

# Untitled Project

Created: today at 17:25

Summary

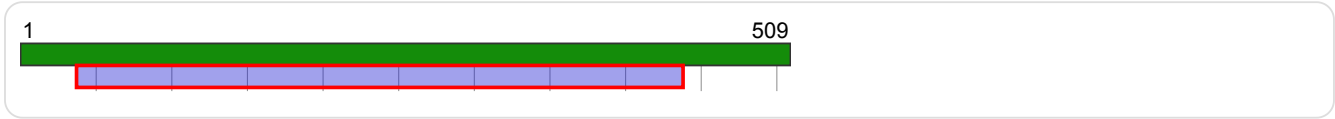
Templates 31

Models 1

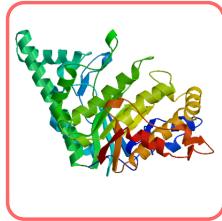
Project Data ▾

## Model Results ⓘ

Order by: GMQE ▾



### Model 01



Structure Assessment

Compare

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**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 LNFRAAPPVTPNYPFLWAWNPAPSEFCLGKFEDEPLDM  
 2pe4.1.A --FRGP-LLPNRPFTTVWNPANTQWCLERHGVVDVDV

Model\_01 SLFSEITGSPRINATGQGVITIFYVDRDLGYYPYIDSI  
 2pe4.1.A SVFDVVANPGQTRFRGPDMTIFYSSQLGTYPIY-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 LRPNHLWGYLLFPDCYNHHYKPGYNGSCFNVEIK  
 2pe4.1.A LRPRGLWGFYGFPCYNYDFLSPNYTGQCPSGIRA

Model\_01 RNDL SWLWNESTALYPSIYL-NTQOSPVAATLYV  
 2pe4.1.A QNDQLGWLWGQSRALYPSIYMPAVLEGTGKSOMYV

Model\_01 RNRVREAIRVSKIPDAKSPLPVFAYTRIVFTDQVL  
 2pe4.1.A QHRVAEAFRVA-VAAGDPNLPVLPYVQI-FYDTTN

Model\_01 KFLSQDELVYTFGETVALGASGIVIWGTL SIMRSM  
 2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLWVSWENTRTK

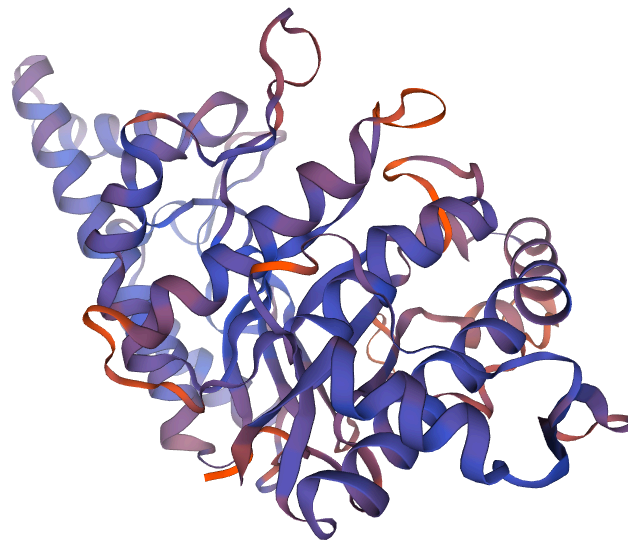
Model\_01 KSCLLLDNYMETILNPYIINVTLAAKMCSQVLCQE  
 2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG

Model\_01 QGVCIRKNWNSSDYHLHLPDNFAIQLE-KGGKFTV  
 2pe4.1.A HGRCVRRTSHPKALLLNPAFSIQLTPGGGPLSL

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.