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NTDB id 600 KW2 RS01765 WP 011675413.1	MATK....KKTNFDDITKKYGA	ERDKALADALALIEKD	FGKGSIMRLGEA	.ANQKVS	VSSGSLALDI	I	ALGAGGYPKGR	74	
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAK....KPKKLEEISKKFGA	EREKALNDALKLI	IEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDI	I	ALGSGGYPKGR	73		
NTDB id 228 SPD RS09265 WP 001085462.1	.MAK....KPKKLEEISKKFGA	EREKALNDALKLI	IEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDI	I	ALGSGGYPKGR	73		
NTDB id 194 SPR RS08825 WP 001085462.1	.MAK....KPKKLEEISKKFGA	EREKALNDALKLI	IEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDI	I	ALGSGGYPKGR	73		
NTDB id 159 SP RS09750 WP 001085462.1	.MAK....KPKKLEEISKKFGA	EREKALNDALKLI	IEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDI	I	ALGSGGYPKGR	73		
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAK....KPKKLDEISKKFGA	EREKALNDALKLI	IEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDI	I	ALGSGGYPKGR	73		
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAK....KPKKLDEISKKFGA	EREKALNDALKLI	IEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDI	I	ALGSGGYPKGR	73		
NTDB id 453 Spy49 1753c ACI62003.1	.MAK....KLKKNEEITKKFGD	ERRKALDDALKN	IEKDFGKGAVMRLGER	.AEQKVQVMSSGSLALDI	I	ALGAGGYPKGR	73		
NTDB id 408 SMU RS09530 WP 002262392.1	.MAK....RIKKTEEITKKFGD	ERRKALDDALKN	IEKDFGKGAVMRLGER	.AEQKVQVMSSGSLALDI	I	ALGAGGYPKGR	73		
NTDB id 115 BSU 16940 NP 389576.2MS	DRQAALDMALKQ	IEKQFGKGSIMKLG	IEK.TDTR	ISTVPSGSLALDI	I	ALGIGGYPRGR	58	
NTDB id 625 LCA RS02525 WP 011374196.1MAKD	ERQAALDAALKK	IEKNFGKGSIMRM	GEK.VDTQVSTVSSGSLALDEALGVGGYPRGR			60		
NTDB id 1313 DR RS12030 WP 010888966.1MSKDATKEISAPTD	AKERSKATETAM	SQIEKAFGKGSIMKLGAE	.SKLDVQVVSTGSLSLDLALGVGGIPRGR			72		
NTDB id 384599 FOU60 RS41795 WP 395808402.1MNKLT	EKLKAVAAAVAA	IEKQFGKGAVMPLG	GEVREQRVAVIPTG	SVGLDRALGVGGYPRGR		62		
NTDB id 1245 Cj1673c YP 002345041.1MDDN	KRKSALDAALKS	LDKTFGKGTILRLGDK	.EVEQID	ISTGSLGLDLALGIGGVPKGR		59		
NTDB id 1211 C694 RS00795 WP 000963128.1MAIDE	DKQKATISLAIKQ	IDKVF	FGKGA	LVR	LDGK.QVEKID	ISTGSLGLDLALGIGGVPKGR	61	
NTDB id 1214 NCTC11637 00359 SQJ03206.1MAIDE	DKQKATISLAIKQ	IDKVF	FGKGA	LVR	LDGK.QVEKID	ISTGSLGLDLALGIGGVPKGR	61	
NTDB id 1334 RAOC RS04870 WP 004918236.1MAK..TETTSPND	DKKKALAAVLEK	LDKT	YGKGTVMKLG	DGTVDHNI	IEVIPS	GSLGLDLALGVGGYPKGR	68	
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKSSKNNT..VVKQVEPEQ	KKQALEAALAQ	IEKQFGKGSIMKLG	QT.EQLDI	ESVSTGSLGLDLALGIGGLPMGR		76		
NTDB id 1130 NGFG RS03960 WP 003688695.1MSD	DKSKALAAALAQ	IEKSF	FGKGA	IMKMDGSQ	QEE	NEVISTGSLGLDLALGVGGLPRGR	60	
NTDB id 1124 NGFG RS03960 WP 003688695.1MSD	DKSKALAAALAQ	IEKSF	FGKGA	IMKMDGSQ	QEE	NEVISTGSLGLDLALGVGGLPRGR	60	
NTDB id 1123 OK783 RS03845 WP 003688695.1MSD	DKSKALAAALAQ	IEKSF	FGKGA	IMKMDGSQ	QEE	NEVISTGSLGLDLALGVGGLPRGR	60	
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....KKAASMSA	EKQKALAAALAQ	IEKQFGKGSIMKMG	DA.EVEPVQVVSTGSLGLDVALGVGGLPRGR			68		
NTDB id 1165 A1552VC RS01565 WP 000344154.1MDEN	KQKALAAALGQ	IEKQFGKGSIMRLG	DN.RAMDVET	ISTGSLSLDI	I	ALGAGGLPMGR	59	
NTDB id 1142 GTF74 RS11550 WP 000344154.1MDEN	KQKALAAALGQ	IEKQFGKGSIMRLG	DN.RAMDVET	ISTGSLSLDI	I	ALGAGGLPMGR	59	
NTDB id 1205 PSJM300 05545 AFN77183.1MDEN	KKRALAAALGQ	IEKQFGKGAVMR	GDH.DRQA	IP	ISTGSLGLDI	I	ALGIGGLPKGR	59
NTDB id 1022 ACIAD RS06390 WP 004925639.1MDDN	KS	KALQAALSQ	IEKQFGKNTVMRLG	DN.TVQAVEAVSTGSLTLDI	I	ALGIGGLPKGR	59	
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDEN	KS	KALQAALSQ	IEKQFGKNTVMRLG	DN.TVQAVEAVSTGSLTLDI	I	ALGIGGLPKGR	59	
consensus		** **	**	**!	*!!*	*!!*	*!!*	*!!*	

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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
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	...	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	...	343
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NTDB id 1165 A1552VC RS01565 WP 000344154.1	...	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	...	354
NTDB id 1205 PSJM300 05545 AFN77183.1	...	347
NTDB id 1022 ACIAD RS06390 WP 004925639.1	...	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349

consensus

 non conserved
 similar
 ≥ 50% conserved