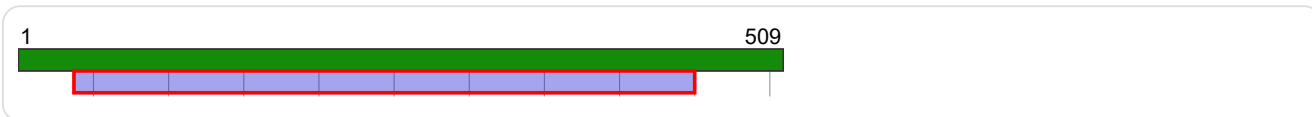


I352

Created: Dec. 4, 2024, 9:44 p.m. at 21:44

Templates Models

Order by: GMQE ▼



Model 01



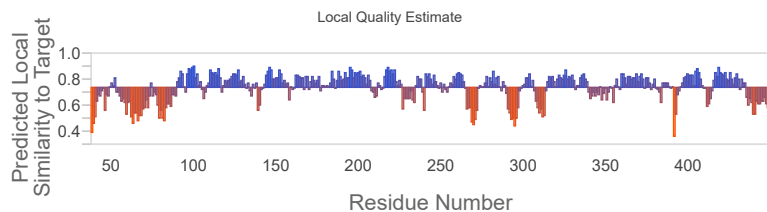
Display files

Oligo-State ⓘ
Monomer

GMQE ⓘ
0.63

QMEANDisCo Global: ⓘ
0.74 ± 0.05

QMEANDisCo Local ⓘ



QMEAN Z-Scores ⓘ

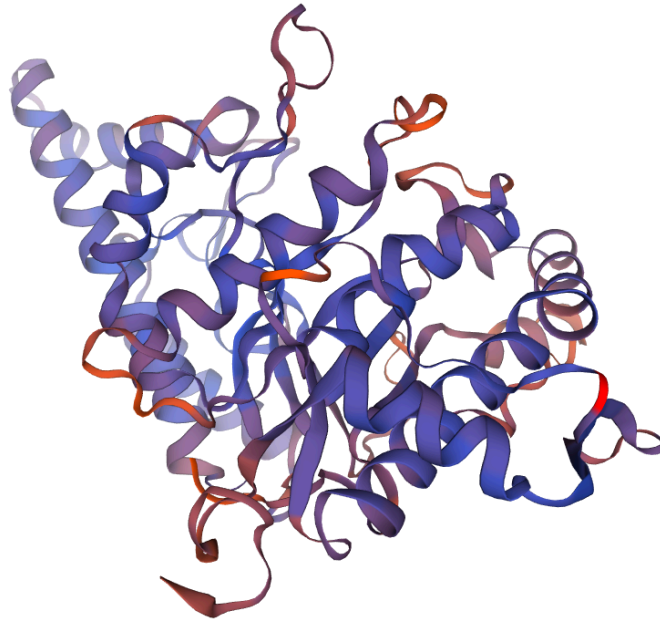
Template
template_upload.1.A Polypeptide

Seq Identity
40.24%



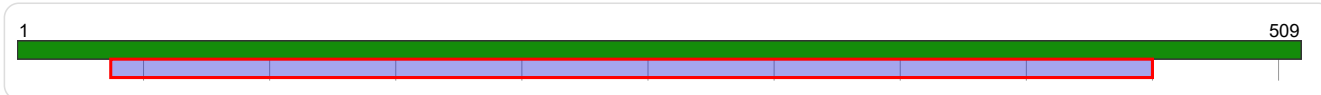
Model-Template Alignment ⓘ

ASP 403 A
Confidence: 0.83

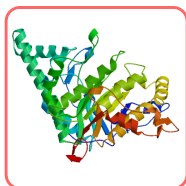


Templates Models

Order by: GMQE



Model 01



Display files

Oligo-State Monomer

GMQE 0.63

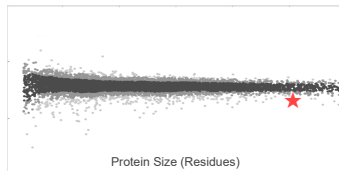
QMEANDisCo Global: 0.74 ± 0.05

QMEANDisCo Local



QMEAN Z-Scores

QMEAN	-3.10
Cβ	-0.07
All Atom	-0.76
solvation	0.41
torsion	-3.39



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	MGVLKFKHIFFRSFKSSGVSQIVFTFLILPCCLTLNFRAPPVDPNVPFLWAWNAPSEFCLGKFDEPLDM	70
template_upload.1.A	-----SFRGPLLPNPPFTTVWNPANTQWCLERHGVDVDDV	34
Model_01	SLFSFETGSPRINATGQGVITFYVDRILGYYPYIDSITGVITVNGGIPOKISLQDHLDKAKKDITFYMPV-DN	139
template_upload.1.A	SVFDVYVANPGQTFRGPDMITFYSSQLGTYPYVTPT-GEPIYFGGLPONASLIAHLARTFQDILAAIPAPDF	103
Model_01	LGMAVTDWEWRPTWARNNKPKDYYKNRSIELVQQQNVQLSLTEATEKAKQEFFEKAGKDFLVETIKLGKL	209
template_upload.1.A	SGLAVTDWEAWRPRWAFENWDTKDIYRQRSRALVQAQHPDWPAPQVEAVAQDQEQGAARAWMAGTLQLGRA	173
Model_01	LRPNHLWGYLFPDCYNHYYKKPGYNGSCFNVEIKRNDLSWLWNESTALYPSIYLNNTQQS-PVAATLYV	278
template_upload.1.A	LRPRGLWGFYGFPCYNYDFLSPNYTGQCPSSGIRAQNDQLGWLWQSRALYPSIYMPAVLEGTGKSQMYV	243
Model_01	RNRVREAIRVSKIPDAKSPLPVFAVTRIVFIDQVLKFLSQDELVYTFGETVALGASGIVIVGTLSIMRSM	348
template_upload.1.A	QHRVAEAFRVAVAA-GDPNLPWLPYVQIFVDTT-NHFLPLDELEHSLGESAAQGAAGVVLVSWENRTRK	311
Model_01	KSCILLDNYMETILNPYIINVTLAAKMCSQVLCEQEGVCFIRKNWNSSDYHLHNPDNFAIQLEK-GGKFTV	417
template_upload.1.A	ESCQAIKEYMDTTLGPFILNVTSGALLCSQALCSGHGREVRRTSHPKALLLNPAFSFIQLTPGGGLSL	381
Model_01	RGKPTLEDLEQFSEKFYCSYSTSCCKEKADYKDDTDAVDVCIADGVCIDAFLLKPPMETEETEPQIFYNASPS	487
template_upload.1.A	RGALSLEDQAQMAVEFKRCYPGWQAPWCERKS-----	414
Model_01	TLSATMFIVSILFLIISSVASL	509
template_upload.1.A	-----	

CYS 437 A
Confidence: 0.60

