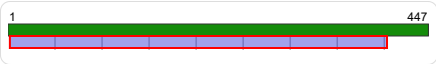


All Projects

Untitled Project Created: today at 18:24

Summary Templates 40 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 42.82%

Coverage

Model-Template Alignment

Model_01 LNFRAPPVDPNPFLWANNAPSEFCLGKPDDEPLDM

2pe4.1.A --FRGP-LIPNDPFTTYVWANTQWCLERHGVDDVDV

Model_01 SLFSFEGSPRINATGGQVITPVVDLGYYPYDSDI

2pe4.1.A SVFDVYANPGQTFRGPDMITPVSSQLGTYPVY-TP

Model_01 TGVLYNGGITPQWISLQDHLDAKKKDITFVMPV-DN

2pe4.1.A TGEIPVFGGLEQNASLTAHLARTEQDILAAIPAPDF

Model_01 LGMAVZDWEWRPTWARNWPKDVKYKNRSIELVQQ

2pe4.1.A SGLAVZDWEAWRPWAPNWDTKDIYRQRSRALVQA

Model_01 QNVQLSLEATEKAKOEFKAGKDFLVETIKLQKL

2pe4.1.A QHPDWPAQVEAVAQDQFGGAARWMAAGTLQLGRA

Model_01 LRPNHLWGYLLFPDCYNHHYKPGYNGSCFNVYIK

2pe4.1.A LRPRLWGFYGFDPDCYNVDFLSPNYTGQCPSGPRA

Model_01 RNDLSLWLNSESTALYPSIYLNIQQSPVATLYV

2pe4.1.A QNDQLGWLWGSRALYPSIYMPVLEGTGKSQMYV

Model_01 RNRVREAIRVSKLRDAKSPLPVFAYRIVETDQVL

2pe4.1.A QHRVAEAFRVA-VAAGDPNLPVLPYQI-KYDTTN

Model_01 KFLSQDELVYTFGETVALGASGLVINGTESQMRSM

2pe4.1.A HFLPLDELEHSLGESAXGAAGVVLWVSWENTRTK

Model_01 KSCLEEDNYMETILNPYIINVTLAAKMSQVLCQE

2pe4.1.A ESCQATKEYMDTTLGPFILNVTSGALLCSQALCSG

Model_01 QGVCIKNWNSSDYLLHNPDMFAIQLE-KGKKTV

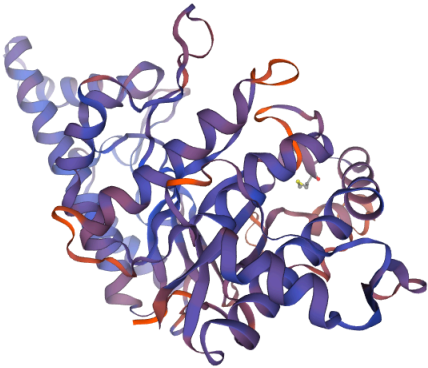
2pe4.1.A HGRGCVRRTSHPKALLLNPAFSTIQLTPGGGPLSL

Model_01 R6KPTCEDLEQFSEKFCSCYSTLSCKEKADV KDT

2pe4.1.A RGALSLEDQAQMAVEFKRCY-----

Model_01 DAVDVCIAIDGVCIDAFLLKPMMEETEPQIFY

2pe4.1.A -----



Cartoon

