

logo

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NTDB id 1018	ACIAD RS01685 WP 004920473.1	MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIEEVHLSPLSLAETISAEFAEYPFDLDDVYDTSQI	80
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NTDB id 1019	ACIAD RS04205 WP 004922051.1	0
NTDB id 1307	DR RS10055 WP 010888596.1	0
NTDB id 1086	NMV RS00245 WP 002218487.1	0
NTDB id 1116	NGFG RS10615 WP 003690200.1	0
NTDB id 1185	GTF74 RS11940 WP 000350195.1	0
NTDB id 1176	A1552VC RS01160 WP 000350195.1	0
NTDB id 125526	HZ99 RS15535 WP 038444143.1	0
NTDB id 1283	ACE17W RS02630 WP 010947729.1	0
NTDB id 1259	GCO85 RS10305 WP 010947729.1	0
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NTDB id 1060	ABD1 RS04380 WP 000355489.1	0
NTDB id 1040	HON27 RS13360 WP 000355489.1	0
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		LKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSKLVKVEPIIVEHDKLERLLSEHFVEETHFNFDTEEL	160
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 NTDB id 1060 ABD1 RS04380 WP 000355489.1
 NTDB id 1040 HON27 RS13360 WP 000355489.1

Y P S S L L E M G F T P E D L A Q P E F Q I F E P V G C H D C R E G Y K G R V G I Y E V M K I T P E I S K I T I M E D G N A L . . E I A A T A E T L G F N N L R 547
 I P K Q S L L E M G F T E Q D L A H P D F R V F Q P V G C L E C R E G Y K G R V G I Y E V M K V T P E I S K I T I M E D G N A L . . E I A A A S E K L G F N N L R 547
 T P D G K G R R A V I E V L I N T P L A A D L I R K G E V H E L K G L M K R S T D L G M Q T F D 316
 T H D G Q S R R A A V E V L I N T P L L S D Y I R Q G E I Y K I K E L M A R S G E L G M Q T F D 319
 R A D G Q G R V L A Y E I L L A N P A V R A L I R E G K T F Q I T S V M Q T G A R E G M V T M D 322
 T H D G N G R V A S H E I L I A N P A V R N L I R E N K I T Q I N S V L Q T G Q A S G M Q T M D 318
 T H D G N G R V A S H E I L I A N P A V R N L I R E N K I T Q I N S V L Q T G Q A S G M Q T M D 318
 R . V G G G R V A C H E I M L A T P A I R N L I R E D K V A Q M Y S I I Q T G A A H G M Q T M E 317
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 K . V G G G R V A A E I M L G T P A I R N L I R E D K V A Q L Y S A I Q T G G A L G M K T L D 317
 K . N K G G R V A A L E I M M C T G A I R N L I R E D K I A Q M Y S S I Q T G Q A K G M Q T L D 317
 K . N K G G R V A A L E I M M C T G A I R N L I R E D K I A Q M Y S S I Q T G Q A K G M Q T L D 317
 K . I G G G R V A A H E I M I G T P A I R N L I R E D K V A Q M Y S A I Q T G G S L G M Q T L D 317
 K . V G G G R V A A H E I M I G T P A I R N L I R E D K V A Q M Y S S I Q T G G S L G M Q T L D 317
 K . E G G G R V A A H E I M I G T P A I R N L I R E G K V A Q M Y S S I Q T G A T H G M T T L D 317
 K . N G G G R V A A H E I M I G I P A I R N L I R E N K V A Q M Y S A I Q T G A N H G M T T L D 317
 K . N G G G R V A A H E I M I G I P A I R N L I R E N K V A Q M Y S A I Q T G A N H G M T T L D 317
 K . N G G G R V A A H E I M I G I P A I R N L I R E N K V A Q M Y S A I Q T G A N H G M T T L D 317

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 NTDB id 1081 FDQ49 RS05335 WP 006581387.1
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R S G L K K V M Q G V T S L Q E I N R V T S E 570
 R S G L K K V M Q G V T S L Q E V N R V T S E 570
 Q A L Y Q L V V D D K I S E Q D A L H S A D S A N D L R L M L K T K R G D D Y . . . G S G S L Q N V K I D M E 368
 Q A L Y N L Y V Q G E I T Y E D A L L H A D S S N D L R L M I K L G S E T D G E H L T S V S . Q G L T L E M S D D D P G R S F R 381
 Q A L F D L Y D H Q H I S Y A D A L K H A D S P N N L R L Q I K L S E E A N K P L D N R E R M S E M E R K M T F D G Q R R N L K 383
 A Y L A G L F R R H L I T Y D K G L E R A I D P K E F A R L A N D A S A G L G G A S M T P A A P P V P A Y N S A S S S T G R G G D F G R G A A P A T 398
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 Q H L T E L V N K N I I S K H T A H E A A L N K S L F 344
 Q H L T E L V N K N I I S K H T A H E A A L N K S L F 344
 M C L K G L V A K G L I S R E N A R E K A K I P E N F 344
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 Q S L K G L V A R G V I S P Q T A R T A A K Q P E S F L 345
 Q S L K G L V A R G V I S P Q T A R T A A K Q P E S F L 345

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NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
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NTDB id 1040 HON27 RS13360 WP 000355489.1	345
consensus		

 non conserved
 similar
 ≥ 50% conserved