All Projects

Untitled Project Created: today at 19:06

Summary Templates 49 Models 1 Project Data

Model Results

Order by: GMQE

Model 01

Structure Assessment

Compare

Download files Display files

Oligo-State	GMQE	QMEANDisCo Global:
Monomer	0.69	0.74 ± 0.05

QMEANDisCo Local

QMEAN Z-Scores

Template

2pe4.1.A Hyaluronidase-1

Seq Identity
42.82%

Coverage

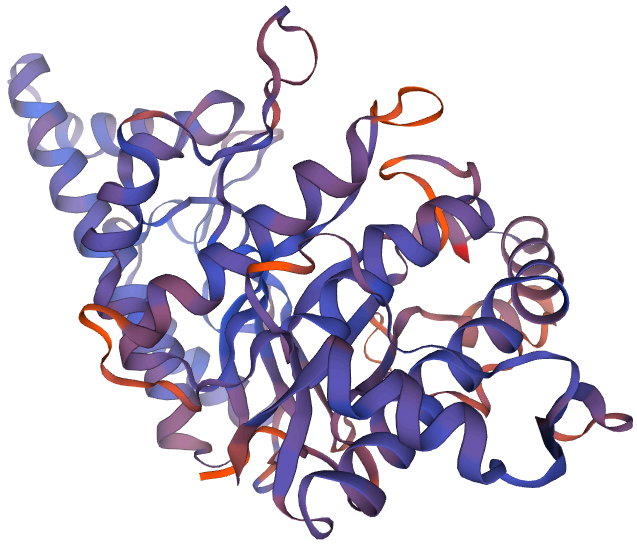
Model-Template Alignment

Halozyme EX2183
Merck v. Halozyme
PGR2025-00046

```
Model_01  LNFRAPPVIPNVPFLWAWNAPSEFCLGKFDEPLDM
2pe4.1.A  --FRGP-LIPNRPFTTVWNANTQWCLERHGVDVDV

Model_01  SLFSFIGSPRINATGQGVTIFYVDRLGYYPYIDSI
2pe4.1.A  SVFDVVANPGQTFRGPDMTIFYSSQLGTYPYY-TP

Model_01  TGVTVNGGIPQKISLQDHLDKAKKDITFYMPV-DN
2pe4.1.A  TGEPVFGGLPQNASLIAHLARTFQDILAAIPAPDF

Model_01  LGMAVIDWEEWRPTWARNWKPKDVYKNRSIELVQQ
2pe4.1.A  SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01  QNVQLSLTEATEKAKQEFEKAGKDFLVETIKLGKL
2pe4.1.A  QHPDWPAPQVEAVAQDQFQGAARAWMAGTLQLGRA

Model_01  LRPNHLWGYYLFPDCYNHHYKKPGYNGSCFNVEIK
2pe4.1.A  LRPRGLWGFYGFPDCYNYDFLSPNYTGQCPSGIRA

Model_01  RNDDLSWLWNESTALYPSIYL-NTQQSPVAATLYV
2pe4.1.A  QNDQLGWLWGQSRALYPSIYMPAVLEGTGKSQMYV

Model_01  RNRVREAIRVSKIPDAKSPLPVFAYTRIVFTDQVL
2pe4.1.A  QHRVAEAFRVA-VAAGDPNLPVLPYVQI-FYDTTN

Model_01  KFLSQDELVYTFGETVALGASGIVIWGTLSIMRSH
2pe4.1.A  HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

Model_01  KSCLLLDNYMETILNPYIINVTLAAKMCSQVLCQE
2pe4.1.A  ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01  QGVCIRKNWNSSDYLHLNPDNFAIQLE-KGGKFTV
2pe4.1.A  HGRCVRRTSHPKALLLLNPASFSIQLTPGGGPLSL

Model_01  RGKPTLEDLEQFSEKFYCSCYSTLSCKEKADVKDT
2pe4.1.A  RGALSLEDQAQMAVEFKCRCY--------------

Model_01  DAVDVCIADGVCIDAFLKPPMETEEPQIFY
2pe4.1.A  ------------------------------
```

MET 310 A

Confidence: 0.69



Cartoon