All Projects

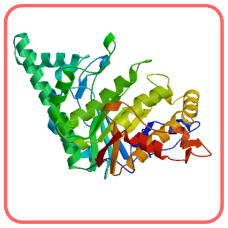
Untitled Project Created: today at 02:33

Summary Templates 45 Models 1 Project Data

Model Results

Order by: GMQE

Model 01



Structure Assessment

Compare

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Oligo-State
Monomer

GMQE
0.69

QMEANDisCo Global:
0.74 ± 0.05

QMEANDisCo Local

QMEAN Z-Scores

Template
2pe4.1.A Hyaluronidase-1

Seq Identity
43.07%

Coverage

Model-Template Alignment

Halozyme EX2189
Merck v. Halozyme
PGR2025-00039

```
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  LGMAVIDWEEWRPTWARNWKPKDVYKARSIELVQQ
2pe4.1.A  SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01  QNVQLSLTEATEKAKQEFEKAGKDFLVETIKLGKL
2pe4.1.A  QHPDWPAPQVEAVAQDQFQGAARAWMAGTLQLGRA

Model_01  LRPNHLWGYYLFPDCYNHHYKKPGYNGSCFNVEIK
2pe4.1.A  LRPRGLWGFYGFPDCYNYDFLSPNYTGQCPSGIRA

Model_01  RNDDLSWLWAESTALYPSIYL-NTQQSPVAATLYV
2pe4.1.A  QNDQLGWLWGQSRALYPSIYMPAVLEGTGKSQMYV

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

Model_01  KFLSQDELVYTFGETVALGASGIVIWGTLSIMRSM
2pe4.1.A  HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

Model_01  KSCLLLKNYMETILNPYIINVTLAAKMCSQVLCQE
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  ------------------------------
```