

Untitled Project Created: today at 19:46

Summary

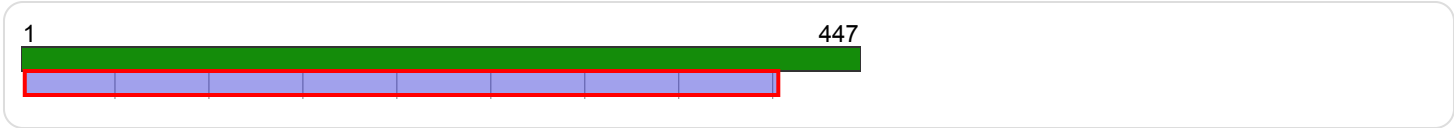
Templates 47

Models 1

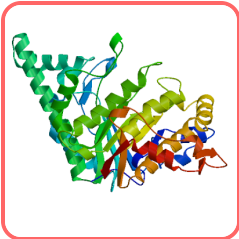
Project Data ▼

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
42.82%
Coverage

Model-Template Alignment

▲

Halozyyme EX2192
Merck v. Halozyyme
PGR2025-00052

Model_01 LNFRAPPVTPNVPFLWAWNPSEFCLGKFDEPLDM
 2pe4.1.A --FRGP-LLPNRPFTTVWVANTQWCLERHGVVDVDV

Model_01 SLFSFEGSPRINATGQGVTFYVDRLGYYPYIDSI
 2pe4.1.A SVFDVVANPGQTFRGPDMTIFYSSQLGTYPYY-TP

Model_01 TGVTVNGGIPQKISLQDHLDKAKKDITFYMPV-DN
 2pe4.1.A TGEIPVFGGLPQNASLIAHLARTEQDILAAIPAPDF

Model_01 LGMAVIDWEWRPTWARNWKPKDVYKNRSIELVQQ
 2pe4.1.A SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01 QNVQLSLTEATEKAKQEFEEKAGKDFLVETIKLGKL
 2pe4.1.A QHPDWPAPQVEAVAQDQFQGAARAWMAGTLQLGRA

Model_01 LRPNHLWGYYLFPDCYNHHYKKPGYNGSCFNVEIK
 2pe4.1.A LRPRGLWGFYGFPCYNYDFLSPNYTGQCPSGIRA

Model_01 RNDPLSWLWNESTALYPSIYLNTIQQSPVAATLYV
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTKSQMYV

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

Model_01 KFLSQDELVYTFGETVALGASGIVTWGTLSTIMRSP
 2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

Model_01 KSCLLLDNYMETILNPYIINVTLAAKMCSQVLCQE
 2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01 QGVCIRKNWNSSDYHLNPDNFAIQLE-KGGKFTV
 2pe4.1.A HGRCVVRTSHPKALLLNPAFSTIQLTPGGGPLSL

Model_01 RGKPTLEDLEQFSEKFYCSYSTLSCKEKADV KDT
 2pe4.1.A RGALSLEDQAQMAVEFKRCY-----

Model_01 DAVDVCIA DGVCIDAFLKPPMETEEPQIFY
 2pe4.1.A -----

