

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLRTAARLLRDRGLFVAETKEPGKGLQAEVRIPALERGPGLK	65
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKKLLKGSVSLLAR.....LTHRVKSK	68
NTDB id 1402	DSB67 RS12670 WP 010643257.1	.MKT.....IAPQLKNYRWKGINSSGKK	TSGNVLAMTEIEEVRERLDAQHIKIKKLLKGSISFITK.....ISHRVKSK	67
NTDB id 1113	AAA85695.1 219..1451()	.MAKNGGFSLFAKKEKRFIFEGRHSASDKLVN	GEVSAFTEEEARKKLLAKRGIRPLQITRVK.....T....SSSKRKITQE	70
NTDB id 1112	NGFG RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRFIFEGRHSASDKLVN	GEVSAFTEEEARKKLLAKRGIRPLQITRVK.....T....SSSKRKITQE	70
NTDB id 1252	GC085 RS07730 WP 011213805.1	.MDK.....NSPPLTLFHYQGINKAGQK	MEGDIQARSLATAKADLRKQGITVNTKVVKKRKPLFDR.....KNKKITQA	67
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAK.....KTQVMPVFAYEGVDRKGIK	LKGELPARNMALAKVTLRKQGITTIREKRKNIFEG....LLKKKKVKPL	69
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVK.....KAQMMPTFAYEGVDRKGIVK	IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG....LFFKKKVTTL	69
NTDB id 1129690	G3R43 RS26265 WP 009315898.1	.MAD.....KALKTSVFIWEGTDRKKGAK	VKGELTGCNPMLVKAHLRKKGINPLKVRKKGISLLG.....AGKKVKPM	66
NTDB id 1198	PSJM300 03950 AFN76868.1	.MAQ.....KAIKNSVFTWEGLRDQGA	IKGELSGVSPALVKAQLRKKGINPQKVRKKSISLFS.....AGKKIKPM	66
consensus		*** * * * * *	! * * * * *	***

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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
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NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1129690 G3R43 RS26265 WP 009315898.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus
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