

NTDB id	1169 A1552VC RS11075 WP 000648511.1	MKATQ.....TLPLKNYRWKGINSNGKK.VSGQLLAISEIEVRDKLKDQHITQIKKLKKGVSVLAR.LTHRVKSKDITI	72
NTDB id	145643 F0RC14 RS13415 WP 005479682.1	.MKTT.....TPQLKNFRWKGVNSGKK.TSGQTLLAMSEIEVRRERLDQAHIKIKKLKKSISFLT.KLSHRVKGKDITV	71
NTDB id	1402 DSB6F RS12670 WP 010643257.1	.MKTII.....APQLKNYRWKGINSNGKK.TSGNVLAMTEIEVRRERLDQAHIKIKKLKKSISFITK.ISHRVKGKDITI	71
NTDB id	1112 NAGFS RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRIFEGRHSAASKLVNGEVSAFTEEAREKKLAKRGIRPLQITRV....KTSSRKKITQEDITV	74
NTDB id	1113 AAA85695.1 219..1451()	.MAKNGGFSLFAKKEKRIFEGRHSAASKLVNGEVSAFTEEAREKKLAKRGIRPLQITRV....KTSSRKKITQEDITV	74
NTDB id	1252 GC085 RS07730 WP 011213805.1	.MDKN.....SPP.LTFHYQGINKAQGK.MEGDIQARS LAIAKADLRKQGI VTNKVVKRKRPFDR.KNKKITQA DITV	71
NTDB id	1198 PSJM300 O3950 AFN76868.1	.MAQQ.....AIKNSVF TWEGLDRQGAK.IKGE LSCVSPALVKAQLRKQGINPQKVREKKSIS.FS..AGKKIKPM DIAL	70
NTDB id	1016 ACTIAD RS01680 WP 004920476.1	MAA KK.....TQVM P VFAYEGVD RKIGIK.LKGELPARNMALAKVT LRKQG ITIRTIREKRKNIFEGLLKKVKV PL DIA I	73
NTDB id	1059 ABD1 RS01610 WP 000279215.1	MAVKK.....AQMMPTFAYEGVD RKG VK.IKGELPAKNMALAKVT LRKQGVTVNRNIREKRKNILEGLFKKKV TTLDITI	73
consensus		* * * * *	

[illegible][illegible]

NTDB id 1169	A1552VC	RS11075	WP 000648511.1	AIAAAITGLKALRKNSFQIRLKTSTRGLKFPITIGNVLAKASTAKFSRTRLAT	SFAAGIPILASLKTITAKTSGNVHVFETA	IN	312
NTDB id 145643	FORC14	RS13415	WP 005479682.1	CSISVVFISARVLSKRSDSFRLMLNRSVLKFPVLGAVLSKAAIAKFSRTRLAT	SFTAGIPILTSLKTTSTKSGNMHYQLAIE		311
NTDB id 1402	DSB67	RS12670	WP 010643257.1	SFVSLISGKIMSQRSDSFRLLAARSSLRVPIILGPVLSKAAIAKFSRTRLAT	SFSAGIPILTALKTTSTKSGNMHYQLAIE		311
NTDB id 1112	NGFG	RS09215	WP 003689811.1	ALGFAIYGFLLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATL	FAAGVPLVDVLDSTAGAAGNLIYEAEATR		314
NTDB id 1113	AAA85695.1	219..1451()		ALGFAIYRFLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATL	FAAGVPLVDVLDSTAGAAGNLIYEAEATR		314
NTDB id 1252	GC085	RS07730	WP 011213805.1	ALGGVIVYSFFYAKNHSLEFAQITDRVMLKLPVIGPILEKAAIARFARTLSIT	FAAGLPLVEALKSVAGATGNIIYAKATD		311
NTDB id 1198	PSJM300	03950	AFN76868.1	CFVGAAYAFKQAHTRSEKFRNWFDRMLLKVPVVGDIYKSSVARFARTLSTT	FAAGVPLVDALDSVAGATGNVVERSATN		310
NTDB id 1016	ACIAD	RS01680	WP 004920476.1	VITGAATAGFLEARKRSKRFRDQDLRLVLRPLIVGDLVYKAI	IARYSRTRLATFAAGVPLIDALASTAGATNNIVYEQAVL		313
NTDB id 1059	ABD1	RS01610	WP 000279215.1	AIGAVIAAFLAELKRSKKFRDGLDKLALKLPFGDLVYKAI	IARYSRTRLATFAAGVPLIDALESTAGATNNVIEYKAVM		313
consensus				*****	*****		

