

 All Projects

Untitled Project

Created: today at 00:35

Summary

Templates

48

Models

1

Project Data ▾

Model Results ⓘ

Order by: GMQE



Model 01



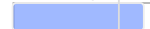
Structure Assessment

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 SLFSFETGSPRIAATGQGVITFYVDRLGYYPYIDSI
 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 LRPRGLWGFYGFPCYNYDFLSPNYTGQCPSGIRA

Model_01 RNDPLSWLWNESTALYPSIYLNTIQSPVAATLYV
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTKSQMYV

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

Model_01 KFLSQDELVYTFGETVALGASGIVTWGTLSTMRSM
 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 DAVDVCIADGVCIDAF LKPPMETEEPQIFY
 2pe4.1.A -----