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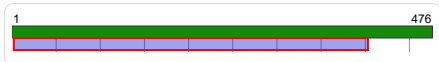
Untitled Project

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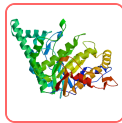
Summary Templates 40 Models 1 Project Data

Model Results

Order by: GMQE



Model 01



Structure Assessment

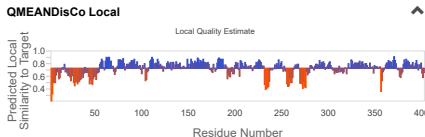
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Oligo-State Monomer

GMQE 0.65

QMEANDisCo Global: 0.73 ± 0.05



QMEAN Z-Scores

Template

Template	Seq Identity
2pe4.1.A Hyaluronidase-1	43.83% Coverage

Model-Template Alignment

```

Model1_01 L NFRAPPIIPNMPFLWANNAPSEFCGLKFDDEPLDM
2pe4.1.A  - -FRGP-LIDPNDPFTIVVWANTQWCLERHGVVDVDV
Model1_01 S L F S L L G S P R I N V T G Q A V I T P V D B L G Y P P D S I
2pe4.1.A  S V F D V A N P G T F R G P D M I T P Y S S L G T Y P V Y - T P
Model1_01 T G V I V N G G I P Q W I S L Q D H L D K A K K D I L F Y M P V - D N
2pe4.1.A  T G E P V F G G L E Q N A S L T A H L A R T E Q D I L A A I P A P D F
Model1_01 L G M A V P D W E E W R P T W A R N W K P K D V Y K N R S T E L V Q Q
2pe4.1.A  S G L A V P D W E A W R P R W A P N W T K D I Y R Q R S R A L V Q A
Model1_01 Q N V Q L N L T E A T E K A K O F E K A G K D F M V E T I K L Q K L
2pe4.1.A  Q H P D W P A P Q V E A V A Q D Q F G G A A R A W M A G T L Q L Q R A
Model1_01 R P N H L W G 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  E 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 P R A
Model1_01 R N D L S W L N E S T A L Y P S I Y L L N I Q S P V A T L Y V
2pe4.1.A  Q N D L G W L W G S R A L Y P S I Y M P V L E G T G K S Q M Y V
Model1_01 R R R V R E A I R V S K L R D A K S P L P F V Y X R I V E T Q V L
2pe4.1.A  Q H R V A E A F R V A - V A G D P N L P Y L P Y Q I - F Y D T T N
Model1_01 K F L S Q D E L V Y T F G E T V A L G S G I V I N G S S I M R S M
2pe4.1.A  H F L P L D E L E H S L G E S A X G A A G V L W V S W E N T R T K
Model1_01 K S C Q L E D N Y M E T I L N P Y I I N V T L A A K M 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
Model1_01 Q G V E I K D W N S S D Y L H L N P D N F A I Q L E - K G G K 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
Model1_01 R G K P T C E D L E Q F S E N F Y C S 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 M A V E F K R C Y - - - - - - - - - - - - - - - -
Model1_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 S Q I F Y N A S P S
2pe4.1.A  - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -
Model1_01 T L S A T M F I W R L E V W D Q G I S R M G F F
2pe4.1.A  - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -

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Halozyyme EX2190
 Merck v. Halozyyme
 PGR2025-00030

