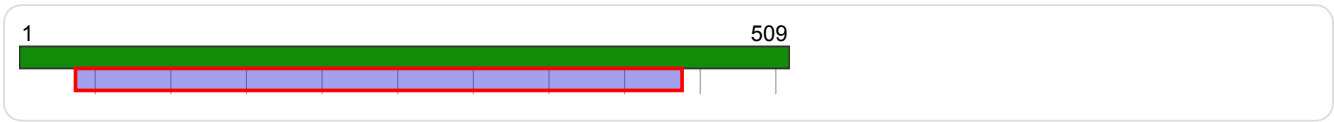


Untitled Project Created: today at 17:25

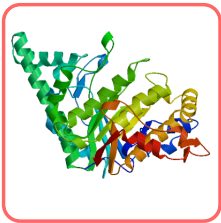
Summary Templates 31 Models 1 Project Data ▾

Model Results ⓘ

Order by: GMQE ▾



Model 01



Structure Assessment

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Oligo-State
Monomer

GMQE
0.62

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

Model_01 MGVLKFKHIFFRSFVKSSGVSQIVFTFLLIPCCLT
 2pe4.1.A -----

Model_01 LNFRAAPPVTPNYPFLWAWNPASEFCLGKFEDEPLDM
 2pe4.1.A --FRGP--LLPNRPFTTVWNPANTQWCLERHGVVDVDV

Model_01 SLFSEITGSPRINATGQGVTIFYVDRLGYYPIYDSI
 2pe4.1.A SVFDVVANPGQTFRGPDMTIFYSSQLGTYPYY--TP

Model_01 TGVTVNGGIPOKISLQDHLDAKAKDITEYMPV-DN
 2pe4.1.A TGEPPVFGGLPQNASLIAHLARTEQDILAAIPAPDF

Model_01 LGMAVIDWEEWRPTWARNWKPKDVYKNRSIELVQO
 2pe4.1.A SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01 QNVQLSLTEATEKAKQEEKAGKDFLVETIKLGKL
 2pe4.1.A QHPDWPAPQVEAVAQDQFQGAARAWMAGTLQLGRA

Model_01 LRPNHLWGYYLFPDCYNHHYKKPGYNGSCFNVEIK
 2pe4.1.A LRPRGLWGIFYGFPDCYNDFLSPNYTGQCPSGIRA

Model_01 RNDLSQLWNESTALYPSIYL-NTQOSPVAATLYV
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTKSOMYV

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

Model_01 KFLSQDELVYTFGETVALGASGIVINGTLSIMRSM
 2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

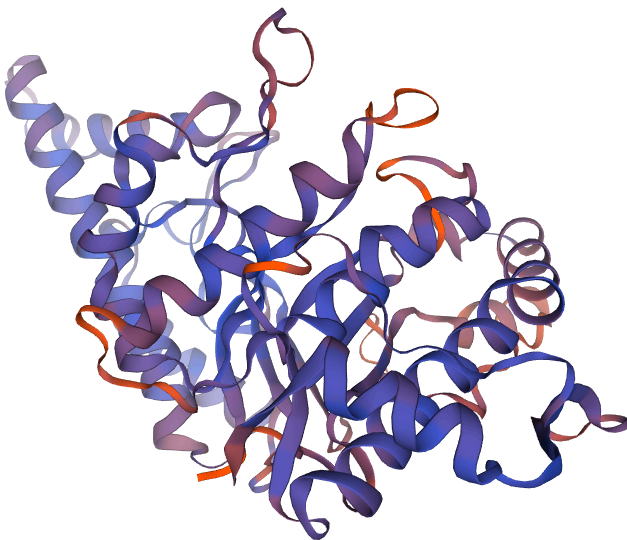
Model_01 KSCLLLDNYPMETILNPYIINVTLAAKMCSQVLCQE
 2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01 QGVCIRKNWNSSDYHLNPDNFAIQLE-KGGKFTV
 2pe4.1.A HGRCVVRTSHPKALLLNPAFSFIQLTPGGGPLSL

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

Model_01 DAVDVCIADGVCIDAFKPPMETEEPQIFYNASPS
 2pe4.1.A -----

Model_01 TLSATMFIVSILFLIISSVASL
 2pe4.1.A -----







Cartoon ▲

 ▲





