

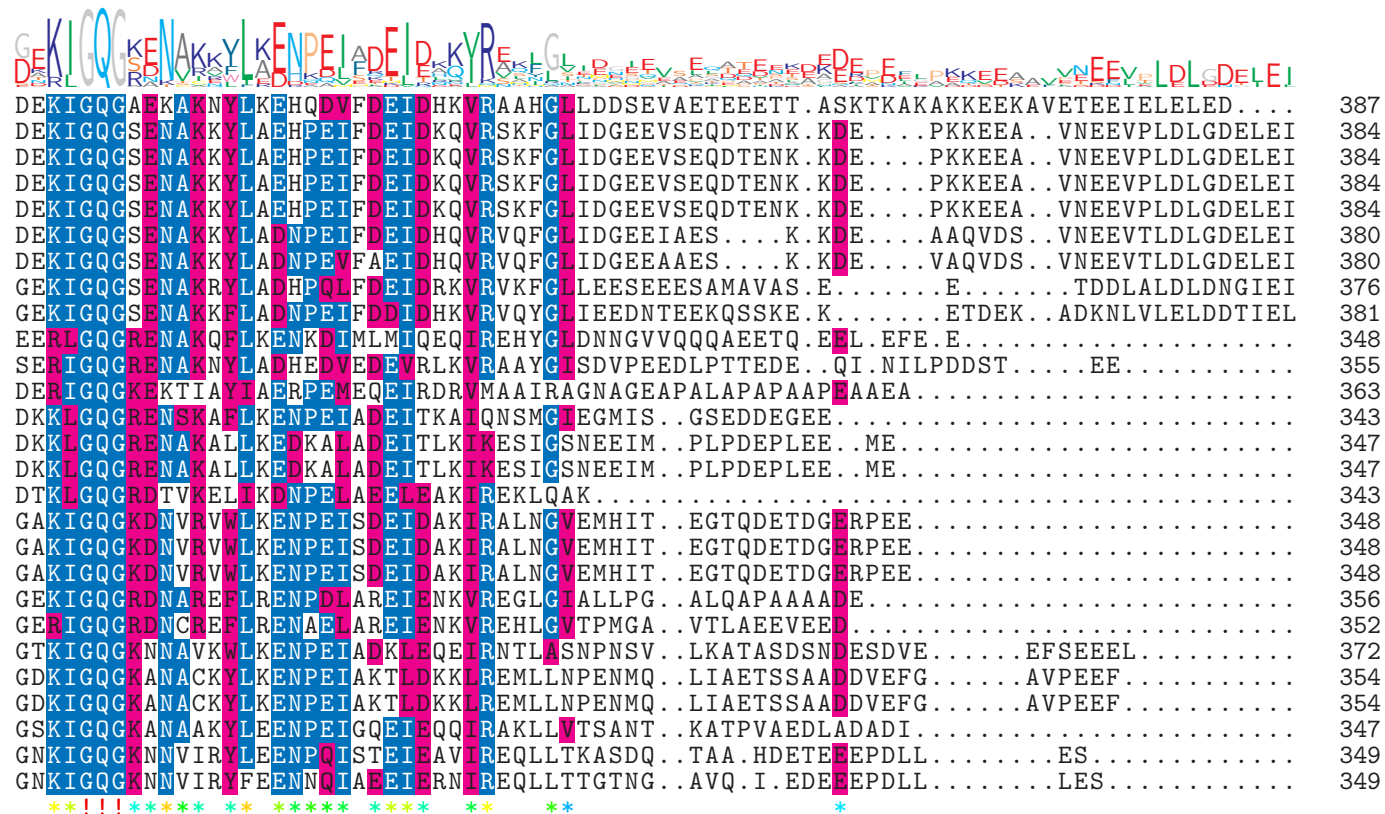
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NTDB id 600 KW2 RS01765 WP 011675413.1	MATKKKT....NFDDITKKYGAERDKALADALALIEKD	FGKGSIMRLGERAA.NQKVS	VSSGSLALDI	ALGAGGYPKGR	74		
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKKPK....KLEEISKKKFGAEREKALNDALKLIEKD	FGKGSIMRLGERA.EQKVQVMSSGSLALDI	ALGSGGYPKGR		73		
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKKPK....KLEEISKKKFGAEREKALNDALKLIEKD	FGKGSIMRLGERA.EQKVQVMSSGSLALDI	ALGSGGYPKGR		73		
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKKPK....KLEEISKKKFGAEREKALNDALKLIEKD	FGKGSIMRLGERA.EQKVQVMSSGSLALDI	ALGSGGYPKGR		73		
NTDB id 159 SP RS09750 WP 001085462.1	.MAKKPK....KLEEISKKKFGAEREKALNDALKLIEKD	FGKGSIMRLGERA.EQKVQVMSSGSLALDI	ALGSGGYPKGR		73		
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKKPK....KLDEISKKKFGAEREKALNDALKLIEKD	FGKGSIMRLGERA.EQKVQVMSSGSLALDI	ALGSGGYPKGR		73		
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKKPK....KLDEISKKKFGAEREKALNDALKLIEKD	FGKGSIMRLGERA.EQKVQVMSSGSLALDI	ALGSGGYPKGR		73		
NTDB id 453 Spy49 1753c ACI62003.1	.MAKKLK....KNEEITKKKFGDERRKALDDALKNIEKD	FGKGVMLGERA.EQKVQVMSSGSLALDI	ALGAGGYPKGR		73		
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKRIK....KTEEITKKKFGDERRKALDDALKNIEKD	FGKGVMLGERA.EQKVQVMSSGSLALDI	ALGAGGYPKGR		73		
NTDB id 115 BSU 16940 NP 389576.2MSDRQAALDMALKQIEKQ	FGKGSIMKLG	EKT.DTRISTVPSGSLALDI	TALGIGGYPRGR	58		
NTDB id 625 LCA RS02525 WP 011374196.1MAKDERQAALDAALKKIEKN	FGKGSIMRMGEKV.DTQVSTVSSGSLALDE	ALGVGGYPRGR		60		
NTDB id 1313 DR RS12030 WP 010888966.1MSKDATKEISAPTDAKERSKATETAMSQIEKA	FGKGSIMKLG	AES.KLDVQVVSTGSLSLDL	ALGVGGIPRGR	72		
NTDB id 1245 Cj1673c YP 002345041.1MDDNKRKSLDAALKS	LDKTFGKGTILRLG	DE.VEQIDSIGTGSVGLDL	ALGIGGVPKGR	59		
NTDB id 1211 C694 RS00795 WP 000963128.1MAIDEDKQKATISLAIKQIDKVF	FGKGVRLGDKQ.VEKIDSISTGSLGLDL	ALGIGGVPKGR		61		
NTDB id 1214 NCTC11637 00359 SQJ03206.1MAIDEDKQKATISLAIKQIDKVF	FGKGVRLGDKQ.VEKIDAISTGSLGLDL	ALGIGGVPKGR		61		
NTDB id 1334 RAOC RS04870 WP 004918236.1MAKTE..TTSPNDKKKALA	AVLEKLDKTYGKGTVMKLG	DGTVDHNI	IEVIPSGSLGLDL	ALGVGGYPKGR	68	
NTDB id 1130 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQIEKS	FGKGAIMKMDGSQ	QEEENLEVISTGSLGLDL	ALGVGGLPRGR	60		
NTDB id 1124 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQIEKS	FGKGAIMKMDGSQ	QEEENLEVISTGSLGLDL	ALGVGGLPRGR	60		
NTDB id 1123 OK783 RS03845 WP 003688695.1MSDDKSKALAAALAQIEKS	FGKGAIMKMDGSQ	QEEENLEVISTGSLGLDL	ALGVGGLPRGR	60		
NTDB id 321156 EAG14 RS22425 WP 121730207.1MDVKGTNP..ALPVNAEKAKALAAALAQIEKQ	FGKGTIMRLGEGEAIEDIQVVSTGSLGLDI	ALGVGGLPRGR		71		
NTDB id 1427 RS RS02760 WP 011000510.1MEDGKK..AASMSAEKQKALAAALAQIEKQ	FGKGSIMKMGDAE.VEPVQVVSTGSLGLD	VALGVGGLPRGR		68		
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKS	SKNNTVV..KQVEPEQKKQALEAALAQIEKQ	FGKGSIMKLGQTE.QLDIESVSTGSLGLDL	ALGIGGLPMGR	76		
NTDB id 1165 A1552VC RS01565 WP 000344154.1MDENKQKALAAALGQIEKQ	FGKGSIMRLG	DNR.AMDVETISTGSLSLDI	ALGAGGLPMGR	59		
NTDB id 1142 GTF74 RS11550 WP 000344154.1MDENKQKALAAALGQIEKQ	FGKGSIMRLG	DNR.AMDVETISTGSLSLDI	ALGAGGLPMGR	59		
NTDB id 1205 PSJM300 05545 AFN77183.1MDENKKRALAAALGQIEKQ	FGKGVMRMGDHD.RQAI	PSISTGSLGLDI	ALGIGGLPKGR	59		
NTDB id 1022 ACIAD RS06390 WP 004925639.1MDDNKS	KALQAALSQIEKQ	FGKNTVMRLG	DNT.VQAVEAVSTGSLTLDI	ALGIGGLPKGR	59	
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDENKS	KALQAALSQIEKQ	FGKNTVMRLG	DNT.VQAVEAVSTGSLTLDI	ALGIGGLPKGR	59	
consensus		** **	**	***!	*!!*****	*	***!!*!!*!!*!!*!!*!!*!!*!!

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 NTDB id 269 KZH43 RS08665 WP 001085462.1
 NTDB id 228 SPD RS09265 WP 001085462.1
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 NTDB id 159 SP RS09750 WP 001085462.1
 NTDB id 499 SM12261 RS08315 WP 078228268.1
 NTDB id 527 SMSK321 RS11065 WP 001085454.1
 NTDB id 453 Spy49 1753c ACI62003.1
 NTDB id 408 SMU RS09530 WP 002262392.1
 NTDB id 115 BSU 16940 NP 389576.2
 NTDB id 625 LCA RS02525 WP 011374196.1
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 NTDB id 1124 NGFG RS03960 WP 003688695.1
 NTDB id 1123 OK783 RS03845 WP 003688695.1
 NTDB id 321156 EAG14 RS22425 WP 121730207.1
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 NTDB id 1142 GTF74 RS11550 WP 000344154.1
 NTDB id 1205 PSJM300 05545 AFN77183.1
 NTDB id 1022 ACIAD RS06390 WP 004925639.1
 NTDB id 1074 ABD1 RS10195 WP 000344167.1
 consensus



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NTDB id 600 KW2 RS01765 WP 011675413.1	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	EIEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	EIEE	388
NTDB id 194 SPR RS08825 WP 001085462.1	EIEE	388
NTDB id 159 SP RS09750 WP 001085462.1	EIEE	388
NTDB id 499 SM12261 RS08315 WP 078228268.1	EIEE	384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	EIEE	384
NTDB id 453 Spy49 1753c ACI62003.1	ED..	378
NTDB id 408 SMU RS09530 WP 002262392.1	ED..	383
NTDB id 115 BSU 16940 NP 389576.2	348
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NTDB id 1313 DR RS12030 WP 010888966.1	363
NTDB id 1245 Cj1673c YP 002345041.1	343
NTDB id 1211 C694 RS00795 WP 000963128.1	347
NTDB id 1214 NCTC11637 00359 SQJ03206.1	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	343
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
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NTDB id 1427 RS RS02760 WP 011000510.1	352
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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