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NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDHQGRL.VEATTEAEDLRTAARLLRDRGTFVAETKEPFGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLPLKNYRWKGINSNQKK.VSCQMLATSEIEVRDKLKDQHIQIKKLLKKSVSILA....RLTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAAPQLKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIKIKKLLKKSISFIT....KISH	62
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSATFEEEBARKKLAKRGIPLQITRVK.....T....SSKR	65
NTDB id 1112	NGFG RS09215 WP 003689811.1	MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSATFEEEBARKKLAKRGIPLQITRVK.....T....SSKR	65
NTDB id 1252	GCO85 RS07730 WP 011213805.1	MDKNSPPLTLTFHYQCINKAGQK.MBCDIQARSLATAKADLRKQGITVTKVVKRKPPLFD....RKNK	62
NTDB id 1140420	DQN96 RS17090 WP 100438366.1	MSVSRSAIKKEPVARATMELQPFVVEGTDKRCVK.MKGEQLAKNANLLRAELRRQGGINPGQVVPKSKPLFG....AAGS	74
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKAIKNSVFTWEGLDRQCAK.IKGELSGVSPALVMAQLRKQGINPQKVRKKSISLFS.....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQVMPVFAYECVDRKCIK.IKGELPARNMALAKVTLRKQGITRTIRKKNIFEG....LLKK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKAQMPTFAYEGVDRKCVK.IKGELPAKNMALAKVTLRKQCVTVRNIREKKNILEG....LFKK	64
consensus		*****	

logo

NTDB id 1029	TT RS02230 WP 011228203.1	GPGLKDLAIFSRQLATMIGACITLLQALAILERQTENRKFRREILKQVHTDVEGCMAFSEALSKHK.IFSRLYVNLVRAGE	139
NTDB id 1169	A1552VC RS11075 WP 000648511.1	RVKSKDITILTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLVVDLVETGE	143
NTDB id 1402	DSB67 RS12670 WP 010643257.1	RVKGGKDITIFTRQIATMTGMTGVPIVQALKLVSENHKKAEMKSLMSVTRAVEACTPMSKAMRTASTHFDPLYTDLIATGE	142
NTDB id 1113	AAA85695.1 219..1451()	KITQEDITVFRQLSTMINAGIPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQCSSLSRAFSNHPKYFDRFYCNLVAAGE	145
NTDB id 1112	NGFG RS09215 WP 003689811.1	KITQEDITVFRQLSTMIKAGIPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQCSSLSRAFSNHPKYFDRFYCNLVAAGE	145
NTDB id 1252	GCO85 RS07730 WP 011213805.1	KITQADITVFSRQLATMIESGIPLVQAFDIVAKQSNKRLKDLTETIKHDIETCLTLAESLIKHPLYFNELECNLVDAGE	142
NTDB id 1140420	DQN96 RS17090 WP 100438366.1	PIKPKDIAFFSRQMATMMKSGVPIVSALDITASGHKNPRMKKLVDITRTDIEGSSMHEATSKHPVQFDELYRNLVRAGE	154
NTDB id 1198	PSJM300 03950 AFN76868.1	KIKPMDIALFTRQMATMMKAGVPLIQSFDIIGECFDPNPMRKLVDLQKQEVAAACNSFASSLRRKPKQYFDDLYCNLVDSGE	141
NTDB id 1016	ACIAD RS01680 WP 004920476.1	KVKPLDIAIFTRQLATMMKAGVPLVQAFEIVAEGLNPAMRDVVLGLKNEVEGCNIFAGALRKYPQYFDNLFCSLIESGE	144
NTDB id 1059	ABD1 RS01610 WP 000279215.1	KVTTLDITIFTRQLATMMKAGVPLVQCFEIVAEGLNPAMREVVLGLKEVEGCSTFASALRKYPQHFDNLFCSLVESGE	144
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logo

NTDB id 1029	TT RS02230 WP 011228203.1	TSGGLDLILDRLASFLEKELELRGKIRSAMTYPVIVFVFAVGVAFLLTGIVPQFAQLTLDLCSELPLLLTRFLIAVSDLL	219
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MSGNLPVFERLATYREKSEQLRKAVIKALIYPSMVVLVALCVSYLMLTMVIPFESMFKCGFAGELPWFQTQVLKLSHVV	223
NTDB id 1402	DSB67 RS12670 WP 010643257.1	QSGNLSQVFERLATYREKSEQLRKAVIKALIYPSMVVLVALCVSYLMLTRVPIPEFEKMTGFCGADLPFAFTQMVNLISAWT	222
NTDB id 1113	AAA85695.1 219..1451()	TGGVLESLLDKLATYKEKTQAIRKKVKLTALTYPVSVIATAICLVFVMMIFVLPFAFKEVYANMGAELPALTQTVMDSDF	225
NTDB id 1112	NGFG RS09215 WP 003689811.1	TGGVLESLLDKLATYKEKTQAIRKKVKLTALTYPVSVIATAICLVFVMMIFVLPFAFKEVYANMGAELPALTQTVMDSDF	225
NTDB id 1252	GCO85 RS07730 WP 011213805.1	KSGSLDIMLDKVATYKEKTEIIRKKIKKALTYPVAVMVALLVTAAGLLIYVVPQFESLFKCGFADLPAMTRAVITMSEFM	222
NTDB id 1140420	DQN96 RS17090 WP 100438366.1	GAGVLETVLDIVATYKENIEALKKIKKALFYPIVVVVAMLVSGIMLVFVVPQFEDVFKSFGAELPAFTQMVVNLISRFM	234
NTDB id 1198	PSJM300 03950 AFN76868.1	QSGSLETLLDRVATYKEKTEALKKIKKANFYPIAVVIVAVIVTATILLIKVVPQFQDVFAFNGAELPAFTLMVIGLSEAL	221
NTDB id 1016	ACIAD RS01680 WP 004920476.1	QSGSLEIMLDRVAVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFGADLPFAFTQLVVNMNSNM	224
NTDB id 1059	ABD1 RS01610 WP 000279215.1	QSGALETMLDRVAVYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVQDLFASFGADLPFAFTQMVVNMNSNM	224
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