

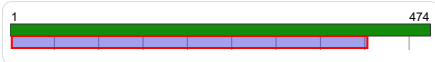
All Projects

Untitled Project Created: today at 17:58

Summary Templates 34 Models 1 Project Data

Model Results

Order by: GMQE



Model 01

Structure Assessment

Compare

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Oligo-State
 Monomer

GMQE
 0.66

QMEANDisCo Global:
 0.73 ± 0.05

QMEANDisCo Local

QMEAN Z-Scores

Template

2pe4.1.A Hyaluronidase-1

Seq Identity: 43.32%
Coverage:

Model-Template Alignment

```

Model1_01 LNFRAAPPVDPNPF1FLWANNAP2SEFC3LGK4FD5EPLDM
2pe4.1.A --FRGP-L1PN2PPFT3IVN4ANT5QWCLERH6GV7DVDV
Model1_01 SLFSE1SGSPRINVTGQD2VIT3YV4DBLGY5PP6DSI
2pe4.1.A SVFDV1YANPG2QTFRGPDM3IT4YSS5QLGTY6Y7TP
Model1_01 TGV1YNGG2IQ3WIS4LQD5HL6DKAK7KDIT8F9MPV-DN
2pe4.1.A TGE1YV2FGGLE3QNAS4L5LAHL6ARTE7QD8IL9AA10IPAPDF
Model1_01 LGM1AV2PDWE3EWRPT4W5ARN6W7KPK8DVY9KNR10ST11EL12V13Q14Q
2pe4.1.A SGL1AV2PDWE3AWR4PR5W6AP7NW8TK9DI10YR11QR12S13R14LV15QA
Model1_01 QNV1QL2ST3TE4ATE5KAK6O7E8FE9KAG10KD11FL12VET13IK14L15K16L
2pe4.1.A QHP1DW2PAP3QVE4AVA5QD6Q7GA8ARA9WM10AG11T12L13Q14L15Q16RA
Model1_01 RPN1HL2WGY3LFP4DCYN5H6YK7KPG8YNS9CF10N11Y12E13IK
2pe4.1.A ER1PR2GL3WGY4GF5PD6CYN7YD8FL9SP10NY11T12G13CP14SG15Z16RA
Model1_01 RND1L2SWL3NE4ST5AL6PS7IY8L9NI10QSP11V12AT13LY14V
2pe4.1.A QND1LG2WL3WGS4R5AL6Y7PS8IY9MP10AV11LE12GT13G14S15Q16MY17V
Model1_01 RNR1VQ2E3IR4V5SK6LR7DA8K9S10PL11P12V13Y14R15IV16ET17Q18VL
2pe4.1.A QHR1V2A3E4FR5V6A7AG8DP9N10L11P12Y13Y14Q15I16-17Y18DT19TN
Model1_01 K1FL2S3Q4DEL5V6Y7TF8GET9V10LG11SG12V13IK14TS15IM16RS17M
2pe4.1.A H1FL2PL3D4LE5H6SL7G8ESA9X10G11A12GV13L14W15SV16EN17TR18TK
Model1_01 K1SC2QL3ED4NY5MT6IL7NP8Y9I10NV11TL12A13AK14MS15Q16VL17C18Q19E
2pe4.1.A E1SC2Q3AI4K5EY6MD7T8LG9FF10IL11NV12TS13G14ALL15C16S17Q18AL19CS20G
Model1_01 Q1GV2E3IK4KN5W6SS7D8YL9HL10NP11DN12FA13IQ14LE15-16K17G18K19KT20V
2pe4.1.A H1GR2CV3R4RT5SH6PK7ALL8LN9P10ASF11SI12QL13TP14GG15PL16SL
Model1_01 R1G2K3PT4LED5LE6Q7F8SE9NY10CS11Y12TL13SC14KE15K16ADV17K18DT
2pe4.1.A R1G2AL3S4LED5QA6MA7VE8FK9RC10Y-----
Model1_01 D1AV2D3V4CI5AD6G7VC8ID9AF10L11K12PP13ME14TE15ES16Q17IF18YN19AS20PS
2pe4.1.A -----
Model1_01 T1LS2AT3M4F5IV6S7IL8FL9II10SS11V12AS13L
2pe4.1.A -----
    
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