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NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK...KTNFDDITKKYGA	ERDKALADALALIEKDFGKGSIMRLGEA	.ANQKVS	VSSGSLALDIALGAGGYPKGR	74						
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK...PKKLEEISKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQ	VMSGSLALDIALGSGGYPKGR	73						
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKK...PKKLEEISKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQ	VMSGSLALDIALGSGGYPKGR	73						
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKK...PKKLEEISKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQ	VMSGSLALDIALGSGGYPKGR	73						
NTDB id 159 SP RS09750 WP 001085462.1	.MAKK...PKKLEEISKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQ	VMSGSLALDIALGSGGYPKGR	73						
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKK...PKKLDEISKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQ	VMSGSLALDIALGSGGYPKGR	73						
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKK...PKKLDEISKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQ	VMSGSLALDIALGSGGYPKGR	73						
NTDB id 453 Spy49 1753c ACI62003.1	.MAKK...LKKNEEITKKFG	DERKALDDALKNIEKDFGKGAVMRLGER	.AEQKVQ	VMSGSLALDIALGAGGYPKGR	73						
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR...IKKTEEITKKFG	DERKALDDALKNIEKDFGKGAVMRLGER	.AEQKVQ	VMSGSLALDIALGAGGYPKGR	73						
NTDB id 115 BSU 16940 NP 389576.2MSDRQAALDMALKQ	IEKQFGKGSIMKLGEK	.TDTRIST	VPSGSLALDTALGIGGYPGR	58						
NTDB id 625 LCA RS02525 WP 011374196.1MAKDERQAALDAALKK	IEKNFGKGSIMRMGEK	.VDTQVST	VSSGSLALDEALGVGGYPGR	60						
NTDB id 1313 DR RS12030 WP 010888966.1MSKDATKEISAPTDAKERSKA	TETAMSQIEKAFGKGSIMKLGAE	.SKLDVQ	VVSTGSLSLDLALGVGGIPGR	72						
NTDB id 1057898 ACFJGY RS07435 WP 379969624.1MSTTDDKKKALQLVLEK	LDKTYGKGTVMTLGENSVDHT	IEVIPSGSLGIDLALGVGGYPGR	62							
NTDB id 1334 RAOC RS04870 WP 004918236.1MAKT.ETTSPNDDKKKALA	AVLEKLDKTYGKGTVMKLGDTVDHN	IEVIPSGSLGLDLALGVGGYPGR	68							
NTDB id 1245 Cj1673c YP 002345041.1MDNKRKSLDAALKS	LDKTFGKGTILRLGDK	.EVEQIDS	IGTGSVGLDLALGIGGVPKGR	59						
NTDB id 1214 NCTC11637 00359 SQJ03206.1MAIDEDKQKAI	SLAIKQIDKVF	FGKGA	LVRLGDK.QVEKIDA	ISTGSLGLDLALGIGGVPKGR	61					
NTDB id 1211 C694 RS00795 WP 000963128.1MAIDEDKQKAI	SLAIKQIDKVF	FGKGA	LVRLGDK.QVEKIDS	ISTGSLGLDLALGIGGVPKGR	61					
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKSSKNNTV.VKQVEP	EQKKQALEAALAQIEKQFGKGSIMKLGQT	.EQLDIES	VSTGSLGLDLALGIGGLPMGR	76						
NTDB id 1130 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQ	IEKSF	FGKGA	IMKMDGSQREENLEVI	ISTGSLGLDLALGVGGLPRGR	60					
NTDB id 1124 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQ	IEKSF	FGKGA	IMKMDGSQREENLEVI	ISTGSLGLDLALGVGGLPRGR	60					
NTDB id 1123 OK783 RS03845 WP 003688695.1MSDDKSKALAAALAQ	IEKSF	FGKGA	IMKMDGSQREENLEVI	ISTGSLGLDLALGVGGLPRGR	60					
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....K.KAASMSAEKQ	KALAAALAQIEKQFGKGSIMKMGDA	.EVEPVQ	VVSTGSLGLDVALGVGGLPRGR	68						
NTDB id 1165 A1552VC RS01565 WP 000344154.1MDENKQKALAAALGQ	IEKQFGKGSIMRLGDN	.RAMDVET	ISTGSLSLDIALGAGGLPMGR	59						
NTDB id 1142 GTF74 RS11550 WP 000344154.1MDENKQKALAAALGQ	IEKQFGKGSIMRLGDN	.RAMDVET	ISTGSLSLDIALGAGGLPMGR	59						
NTDB id 1022 ACIAD RS06390 WP 004925639.1MDDNKSKALQAALSQ	IEKQFGKNTVMRLGDN	.TVQAVE	AVSTGSLTLDIALGIGGLPKGR	59						
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDENKSKALQAALSQ	IEKQFGKNTVMRLGDN	.TVQAVE	AVSTGSLTLDIALGIGGLPKGR	59						
NTDB id 1205 PSJM300 05545 AFN77183.1MDENKSRALAAALGQ	IEKQFGKGAVMRMGDH	.DRQAIPS	ISTGSLGLDIALGIGGLPKGR	59						
consensus		*** **	**	***!!	*****	*	***!!	***!!	!!!	!!!	!!!

logo

NTDB id 600 KW2 RS01765 WP 011675413.1	...	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	IEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	IEE	388
NTDB id 194 SPR RS08825 WP 001085462.1	IEE	388
NTDB id 159 SP RS09750 WP 001085462.1	IEE	388
NTDB id 499 SM12261 RS08315 WP 078228268.1	IEE	384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	IEE	384
NTDB id 453 Spy49 1753c ACI62003.1	D..	378
NTDB id 408 SMU RS09530 WP 002262392.1	D..	383
NTDB id 115 BSU 16940 NP 389576.2	...	348
NTDB id 625 LCA RS02525 WP 011374196.1	...	355
NTDB id 1313 DR RS12030 WP 010888966.1	...	363
NTDB id 1057898 ACFJGY RS07435 WP 379969624.1	...	333
NTDB id 1334 RAOC RS04870 WP 004918236.1	...	343
NTDB id 1245 Cj1673c YP 002345041.1	...	343
NTDB id 1214 NCTC11637 00359 SQJ03206.1	...	347
NTDB id 1211 C694 RS00795 WP 000963128.1	...	347
NTDB id 1389 A4U84 RS07560 WP 021112712.1	...	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	...	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	...	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	...	348
NTDB id 1427 RS RS02760 WP 011000510.1	...	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	...	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	...	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	...	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349
NTDB id 1205 PSJM300 05545 AFN77183.1	...	347

consensus

 non conserved
 similar
 ≥ 50% conserved