

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLRTAARLRDRGLFVAETKEPGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKKLKKGVSLSLLAR....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAAPQLKNYRWKGINSSGKK	TSGNVLAMTEIEVRRLDAQHIKIKKLKKGISISFITK....ISH	62
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVN	GEVSFAFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVN	GEVSFAFTEEEARKKLAKRGIRPLQITRVK....T....SSKR	65
NTDB id 291494	DGN11 RS06755 WP 007973903.1	MSVARSSIIKKQPVDRNVSEIPFVWEGTDKRGVK	MKGEQQARNANMLRAELRRQGITPTVVKTKPKPLFGA....AGK	74
NTDB id 1252	GC085 RS07730 WP 011213805.1	MDKNSPPLLTFHYQGINKAGQK	MEGDIQARSLAIKADLRKQGIVTNKKVKKRKPLFDR....KNK	62
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKAIAKNSVFTWEGLDKRGAK	IKGELSGVSPALVKAQLRKQGINPQKVRKKSISLFS....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYEGVDRKGIK	LKGELPARNMALAKVTLRKQGITTRTIREKRKNIFEG...LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKAQMMPTFAYEGVDRKGVK	IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG...LFKK	64
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[illegible]

NTDB id	1029	TT RS02230 WP	O11228203.1	TSGGLDLILDRLASFLKEELELRGKIRSAMTYPVIVFVFAVGVAIFLLTGIVPQFAQLITDGLSELPLLTRFLIAVSDDL	219
NTDB id	1169	A1552VC RS11075 WP	000648511.1	MSGNLPEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMLTMVIPEFESMFKGFGAELPWFTQQVKLSHWV	223
NTDB id	1402	DSB67 RS12670 WP	010643257.1	QSCNLSQVFERLATYREKSEQLRAKVIKALIYPAMVVLVALGVSYLMLTRVIPEFEKMFTGFADLPAPTQMVLINLSAWT	222
NTDB id	1112	NNGF RS09215 WP	003689811.1	TGGVLESLLDKLATYKEKTQAIRKKVKLTALTYPVSVIAVAIGLVFMVMIFVLPAFKEVYANMGAELPPLTQTVMMSDF	225
NTDB id	1113	AAA85695.1	219..1451()	TGGVLESLLDKLATYKEKTQAIRKKVKLTALTYPVSVIAVAIGLVFMVMIFVLPAFKEVYANMGAELPALTOTVMMSDF	225
NTDB id	291494	DGN11 RS06755 WP	007973903.1	GAGVLETLDVTANYKENIEALKKKIKKALFYPMVMMAVAIIVSGILLVFPVPQFEDVFKGFGAELPAFTQMIVAASRFM	234
NTDB id	1252	GCO85 RS07730 WP	011213805.1	KSGSLDIMLDKVATYKEKETIKKKIKKALTYPITAVMVALLLVTAGLLIYVVPQFESLFKGFADLPAMTRAIVITMSEFM	222
NTDB id	1198	PSJM300 O3950 AFN76868.1		QCSGLETLIDRVATYKEKTEALKAKIKKANMPYIAVVLVAVIVTAILLIKVVPQFQDVVANFGAELPAFTLMVIGLSEAL	221
NTDB id	1016	ACIAD RS01680 WP	004920476.1	QCSGLEIMLDRVAVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIIILMVKVPVPFKDLFTSFGADLPAPTQLVNNMSNWM	224
NTDB id	1059	ABD1 RS01610 WP	000279215.1	QSAGLEIMLDRVATYKEKSELLKQKIKKAMKYPATVIVVAIVVTIIILMVKVPVPFDLDFASFGADLPAPTQMNVNSKWM	224
consensus				* ! ***** * ** *** **** ***** * * * * * * * * * * *	

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