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NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK....KTNFDDITKKYGAERDKALADALALIEKD	FGKGSIMRLGER	.ANQKVS	VSSGSLALDI	ALGAGGYPKGR	74
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK....PKKLEEISKKFGAEREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALDI	ALGSGGYPKGR		73
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKK....PKKLEEISKKFGAEREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALDI	ALGSGGYPKGR		73
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKK....PKKLEEISKKFGAEREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALDI	ALGSGGYPKGR		73
NTDB id 159 SP RS09750 WP 001085462.1	.MAKK....PKKLEEISKKFGAEREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALDI	ALGSGGYPKGR		73
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKK....PKKLDEISKKFGAEREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALDI	ALGSGGYPKGR		73
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKK....PKKLDEISKKFGAEREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALDI	ALGSGGYPKGR		73
NTDB id 453 Spy49 1753c ACI62003.1	.MAKK....LKKNEEITKKFGERDKALDDALKNIEKD	FGKGVMLRGER	.AEQKVQVMSSGSLALDI	ALGAGGYPKGR		73
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR....IKKTEEITKKFGERDKALDDALKNIEKD	FGKGVMLRGER	.AEQKVQVMSSGSLALDI	ALGAGGYPKGR		73
NTDB id 115 BSU 16940 NP 389576.2MSDRQAALDMALKQIEKQFGKGSIMKLG	IEK	.TDTRISTVPSGSLALDI	TALGIGGYPRGR		58
NTDB id 625 LCA RS02525 WP 011374196.1MAKDERQAALDAALKKIEKNFGKGSIMRM	GEK	.VDTQVSTVSSGSLALDE	ALGVGGYPRGR		60
NTDB id 1313 DR RS12030 WP 010888966.1MSKDATKEISAPTDAKERSKATETAMSQIEKA	FGKGSIMKLGAE	.SKLDVQVVSTGSLSLDL	ALGVGGIPRGR		72
NTDB id 1245 Cj1673c YP 002345041.1MDNKRKSLDAALKS	LDKTFGKGTILRLGDK	.EVEQIDSIGTGSVGLDL	ALGIGGVPKGR		59
NTDB id 320192 D4J22 RS07055 WP 139547512.1MAIDEDKQKAIISLAIKQIDKVF	FGKGVMLRGER	.QVEKIDAI	ISTGSLGLDL	ALGIGGVPKGR	61
NTDB id 1211 C694 RS00795 WP 000963128.1MAIDEDKQKAIISLAIKQIDKVF	FGKGVMLRGER	.QVEKIDAI	ISTGSLGLDL	ALGIGGVPKGR	61
NTDB id 1214 NCTC11637 00359 SQJ03206.1MAIDEDKQKAIISLAIKQIDKVF	FGKGVMLRGER	.QVEKIDAI	ISTGSLGLDL	ALGIGGVPKGR	61
NTDB id 1334 RAOC RS04870 WP 004918236.1MAKT..ETTSPNDDKKKALAAVLEKLDKT	YKGTVMKLG	DGTVDHNIEV	IPSGSLGLDL	ALGVGGYPKGR	68
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKSSKNNTV..VKQVEPEQKKQALEAALA	QIEKQFGKGSIMKLGQT	.EQLDIESVSTGSLGLDL	ALGIGGLPMGR		76
NTDB id 1130 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQIEKSFGKGAIMKMDGS	QVEENLEVISTGSLGLDL	ALGVGGLPRGR		60	
NTDB id 1124 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQIEKSFGKGAIMKMDGS	QVEENLEVISTGSLGLDL	ALGVGGLPRGR		60	
NTDB id 1123 OK783 RS03845 WP 003688695.1MSDDKSKALAAALAQIEKSFGKGAIMKMDGS	QVEENLEVISTGSLGLDL	ALGVGGLPRGR		60	
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....K..KAASMSAEKQKALAAALA	QIEKQFGKGSIMKMGDA	.EVEPVQVVSTGSLGLDV	ALGVGGLPRGR		68
NTDB id 1165 A1552VC RS01565 WP 000344154.1MDENKQKALAAALGQIEKQFGKGSIMRLG	DN	.RAMDVETISTGSLSLDI	ALGAGGLPMGR		59
NTDB id 1142 GTF74 RS11550 WP 000344154.1MDENKQKALAAALGQIEKQFGKGSIMRLG	DN	.RAMDVETISTGSLSLDI	ALGAGGLPMGR		59
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDENKSKALQAALSQIEKQFGKNTVMRLG	DN	.TVQAVEAVSTGSLTLDI	ALGIGGLPKGR		59
NTDB id 1022 ACIAD RS06390 WP 004925639.1MDDNKSKALQAALSQIEKQFGKNTVMRLG	DN	.TVQAVEAVSTGSLTLDI	ALGIGGLPKGR		59
NTDB id 1205 PSJM300 05545 AFN77183.1MDENKRRALAAALGQIEKQFGKGVMLR	GDH	.DRQAIPSISTGSLGLDI	ALGIGGLPKGR		59
consensus	*** ** *	***! *!*****	*	***!****!!	!! *!*	

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consensus

 non conserved
 similar
 ≥ 50% conserved