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NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKN	72
NTDB id 1402 DSB67 RS12670 WP 010643257.1	APPLTKFYEINSAGKVGKESAEAKLRKQGLPSKIKKKRKSLEFRSkkKITQDITY	71
NTDB id 1113 AAA85695.1 219..1451()	MAKNGGFSFLFAKKEKRFIFEGRHSASDKLVNGEVSFAFTEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	74
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFSFLFAKKEKRFIFEGRHSASDKLVNGEVSFAFTEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	74
NTDB id 173683 A1D13 RS07680 WP 027225016.1	MDKN.....SPPLLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKQGIVTNKVVKRRKPLFDR.KNKKITQADITV	71
NTDB id 1252 GC085 RS07730 WP 011213805.1	MDKN.....SPPLLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKQGIVTNKVVKRRKPLFDR.KNKKITQADITV	71
NTDB id 1198 PSJM300 03950 AFN76868.1	MAQK.....AIKNSVFTWGLDRQGAQK.IKGELSGVSPALVKAQLRKQGINPQKVRKKKSISLFS..AGKKIKPMDIAL	70
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAAKK.....TQVMPVFAYEGVDRKGIK.LKGELPARNMALAKVTLRKQGITIRTIREKRKNIFEGLLKKVKPLDIAI	73
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAVKK.....AQMMPTFAVEGVDRKGVK.IKGELPAKNMALAKVTLRKQGITVRNIREKRKNILEGLFKKKVTTLDTITI	73
consensus	*** * * * * ! * * ! * ! * * * * ! * * * * ! * * * * ! * * * * ! * * * *	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	FTRQLATMkaGyPLVQAFIVAGHNPMEKMTIKVEGGLAALRKHPYFDLYCNVAGFESGGLIEM	152
NTDB id 1402 DSB67 RS12670 WP 010643257.1	LTRQLATMITTGVPIVQALKLVGDNHRKAEMKSIILAQITKSVEAGTPLSKAMRTASAHFDTLYVDLIVETGEMSGNLPVEVF	151
NTDB id 1113 AAA85695.1 219..1451()	FTRQIATMLMTGVPIVQALKLVSENHKKAEKMSILMSVTRAVEAGTPMSKAMRTASTHFDPLYTDLIATGEQSGNLSQVF	154
NTDB id 1112 NGFG RS09215 WP 003689811.1	FTRQLSTMINAGLPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVLESLL	154
NTDB id 173683 A1D13 RS07680 WP 027225016.1	FTRQLSTMIKAGLPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGGVLESLL	151
NTDB id 1252 GC085 RS07730 WP 011213805.1	FSRQLATMIESGIPLVQAFDIVAKGQSNKRLKDLIETIKHDIETGLTLAESLIKHPLYFNELFCNLVDAGEKSGSLDIML	151
NTDB id 1198 PSJM300 03950 AFN76868.1	FSRQLATMIESGIPLVQAFDIVAKGQSNKRLKDLIETIKHDIETGLTLAESLIKHPLYFNELFCNLVDAGEKSGSLDIML	150
NTDB id 1016 ACIAD RS01680 WP 004920476.1	FTRQMATMMKAGVPLQSFDIIGEGFDNPNMRKLVDDLKQVEVAGNSFASSLRKKPQYFDDLYCNLVDSGEQSGSLETLL	153
NTDB id 1059 ABD1 RS01610 WP 000279215.1	FTRQLATMMKAGVPLVQAFEIVAEGLENPAMRDVVLGLKNEVEGGNTFAGALRKYPQYFDNLFCSLIESGEQSGSLEIML	153
consensus	***** ! * * * * ! * * * * ! * * * * ! * * * * ! * * * * ! * * * * ! * * * * ! * * * * ! * * * *	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	DRVATYKEKSELKKKKIKKALYPVIVVAVVGTALLIKVVPFESLFAEGAD
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