

T F K E M P T K E G S Y L S S D L F K G V G K K T A Q N I V N T L G D S A I N K D D H S V L Y K V S R S K K K Q K T L A G Q L S R N Q G S E Q I M I S L N Q
H F K E M P T K E A Q L S S D L F K G V G K K T A E E I V K K L G D S A I N K D A A A A E D P G L S K K K A D Q T A E A T Q A H E L E K I M I R L H D

T F E K E M P Q T K E A I I S Y L S S D L F K G V G K K T A Q N I V N T L G D N A I N D I L D D H S V L E K V S G L S K K K Q K Q I A E Q I S A N Q E S E K I M I R L H D
H F K K E I P T T K E G I I Q Y L S S D L F E G I G K K T A E E I V K K L G D S A I N K I L A D A S V L Y D V P R L S K K K A D T L A G A L Q R H Q G L E Q I M I S L N Q

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F G F G P Q L S M K A Y Q F V L E D T L T K L Q R N P V Q L Y V D Y K G I F N K A D Q L G S R M G L S Y N H P E R V K A A L Y T L E T T C L S Q G H T Y L P T N W Y I E E Q L
L G F G P K L S M A I Y Q F Y L G D T L T I I D R N P Y Q L T Y D I K G I G F N K A D Q L A R N I G I T A Y N D N E R L K A A L Y T L E E E C I K Q G H T Y L P I N V Y I
F G F G P Q L S M K I Y Q A Y E S E T L E K I Q E N P Y Q L V K D V E G I G F G K A D E L G S R M L G S N H P E R V K A A L Y T L E T T C L S E G H T Y I E T E Q L

[illegible][illegible]

RLTEVYRQQEGSSVELAHQMKNGQLPNTLQRTKDRSFRCQGSQIPTWEKWTSAVYNKGYTMDIQVLAPMYRGKAGINRNLN
 NLTEVYRQQDGSSIIELAHRMKLGKQKILDITQRFHDRSFINCQANQIPTVVEKVVTSAVYNKGYTMDIQVLAPMYRGKAGIKRNLN
 RLTDITYRQAEGSSVELAHQMKNGLLPNNLTAPTDRSFIRCGSQIKEVVEKVVTANALKKGYTAKDIQVLAPMYRGKAGINELN

586
595

logo

NTDB id 266553 RK88 RS03090 WP 001283316.1
NTDB id 355 BSU 27480 NP 390625.1
consensus



671
680

logo

NTDB id 266553 RK88 RS03090 WP 001283316.1
NTDB id 355 BSU 27480 NP 390625.1
consensus



756
762

logo

NTDB id 266553 RK88 RS03090 WP 001283316.1
NTDB id 355 BSU 27480 NP 390625.1
consensus



825
798

X non conserved
X similar
X ≥ 50% conserved