

NTDB id	1169 A1552VC RS11075 WP 000648511.1	MKAT.....QTLP LKNYRWKGINSNGKK.VSCQMLAISEIEVRDKLKDQHITQIKKLKKSVS.LAR.LTHRVKSKDITI	72
NTDB id	992836 AAIA71 RS03635 WP 005446256.1	.MKT.....IAPQLKNYRWKGINSNGKK.TSGNVLAMTEVEVRERLDAHQHIKI KKLKKSISFITK.ISHRVKVGKDITV	71
NTDB id	1402 DS867 RS12670 WP 010643257.1	.MKT.....IAPQLKNYRWKGINSNGKK.TSGNVLAMTEIEVRERLDAHQHIKI KKLKKSISFITK.ISHRVKVGKDITI	71
NTDB id	1113 AAA8569S.1 219..1451()	.MAKNGGFSLF AKKEKR FIFEGRHSASDKLVNGEVS AFTEE EARKKLAKRGIRPLQITRV....KTSSRKKITQEDITV	74
NTDB id	1112 NCFG RS09215 WP 003689811.1	.MAKNGGFSLF AKKEKR FIFEGRHSASDKLVNGEVS AFTEE EARKKLAKRGIRPLQITRV....KTSSRKKITQEDITV	74
NTDB id	1252 GC085 RS07730 WP 011213805.1	.MDKN.....SPPLLTFHYQGINKAGQK.MEGDIQARS LAIAKADLRKQGI VTNKVVKRRKP.FDR.KNKKITQA DITV	71
NTDB id	1198 PSJM300 O3950 AFN76868.1	.MAQQ.....AIKNSVF TWEGL DRQGAK.IKGEL SCV SPALVKAQL RKQGINP QKV EKKSIS.FS..AGKKIKPM DIAL	70
NTDB id	1016 ACTAD RS01680 WP 004920476.1	MAAKK.....TQVM P VFAYEGVD RKG I.KGELPARNMALA KVTL RKQGITIRTIRE KRKNIFE GLLK KKVK PL DIAI	73
NTDB id	1059 ABD1 RS01610 WP 000279215.1	MAVKK.....AQMMPTFAYEGVD RKG I.KGELPAKNMALA KVTL RKQGVTVRNIREKRKNILEGLFKKKVTLLDITI	73
consensus		* ** * * * * *	

[illegible][illegible]

NTDB id 1169	A1552VC	RS11075	WP 000648511.1	AIAAAITFLKALRKNSFQIRLKTSTRGLKFPITIGNVLAKASTAKFSRTRLAT	SFAAGIPILASLKTITAKTSGNVHVFETA	IN	312
NTDB id 992836	AAIA71	RS03635	WP 005446256.1	SFVSLIISGKIMSQRSDSFRLLTARSSLRVPILGPVLSKAAIAKFSRTRLAT	SFSAGIPILTALKTTSKTSGNMHHYQLAIE		311
NTDB id 1402	DSB67	RS12670	WP 010643257.1	SFVSLIISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTRLAT	SFSAGIPILTALKTTSKTSGNMHHYQLAIE		311
NTDB id 1113	AAA85695.1	219..1451()		ALGFATYRFLKLKARSIKIQRRMDATILLRMPIFGDIVRKGTIARWGRTTATL	FAAGVPLVDVLDSTAGAAGNLIYEEATR		314
NTDB id 1112	NGFG	RS09215	WP 003689811.1	ALGFATYRFLKLKARSIKIQRRMDATILLRMPIFGDIVRKGTIARWGRTTATL	FAAGVPLVDVLDSTAGAAGNLIYEEATR		314
NTDB id 1252	GC085	RS07730	WP 011213805.1	ALGGVIVYSFFYAKNHSLEFAQITDRVMLKLPVIGPILEKAAIARFARTLSIT	FAAGLPLVEALKSVAGATGNIIYAKATD		311
NTDB id 1198	PSJM300	03950	AFN76868.1	CFVGAAYAFKQAHTRSEKFRNWFDRMLKVPVVGDIYKSSVARFARTLSTT	FAAGVPLVDALDSVAGATGNVVERSATN		310
NTDB id 1016	ACIAD	RS01680	WP 004920476.1	VIGAAIAGFLEARKRSKRFRDQDLRLVLRLPVIGDLVYKAI	IARYSRTRLATFAAGVPLIDALASTAGATINN	TVYEQAVL	313
NTDB id 1059	ABD1	RS01610	WP 000279215.1	AIGAVIAAFLAEEKRSKKFRDGLDKLALKLPIFGDLVYKAI	IARYSRTRLATFAAGVPLIDALESTAGATINN	VIYEKAVM	313
consensus				*****	*****	*****	*****

