☰ All Projects

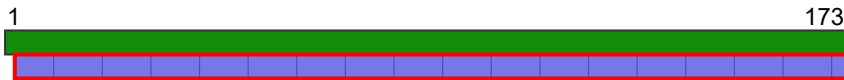# 7BF4 mutated.fasta Created: today at 12:42

Summary   Templates 16   Models 1   Project Data ▾
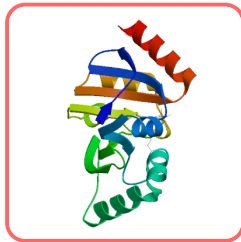
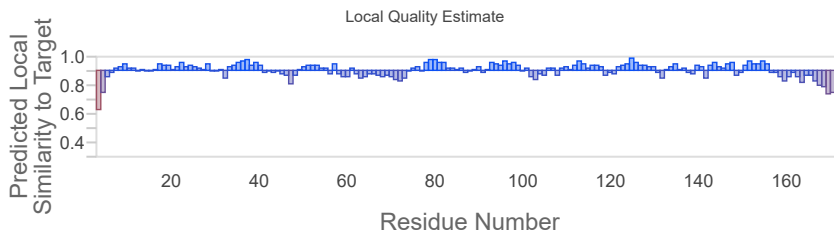## Model Results ❓

Order by: GMQE ▾
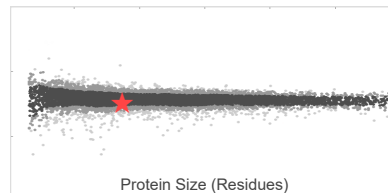

Model 01

Structure Assessment

Compare ☐

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Oligo-State	**GMQE**	**QMEANDisCo Global:**
Monomer	0.96	0.90 ± 0.07

Ligands
1 x APR ⬈ ✔

QMEANDisCo Local ⌃

QMEAN Z-Scores ⌃

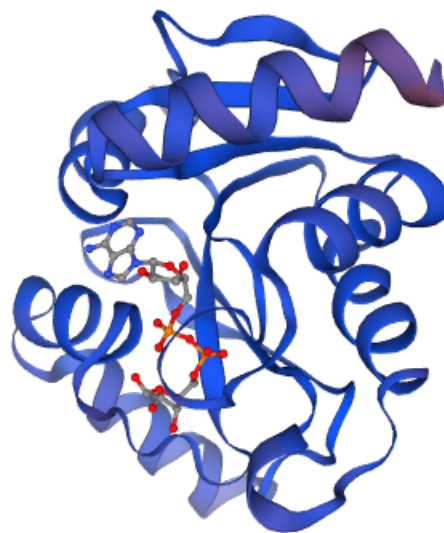| QMEAN | -0.19 |
Cβ	0.19
All Atom	0.50
solvation	0.10
torsion	-0.30

Template ⌄

6z5t.1.A Replicase polyprotein 1ab

SARS-CoV-2 Macrodomain in complex with ADP-ribose

Seq Identity
98.83%
Coverage

Model-Template Alignment ⌃

```
Model_01  GHMVNSFSGYLKLTDNVYIKNADIVEEAKKVKPTV   35
6z5t.1.A  --EVNSFSGYLKLTDNVYIKNADIVEEAKKVKPTV   35

Model_01  VVNAANVYLKHGGGVAAALNKATNNAMQVESDDYI   70
6z5t.1.A  VVNAANVYLKHGGGVAGALNKATNNAMQVESDDYI   70

Model_01  ATNGPLKVGGSCVLSGHNLAKHCLHVVGPNVNKGE  105
6z5t.1.A  ATNGPLKVGGSCVLSGHNLAKHCLHVVGPNVNKGE  105

Model_01  DIQLLKSAYENFNQHEVLLAPLLSAGIFGADPIHS  140
6z5t.1.A  DIQLLKSAYENFNQHEVLLAPLLSAGIFGADPIHS  140

Model_01  LRVCVDTVRTNVYLAVFDKNLYDKLVSSFLEMK    173
6z5t.1.A  LRVCVDTVRTNVYLAVFDKNLYDKLVSSFLEMK    173
```


| ✏ | ⚙ | Cartoon ▲ | 📷 ▲ | ▶ | ⏏ | ↻ |