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Untitled Project Created: today at 19:50

Summary

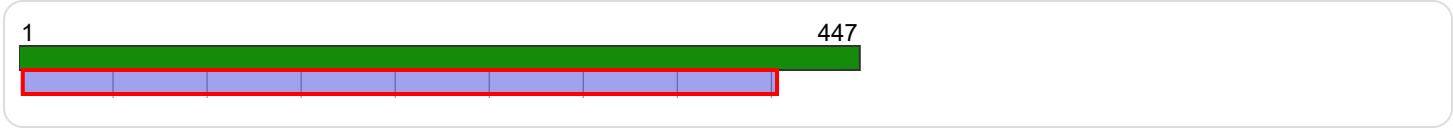
Templates 48

Models 1

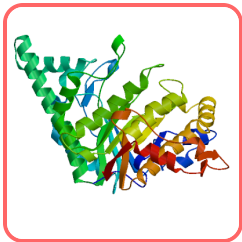
Project Data ▼

Model Results ⓘ

Order by: GMQE ▼



Model 01



Structure Assessment

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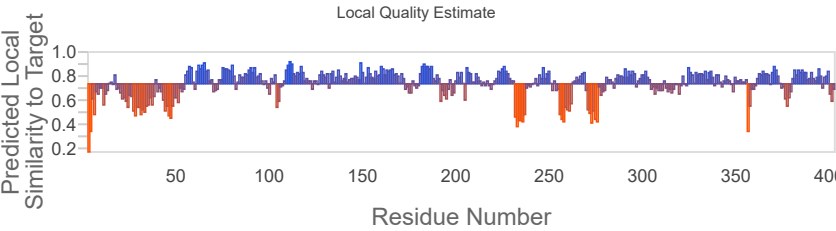
Display files 📄

Oligo-State ⓘ
Monomer

GMQE ⓘ
0.69

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

Halozyne EX2197
Merck v. Halozyne
PGR2025-00046

Model_01 L N F R A P P V I P N V P F L W A W N A P S E F C L G K F D E P L D M
 2pe4.1.A - - F R G P - L L P N R P F T T V W N A N T Q W C L E R H G V D V D V

Model_01 S L F S E I G S P R I N A T G Q G V T I F Y V D R L G Y Y P Y I D S I
 2pe4.1.A S V F D V V A N P G Q T F R G P D M T I F Y S S Q L G T Y P Y Y - T P

Model_01 T G V T V N G G I P Q K I S L Q D H L D K A K K D I T E Y M P V - D N
 2pe4.1.A T G E P V F G G L P Q N A S L T A H L A R T F Q D I L A A I P A P D F

Model_01 L G M A V I D W E E W R P T W A R N W K P K D V Y K N R S I E L V Q Q
 2pe4.1.A S G L A V I D W E A W R P R W A F N W D T K D I Y R Q R S R A L V Q A

Model_01 Q N V Q L S L T E A T E K A K Q E F E K A G K D F L V E T I K L G K L
 2pe4.1.A Q H P D W P A P Q V E A V A Q D Q F Q G A A R A W M A G T L Q L G R A

Model_01 L R P N H L W G Y Y L F P D C Y N H H Y K K P G Y N G S C F N V E I K
 2pe4.1.A L R P R G L W G F Y G F P D C Y N Y D F L S P N Y T G Q C P S G I R A

Model_01 R N D D L S W L W N E S T A L Y P S I Y L N T Q Q S P V A A T I Y V
 2pe4.1.A Q N D Q L G W L W G S R A L Y P S I Y M P A V L E G T G K S O M Y V

Model_01 R N R V R E A I R V S K I P D A K S P L P V F A Y T R I V E T D Q V L
 2pe4.1.A Q H R V A E A F R V A - V A A G D P N L P V L P Y V Q I - F Y D T T N

Model_01 K F L S Q D E L V Y T F G E T V A L G A S G I V I N G T L S I M R S T
 2pe4.1.A H F L P L D E L E H S L G E S A A Q G A A G V V L W V S W E N T R T K

Model_01 K S C L L L D N Y M E T I L N P Y I I N V T L A A K M C S Q V L C Q E
 2pe4.1.A E S C Q A I K E Y M D T T L G P F I L N V T S G A L L C S Q A L C S G

Model_01 Q G V C I R K N W N S S D Y L H L N P D N F A I Q L E - K G G K F T V
 2pe4.1.A H G R C V R R T S H P K A L L L N P A S F S I Q L T P G G G P L S L

Model_01 R G K P T L E D L E Q F S E K F Y C S C Y S T L S C K E K A D V K D T
 2pe4.1.A R G A L S L E D Q A Q M A V E F K C R C Y - - - - -

Model_01 D A V D V C I A D G V C I D A F L K P P M E T E E P Q I F Y
 2pe4.1.A - - - - -







Cartoon ▲

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