

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

Sequence logo:

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

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

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

[illegible]

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592

logo

NTDB id 54539 BN389 RS07610 WP 003727413.1
NTDB id 355 BSU 27480 NP 390625.1
consensus



678
676

logo

NTDB id 54539 BN389 RS07610 WP 003727413.1
NTDB id 355 BSU 27480 NP 390625.1
consensus



763
761

logo

NTDB id 54539 BN389 RS07610 WP 003727413.1
NTDB id 355 BSU 27480 NP 390625.1
consensus



798
798

- X non conserved
- X similar
- X $\geq 50\%$ conserved