

[illegible]

NTDB id 1399 DSB67 RS15600 WP 010645755.1	AASHHHLLTADALYPPLLKQTVAPPPLLFVKGEASC.....LPQQQTAMVGSRNASVDGLQHTRTFASDLVQHDLI	146
NTDB id 1152 A1552VC RS00200 WP 000654772.1	SAANNRYIVTPHCPLYRRLKELNSSPPVLFIIEGIWEA.....VHDPAAVAIVGSRNASVDGRQIARQFATELAQSGLV	144
NTDB id 1314 DR RS00625 WP 010886768.1	AAERGVTLLGRGLPGYPAALEALGDPPAVLWVRGAGGAELLAGLGTVPHSVGVIGTRAASPHALALTRTIAGELAAAGVL	160
NTDB id 68824 ABWW09 RS09345 WP 353300258.1	AEAMGVRRLLGLWEEGFPEGLKALPQPPTHLYLKGELEP.....PEREAAVALVGTRRASPWALAFARRLARELAEAGLW	126
NTDB id 1037 TT RS09535 WP 011174226.1	AEALGVRRLLGLWEEGFPEGLKALPQPPTHLYLKGELEP.....PEREAAVALVGTRRASPWALAFARKLARELS EAGLW	126
consensus	* * * * *	

[illegible]

NTDB id 1399 DSB67 RS15600 WP 010645755.1	SGLSLGVVVEAAEKSGSLITARYALEQGREVFALPASINAPNASGGNQLIQNGACLVQNTQEVLNEIQSLLDWSINQSL	301
NTDB id 1152 A1552VC RS00200 WP 000654772.1	SGLSLGVVVVEAAEKSGSLITARYALEQGREVFVVPGSIFNAASQGSNQLIRQGACLVQSVQQIHQELKNALTWSLSEQV	299
NTDB id 1314 DR RS00625 WP 010886768.1	AALSAGVLVVEGERKSGSLITATHALDCGRTVFAVPGRAGDPRASGPHALIRDGAVALTESAQDVLTEL...NWGLAPAP	312
NTDB id 68824 ABWW09 RS09345 WP 353300258.1	AGLCRAVVVVVEAPLDSCALITARHALELGKEVLAVPGRPTDERSLGANRLIQDGAYPVLSAEDVLSYL...GMSARPKP	273
NTDB id 1037 TT RS09535 WP 011174226.1	AGLCRAVVVVVEAPLDSCALITARHALELGKEVLAVPGRPMDESLGANRLIQDGAYPVLSAEDVLSYL...GMSARPKP	273
consensus	* * * * *	

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NTDB id 1314 DR RS00625 WP 010886768.1
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NTDB id 1037 TT RS09535 WP 011174226.1
consensus

 non conserved
 similar
 $\geq 50\%$ conserved