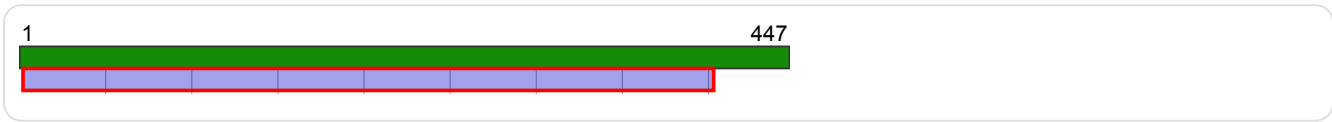


Untitled Project Created: today at 03:07

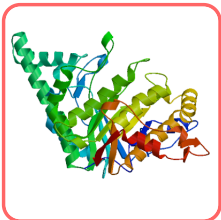
Summary Templates 45 Models 1 Project Data ▾

Model Results ⓘ

Order by: GMQE ▾



Model 01



Structure Assessment

Compare ☐

Download files ⬇

Display files 📄

Oligo-State
Monomer

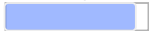
GMQE
0.69

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

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

Model_01 SLFSFETGSPRIAATGQGVTIFYVDRLLGYYPYIDSI
 2pe4.1.A SVFDVVANPGQTFRGPDMTIFYSSQLGTYPYY-TP

Model_01 TGVTVNGGIPQKISLQDHLDKAKKDITFYMPV-DN
 2pe4.1.A TGEIPVFGGLPQNASLIAHLARTEQDILAAIPAPDF

Model_01 LGMAVIDWEEWRPTWARNWKPKDVYKARSIELVQQ
 2pe4.1.A SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01 QNVQLSLTEATEKAKQEFEEKAGKDFLVETIKLGKL
 2pe4.1.A QHPDWPAPQVEAVAQDQFQGAARAWMAGTLQLGRA

Model_01 LRPNHLWGYYLFPDCYNHHYKKPGYNGSCFNVEIK
 2pe4.1.A LRPRGLWGFYGFPCYNDFLSPNYTGQCPSGIRA

Model_01 RNDPLSWLWAFSTALYPSIYLNTQQSPVAATLYV
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTGKSQMYV

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

Model_01 KFLSQDELVYTFGETVALGASGIVTWGTLSTIMRSM
 2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

Model_01 KSCLLLLKNYMETIILNPYIINVTLAAKMCSQVLCQE
 2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01 QGVCIRKNWNSSDYHLNPDNFAIQLE-KGGKFTV
 2pe4.1.A HGRCVVRTSHPKALLLLNPASFSQLTPGGGPLSL

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

Model_01 DAVDVCIADGVCIDAFLLKPPMETEEPQIFY
 2pe4.1.A -----

