

All Projects

Untitled Project Created: today at 02:11

Summary Templates 48 Models 1 Project Data

Model Results

Order by: GMQE

1447

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

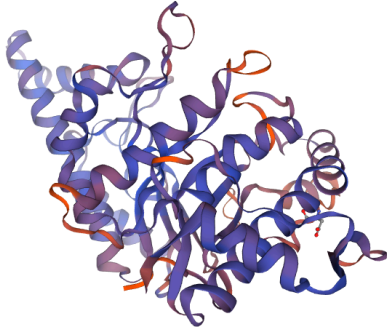
Template 2pe4.1.A Hyaluronidase-1

Seq Identity 43.07%

Coverage

Model-Template Alignment

Model_01 LNFRAPPVDPNPFLLWANNAPSEFCLGKPDDEPLDM
2pe4.1.A --FRGP-LIPNDPFFTIVNNANTQWCLERHGVDDVDV
Model_01 SLFSFEGSPRIAATGGQVTIPVDNLGYYPYDPSI
2pe4.1.A SVFDVYANPGQTFRGPDMITIPYSSQLGTYPIYV-TP
Model_01 TGVITVNGGITPQWISLQDHLDKAKKDITFYMPV-DN
2pe4.1.A TGEIPVFGGLPQNASLIAHLARTFQDILAAIPAPDF
Model_01 LGMAVTDWEWRPTWABNNWPKDVKYKARSTELVQQ
2pe4.1.A SGLAVTDWEAWRPRWAEINWOTKDIYRQSRALVQA
Model_01 QNVQLSLTEATEKAKQEFKAGKDFLVETIKLQKL
2pe4.1.A QHPDWPAAPQVEAVAQDQFGGAARAWMAGTLQLQRA
Model_01 LRPNHLWGYLLPPDCYNHHYKKPGYNGSCFNVDIK
2pe4.1.A LRPRLWGFYGFDPDCYNVDFLSPNYTGQCPSGTRA
Model_01 RNDPLSHLWNESTALYPSIYL-NIQQSPVAAILYV
2pe4.1.A QNDQLGWLWGSRALYPSIYMPAVLEGTGKSGHYV
Model_01 RNRVREAIRVSKIPDAKSPLPVFAYTRIVTDQVL
2pe4.1.A QHRVAEFRVA-VVAGDPNLPVLPVQI-FYDTTN
Model_01 KFLSDDELVYTFGETVALGASGVITVGTLSNMRSM
2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLVSWENTRTK
Model_01 KSCLLLDNYMETILNPYIINVTLAAKMCSQVLCQE
2pe4.1.A ESCQATKEYMDTTLGPFILNVTSGALLCSQALCSG
Model_01 QGVCTIKKNWSSDYHLNPDNFAIQIE-KGKKTV
2pe4.1.A HGRKVRKRTSHPKALLLNPAFSTIQLTPGGGPLYSL
Model_01 RLKPKTEEDLEQFSEKPYLSLYSTLSCKEKADV KDT
2pe4.1.A RGALSTEDQAQMAVEFKRCY-----
Model_01 DAVDVCIAIDGVCIADFLKPPMETEEPQIFY
2pe4.1.A -----



Cartoon

https://swissmodel.expasy.org/interactive/ykcvPM/models/

Halozyme EX2194
Merck v. Halozyme
PGR2025-00050

1/2

