

Untitled Project

Created: today at 17:25

Summary

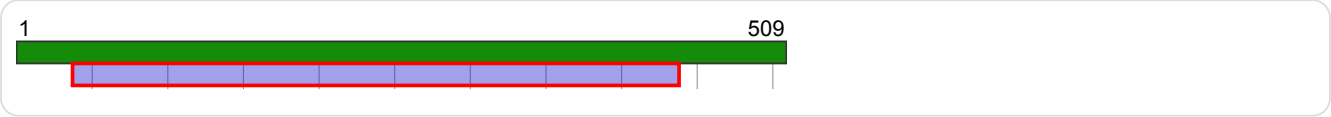
Templates 31

Models 1

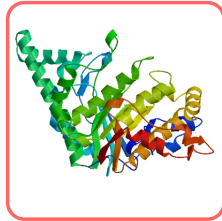
Project Data ▾

Model Results ⓘ

Order by: GMQE ▾



Model 01



Structure Assessment

Compare

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Display files 📄

Oligo-State
Monomer

GMQE
0.62

QMEANDisCo Global:
0.74 ± 0.05

QMEANDisCo Local

QMEAN Z-Scores

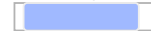
Template

2pe4.1.A Hyaluronidase-1

Seq Identity

42.82%

Coverage



Model-Template Alignment

Model_01 MGVLKFKHIFFRSFVKSSGVSQIVFTFLLI PCCLT
 2pe4.1.A -----

Model_01 LNFRA P P V I P N Y P F L W A W N A P S E F C L G K F D E P L D M
 2pe4.1.A --FRGP-LLPNRPFITTVWVANTQWCLERHGVVDVDV

Model_01 SLFSEITGSPRINATGQGVTIFYVDRDLGYYPYIDSI
 2pe4.1.A SVFDVVANPGQTRFRGPDMTIFYSSQLGTYPYY-TP

Model_01 TGVTVNGGIPQKISLQDHLDKAKKDITFYMPV-DN
 2pe4.1.A TGEPPVFGGLPQNASLIAHLARTEFDILAAIPAPDF

Model_01 LGMAVIDWEEWRPTWARNWKPKDVYKNRSIELVQO
 2pe4.1.A SGLAVIDWEAWRPRWAFNWDTKDIYRQRSRALVQA

Model_01 QNVQLSLTEATEKAKQEFKAGKDFLVETIKLGKL
 2pe4.1.A QHPDWPAPQVEAVAQDQFQGAARAWMAGTLQLGRA

Model_01 LRPNHLLWGYLLFPDCYNHHYKPGYNGSCFNVEIK
 2pe4.1.A LRPRGLWGFYGFPCYNYDFLSPNYTGQCPSGIRA

Model_01 RNDL SWLWNE STALYPSIYL-NTQOSPVAATLYV
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTGKSOMYV

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

Model_01 KFLSQDELVYTFGETVALGASGIVIWGTL SIMRSM
 2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

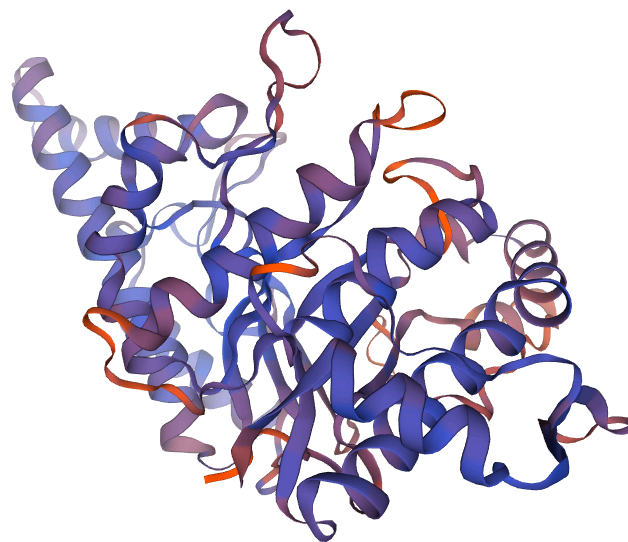
Model_01 KSCLLLDNYMETILNPYIINVTLAAKMCSQVLCQE
 2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01 QGVCIRKNWNSSDYHLHLPDNFAIQLE-KGGKFTV
 2pe4.1.A HGRCVVRTSHPKALLLNPA SF SIQLTPGGGPLSL

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

Model_01 DAVDVCIA DGVCIDAF LKPPMETEEPQIFYNASPS
 2pe4.1.A -----

Model_01 TLSATMFIVSILFLIISVASL
 2pe4.1.A -----



Interactive toolbar with icons for editing, settings, view mode (Cartoon), camera, play, zoom, and refresh.