

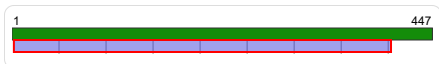
All Projects

Untitled Project Created: today at 17:18

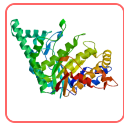
Summary Templates 50 Models 1 Project Data

Model Results

Order by: GMQE



Model 01



Structure Assessment

Compare

Download files

Display files

Oligo-State	GMQE	QMEANDisCo Global:
Monomer	0.69	0.74 ± 0.05

QMEANDisCo Local

QMEAN Z-Scores

Template

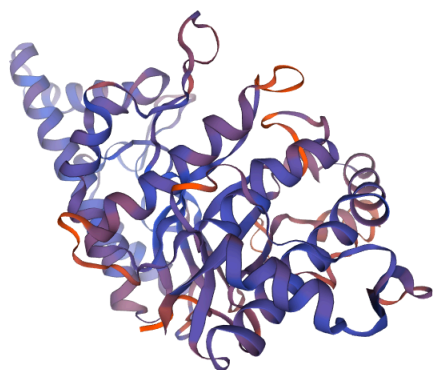
2pe4.1.A Hyaluronidase-1	Seq Identity
	42.82%
	Coverage

Model-Template Alignment

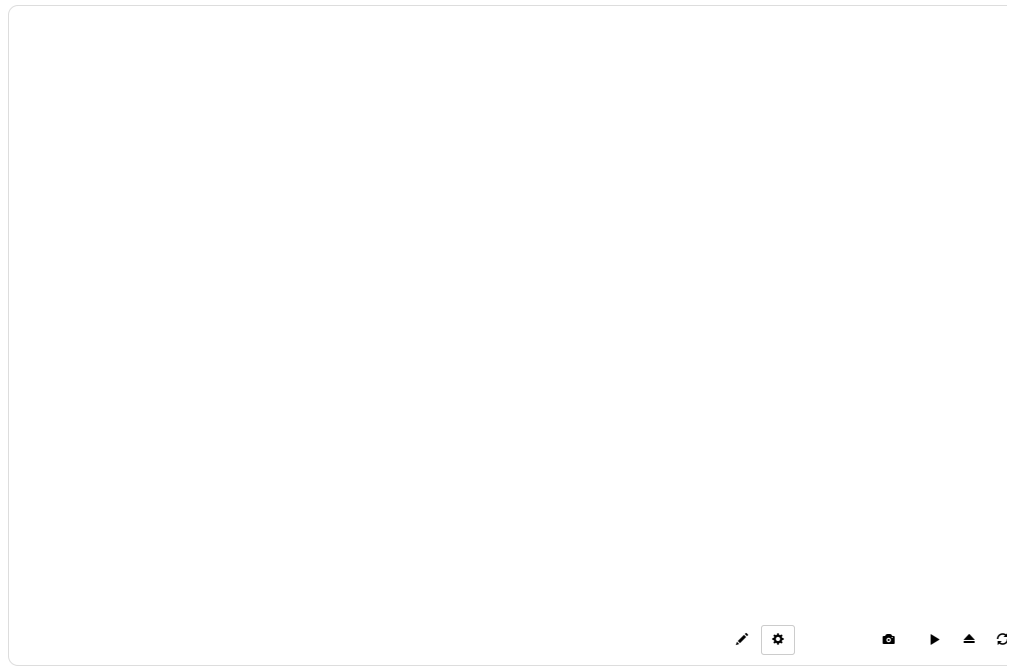
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Model_01 LNFRAPPVDPNPFLLWANNAPSEFCGLGKFDDEPLDM
2pe4.1.A --FRGP-LIPNDPFFITVWVANNIQWCLERHGVDDVDV
Model_01 SLFSEKGSPRINATGQQVITPVVDBLGYPPYDSDI
2pe4.1.A SVFDVYANPGQTFRGPDMITFVSSQLGTYPYV-TP
Model_01 TGVIVNGGIPQWISLQDHLDKAKKDITFVMPV-DN
2pe4.1.A TGEIPVFGGLEQNASLTAHLARTEQDILAAIPAPDF
Model_01 LGMAVZDWEWRPTWARNNWPKDYVKNRSTELVQQ
2pe4.1.A SGLAVZDWEAWRPRWAPNWDTKDIYRQRSRALVQA
Model_01 QNVQLSLEATEKAKQEFKAGKDFLLETIKLQKL
2pe4.1.A QHPDWPARQVEAVAQQDQGAARAWMAGTLQLQRA
Model_01 RPNHLWGYLPPDCYNHHYKPGYNGSCFNVEIK
2pe4.1.A ERPRGLWGFYGFPCYNYDFLSPNYTGQCPGZRA
Model_01 RNDLSLWLNNESTALYPSIYLNIQQSPVAATLYV
2pe4.1.A QNDLGLWLGQSRALYPSIYMPVLEGTGKSOYV
Model_01 RNRVREAIRVSKLRDAKSPLPWFAYRIVETQVL
2pe4.1.A QHRVAEFRVA-VAAGDPNLPVLPYQI-KYDTTN
Model_01 KFLSQDELVYTFGETVALGASGVILKGTESIMRSM
2pe4.1.A HFLPLDELEHSLGESAAQGAAGVVLHVSWENTRTK
Model_01 KSCLELDNYMHTILNPYIINVTLAAKMESQVLCQE
2pe4.1.A ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG
Model_01 QGVCIKNWNSDYLHLNPDNFAIQLEKGGKKTIV
2pe4.1.A HGRRCVRRISHPKALLLNPAFSIQLTPGGGLSL
Model_01 RQKPTLEDLEQFSEKFCSCYSLTSCKEKADVKDT
2pe4.1.A RGALSLEDQAQMAVEFKRCY-----
Model_01 DAVDVCIDGVCIIDAFLKPPMETEEPQIFY
2pe4.1.A -----

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Halozyme EX2184
Merck v. Halozyme
PGR2025-00017



Halozyme EX2184
Merck v. Halozyme
PGR2025-00017