

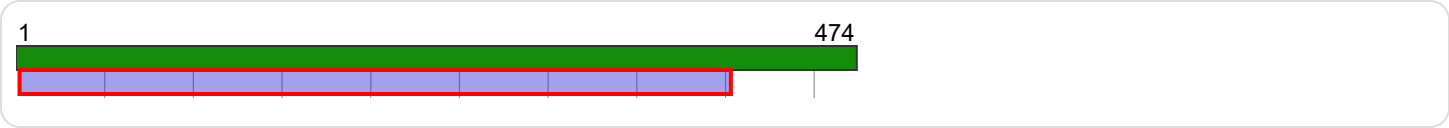
 All Projects

Untitled Project Created: today at 18:31

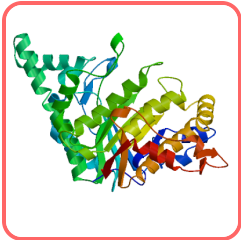
Summary Templates 37 Models 1 Project Data ▼

Model Results

Order by: GMQE ▼



Model 01



Structure Assessment

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Oligo-State
Monomer

GMQE
0.66

QMEANDisCo Global:
0.73 ± 0.05

QMEANDisCo Local
QMEAN Z-Scores

▼
▼

Template
2pe4.1.A Hyaluronidase-1

▼

Seq Identity
43.83%
Coverage



Model-Template Alignment

▲

Halozyyme EX2192
Merck v. Halozyyme
PGR2025-00053

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

Model_01 SLFSLTGSPRINVTGQGVTIFYVDRLGYYPYIDSI
 2pe4.1.A SVFDVVPANPGQTFRGPDMTIFYSSQLGTYPYY-TP

Model_01 TGVTVNGGIPQKISLQDHLDKAKQDITEYMPV-DN
 2pe4.1.A TGEIPVFGGLPQNASLIAHLARTFQDILAAIPAPDF

Model_01 LGMAVIDWEWRPTWARNWKPKDVYKNRSIELVQQ
 2pe4.1.A SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01 QNVQLSLAEATEKAKQEFEEKAGKDFMVETIKLGKL
 2pe4.1.A QHPDWPAQVEAVAQDQFGAARAWMAGTLQLGRA

Model_01 LRPNHLWGYLLFPDCYNHHYKKPGYNGSCFNVEIK
 2pe4.1.A LRPRLWGFYGFDCYNYDFLSPNYTGQCPSGIRA

Model_01 RNDLSWLWNESTALYPSIYLNTQQSPVAATLYV
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTGKSQMYV

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

Model_01 KFLSRDELVYTLGETVALGASGIVINGSLSIVRSM
 2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

Model_01 KSCLLLKNYMETILNPYIINVTLAAKMCSQVLCQE
 2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01 QGVCIRKDNSSDYHLNPDNFAIQLE-KGGKFTV
 2pe4.1.A HGRCVRRITSHPKALLLNPAFSIQLTGPGGPLSL

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

Model_01 DAVDVCIADGVCIDAF LKPPKETEESEQIFYNASPS
 2pe4.1.A -----

Model_01 TLSATMFIVSILFLIISVVSL
 2pe4.1.A -----

