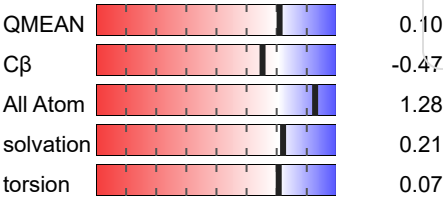
GMQE  
0.91

QMEANDisCo 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	MESVAD	IQQLEEE	VDESDVRIYFNEKS	SGGKISID	35
2qac.1.A	- - SVAD	IQQLEEK	VDESDVRIYFNEKS	SGGKISID	35
Model_01	NASYNARK	LGLAPSS	IDEKKIKELYGDN	ITYEQYL	70
2qac.1.A	NASYNARK	LGLAPSS	IDEKKIKELYGDN	ITYEQYL	70
Model_01	EYLSI	CVHDKDN	VEELIKMFAHFDNNCT	GYLTKSQ	105
2qac.1.A	EYLSI	CVHDKDN	VEELIKMFAHFDNNCT	GYLTKSQ	105
Model_01	MKNILTT	WGDALTD	QEAIDALNAPSS	EDNIDYKLE	140
2qac.1.A	MKNILTT	WGDALTD	QEAIDALNAPSS	EDNIDYKLE	140
Model_01	CED	ILQ			146
2qac.1.A	CED	ILQ			146