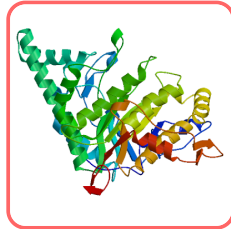


Templates Models

Order by: GMQE



Model 01



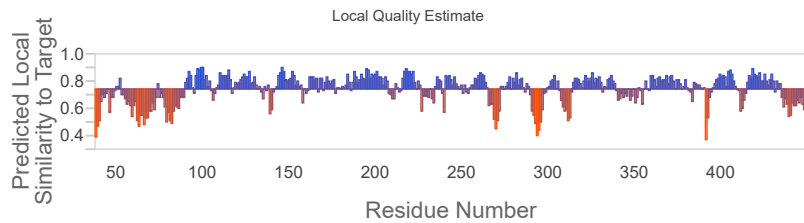
Display files

Oligo-State Monomer

GMQE 0.64

QMEANDisCo Global: 0.74 ± 0.05

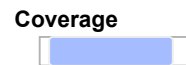
QMEANDisCo Local



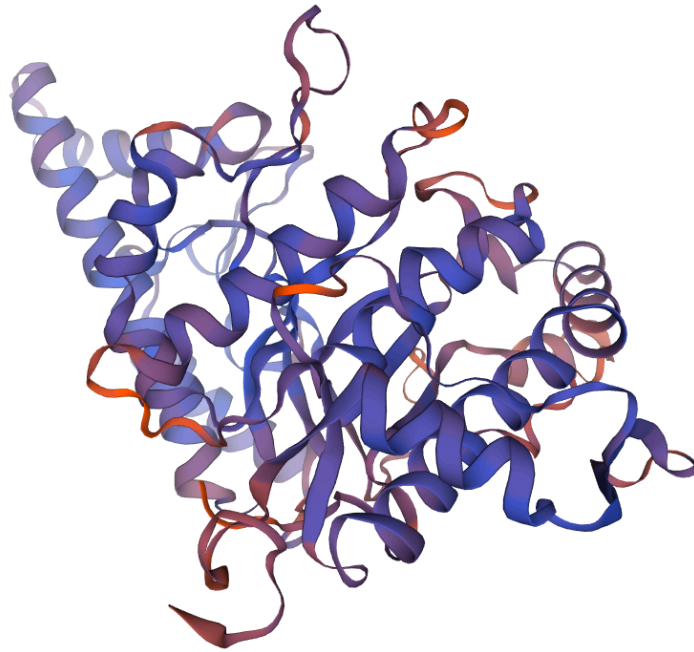
QMEAN Z-Scores

Template template_upload.1.A Polypeptide

Seq Identity 40.24%

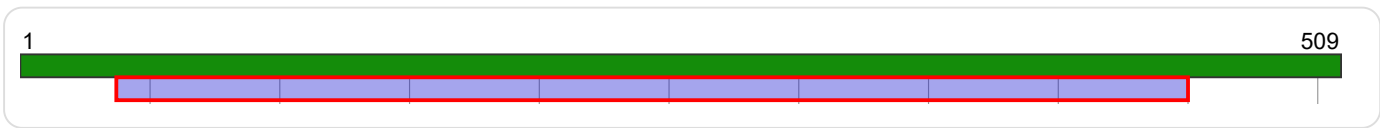


Model-Template Alignment

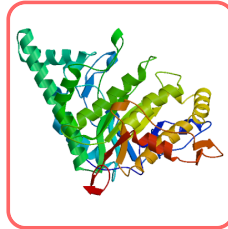


Templates Models

Order by: GMQE



Model 01



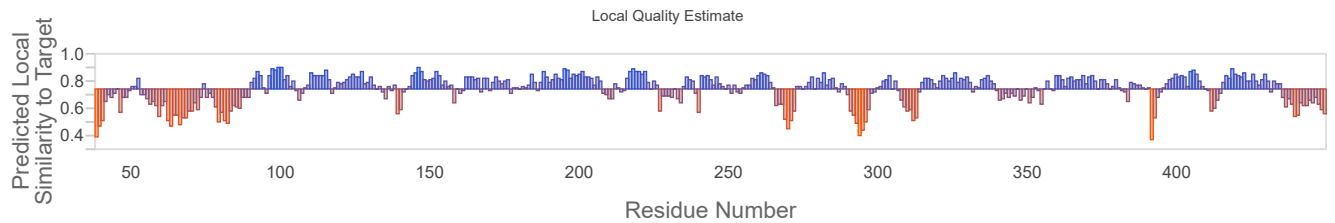
Display files

Oligo-State Monomer

GMQE 0.64

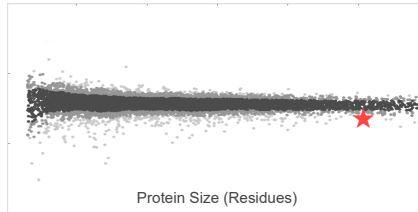
QMEANDisCo Global: 0.74 ± 0.05

QMEANDisCo Local



QMEAN Z-Scores

QMEAN	-2.65
Cβ	-0.41
All Atom	-0.84
solvation	0.41
torsion	-2.84



Template

template_upload.1.A Polypeptide

Seq Identity 40.24%



Biounit Oligo State Monomer QSQE 0.00

Method Unknown
Seq Similarity 0.40
Coverage 0.81
Range 38-450

Model-Template Alignment

Model_01	MGVLKFKHIFFRSFKSSGVSQIVFTFLIPCCLTNFRAPPVTPNYPFLWAWNA	55
template_upload.1.A	-----SFRGPLLPNRPFTTVWNA	19
Model_01	PSEFCLGKFDDEPLDMSLFSFETGSPRINATGQGVTFYVDR LGYYPYIDSITGVTV	110
template_upload.1.A	NTQWCLERHGVDVDVSVFDVYANPGQTFRGPDMTIFYSSQLGTYPYTPT-GEPV	73
Model_01	NGGIPOKISLQDHLDAKAKDITFVMPV-DNLGMAVTDWEWRPTWARNWPKPKDVY	164
template_upload.1.A	FGGLPQNASLIAHLARTEQDILAAIPAPDFSGLAVIDWEAWRPRWAENWDTKDIY	128
Model_01	KNRSIELVQQONVQLSLTEATEKAKQEFKAGKDFLVEITIKLGKLLRPNHLWGYV	219
template_upload.1.A	RQRSRALVQAQHPDWPAQVEAVAQDQEQGAARAWMAGTLQLGRALRPRGLWGFV	183
Model_01	LFPDCYNHHYKPKGYNGSCFNVEIKRNDLSWLWNESTALYPSIYLNTOQS-PVA	273
template_upload.1.A	GFPDCYNYDFLSPNYTGQCPSGIRAQNDQLGWLWQSRALYPSIYMPAVLEGTGK	238
Model_01	ATLYVRNRVREAIRVSKIPDAKSPLPVFAYTRIVFTDQVLKFLSQDELVYTFGET	328
template_upload.1.A	SQMYVQHRVAEAFRVAVAA-GDPNLPVLPYVQIFYD TT-NHFLPLDELEHSLGES	291
Model_01	VALGASGIVTNGTLSIMRSMKSCMLLDNYMETILNPYIINVTLAAKMCSQVLCQE	383
template_upload.1.A	AAQGAAGVVLWVSWENTRTKESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSG	346
Model_01	QGVCTIRKNWNSDYHLNPDNFATOLEK-GGKETVRGKPTLEDLEQFSEKFCYCS	437
template_upload.1.A	HGRCVRRRTSHPKALLLNPAFSIQLTGGGGLSLRGALSLEDQAQMAVEFKCRC	401
Model_01	YSTLSCKEKADVKDDTDAVDVCIADGVCIDAFKPPMETEEPQIFYNASPSTLSAT	492
template_upload.1.A	YPGWQAPWCERKS-----	414
Model_01	MFIVSILFLIISSVASL	509
template_upload.1.A	-----	

