

logo

MEALIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPHTPV...KETNLQ.AEVVAISK...DSVS.EKEVK

NTDB id 105 BSU 25590 NP 390437.1MNWLN...QHKKAIILAASAAVFTAIM.IFLATGKNKEPV.....KQAVPTE.....TENTVV	49
NTDB id 1127 NGFG RS10560 WP 003703428.1	0
NTDB id 1126 NGFG RS09375 WP 003703428.1	0
NTDB id 1125 NGFG RS07215 WP 003703428.1	0
NTDB id 1108 NGFG RS06525 WP 003703428.1	0
NTDB id 1242 Cj0011c YP 002343483.1	0
NTDB id 1382 A4U84 RS07375 WP 062924234.1	0
NTDB id 1362 HI 1008 AAC22669.1	0
NTDB id 1041 HON27 RS14740 WP 005124713.1MQYSMRFWKNKYFLLIMLCLGVISPSVC.....	28
NTDB id 1069 ABD1 RS03060 WP 043960841.1MQYSMRFWKNKYFVLLIMLCLGVISPSVC.....	28
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	0
NTDB id 1150 A1552VC RS08580 WP 001166105.1	0
NTDB id 1294 VP RS04495 WP 021449594.1	0
NTDB id 1395 DSB67 RS04445 WP 010649155.1	0
NTDB id 613 LCA RS05340 WP 231952379.1MMWQDYWQRYRRFVWVPCLVLLGIAGGFYWHQSQQKQKVT.....QNEQLLATSASK.....TKPAT	58
NTDB id 1014 ACIAD RS13855 WP 004924483.1MSIR.....HFFTFFWATLL.....	15
NTDB id 36 MW RS08230 WP 001793467.1	MVLLYQFLLRYKDFLTQWK.LYIISAVVLIMVLIGFIFWRQDDYTSRN....FENKDNALKQSTSE..NSSLSKVEDVQ	72
NTDB id 20 SA RS08005 WP 001802022.1	MVLLYQFLLRYKDFLTQWK.LYIISAVVLIMVLIGFIFWRQDDYTSRN....FENKDTALKQSTSE..NNSLSKLEDVQ	72
NTDB id 300 STER RS07485 WP 011681496.1MKEKILAYVKDNR.LF.VSVIAVLMVIFCFLLWMTCGAGNSMEAETSYPDVT.ALSTSSSKQSSQSLS.EASSQ	70
NTDB id 595 KW2 RS08725 WP 021037687.1MEKILDKLLKEYW.KI.MLLVVCALIAGGIFYIFTNSPKPA.....ETLSVE.NLSSSSTK...SSVS.K....	57
NTDB id 1116097 E7X43 RS09315 WP 243114866.1	0
NTDB id 521 SMSK321 RS04960 WP 000443804.1MESIIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPHTPV....KETNLQ.AEVVAISK...DSST.EKDVK	61
NTDB id 493 SM12261 RS04300 WP 000387351.1MEAIIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPQTPV....KETNLQ.AEVVAVSK...DSVS.EKEVK	61
NTDB id 263 KZH43 RS04195 WP 000387330.1MEAIIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPQTPV....KETNLQ.AEVAAVSK...DLVS.EKEVN	61
NTDB id 222 SPD RS04535 WP 000387330.1MEAIIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPQTPV....KETNLQ.AEVAAVSK...DLVS.EKEVN	61
NTDB id 188 SPR RS04295 WP 000387330.1MEAIIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPQTPV....KETNLQ.AEVAAVSK...DLVS.EKEVN	61
NTDB id 153 SP RS04730 WP 000387344.1MEAIIEKIKEYK.II.VICTGLGLLVGGFFLLKPAPQTPV....KETNLQ.AEVAAVSK...DSST.EKEVK	61
consensus		

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NTDB id 105 BSU 25590 NP 390437.1
 NTDB id 1127 NGFG RS10560 WP 003703428.1
 NTDB id 1126 NGFG RS09375 WP 003703428.1
 NTDB id 1125 NGFG RS07215 WP 003703428.1
 NTDB id 1108 NGFG RS06525 WP 003703428.1
 NTDB id 1242 Cj0011c YP 002343483.1
 NTDB id 1382 A4U84 RS07375 WP 062924234.1
 NTDB id 1362 HI 1008 AAC22669.1
 NTDB id 1041 HON27 RS14740 WP 005124713.1
 NTDB id 1069 ABD1 RS03060 WP 043960841.1
 NTDB id 1159 ABDM36 RS04765 WP 001166105.1
 NTDB id 1150 A1552VC RS08580 WP 001166105.1
 NTDB id 1294 VP RS04495 WP 021449594.1
 NTDB id 1395 DSB67 RS04445 WP 010649155.1
 NTDB id 613 LCA RS05340 WP 231952379.1
 NTDB id 1014 ACIAD RS13855 WP 004924483.1
 NTDB id 36 MW RS08230 WP 001793467.1
 NTDB id 20 SA RS08005 WP 001802022.1
 NTDB id 300 STER RS07485 WP 011681496.1
 NTDB id 595 KW2 RS08725 WP 021037687.1
 NTDB id 1116097 E7X43 RS09315 WP 243114866.1
 NTDB id 521 SMSK321 RS04960 WP 000443804.1
 NTDB id 493 SM12261 RS04300 WP 000387351.1
 NTDB id 263 KZH43 RS04195 WP 000387330.1
 NTDB id 222 SPD RS04535 WP 000387330.1
 NTDB id 188 SPR RS04295 WP 000387330.1
 NTDB id 153 SP RS04730 WP 000387344.1
 consensus

KEEKEEELERDLITVDVKGAVKSPGIYDLSV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG.EEAS
 KQEANNDESNETIVIDIKGAVQHHPGVYEMRT.GDRVSQAIEKAGGTSEQADEAQVNLAEITQDGTVVYIPKKGEETA VQQ 128
MSRGGATPY....RFLLRIRHIVARCGLLFATLKGKTMKKMF 37
MSRGGATPY....RFLLRIRHIVARCGLLFATLKGKTMKKMF 37
MSRGGATPY....RFLLRIRHIVARCGLLFATLKGKTMKKMF 37
MSRGGATPY....RFLLRIRHIVARCGLLFATLKGKTMKKMF 37
MFKKLSRYTFSLLFG.LFSATTFAEEKVTT 28
MKTLLFTSVVLCGALVVSSSFAEEKATX 27
AQSFDQNFQEWKAK..QQMYDQKLNITKS..... 55
AQSFDQNFQEWKVK..QQMYDQKLNISKS..... 55
MQIKTKIVT..LFLSLCLPTLPLLAN..AE... 26
MQIKTKIVT..LFLSLCLPTLPLLAN..AE... 26
MK..WITLCLVVEAPLSL..AA... 19
MK..WMTLCLLILAPMSW..AE... 19
 KKTHQAATKKKGGFVDIKGAVVYPIYVSETRLFEVTKNAGGPTDAADTKRLNLAQKLT DQLIYIPKKG..EVVTD 136
TLFSMQIVFAQQFDQYDAWKTQ..QKAHDTRL.....NSS 49
 VKDGDNSKNKGPVYVDVKGAVKHPNVYKMTS.KDRVVDLIDKAQ.LLEDADVSRINLSEKLT DQKMIFIPHKG..QKNVE 148
 VKDGDNSKNKGPVYVDVKGAVKHPNVYKMTS.KDRVVDLIDKAQ.LLDDADVSRINLSEKLT DQKMIFIPHKG..QKNVE 148
 SKTEGSEKVKSKVTVDVKGAVVNPVYTLKA.GARVTDVIQEAAGMTEADAKSVNLAASLSDEEVIYVANKD..ENSV 147
 FNSSSSEKNKNEIMVDLKGAVAKPNVYQISS.DERLVDLIRQAAGGFTDQADQKSLNLSAKLKDEEVIYV PKVG..ESSS 134
 0
 KEEKEEPVEQDLITVDVKGAVKSPGIYDLSV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG..EESAS 138
 KEEKEEPLEQDLITVDVKGAVKSPGIYDLPV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG..EEAAS 138
 KEEKEEPLEQDLITVDVKGAVKSPGIYDLPV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG..EEAVS 138
 KEEKEEPLEQDLITVDVKGAVKSPGIYDLPV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG..EEAVS 138
 KEEKEEPLEQDLITVDVKGAVKSPGIYDLPV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG..EEAVS 138
 KEEKEEPLEQDLITVDVKGAVKSPGIYDLPV.GSRVNDAYQKAGGLTEQADSKSLNLAQKVSD EALVYVPTKG..EEAVS 138

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NTDB id 105 BSU 25590 NP 390437.1	KVSGIGEKSEFEKIKSSITVK.....	205
NTDB id 1127 NGFG RS10560 WP 003703428.1	KVKGIGPAVLAKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1126 NGFG RS09375 WP 003703428.1	KVKGIGPAVLAKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1125 NGFG RS07215 WP 003703428.1	KVKGIGPAVLAKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1108 NGFG RS06525 WP 003703428.1	KVKGIGPAVLAKLKDQASVGAPAPKGPAPVLPVAVKK	131
NTDB id 1242 Cj0011c YP 002343483.1	KVKGIGDKLFEKIKNDITIE.....	79
NTDB id 1382 A4U84 RS07375 WP 062924234.1	EVSGIGKATLEKNRDCIVIE.....	113
NTDB id 1362 HI 1008 AAC22669.1	EVQGIGKATLEKNRDRIF.....	112
NTDB id 1041 HON27 RS14740 WP 005124713.1	NVKGIGPAIFFEKNKSRLAL.....	136
NTDB id 1069 ABD1 RS03060 WP 043960841.1	NVKGIGPAIFFEKNKSRLAL.....	136
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	NVKGIGEATVRNNAARILL.....	103
NTDB id 1150 A1552VC RS08580 WP 001166105.1	NVKGIGEATVRNNAARILL.....	103
NTDB id 1294 VP RS04495 WP 021449594.1	NVKGIGEATIKKNEDRILL.....	94
NTDB id 1395 DSB67 RS04445 WP 010649155.1	KVKGIGEATVRKNEDRILL.....	95
NTDB id 613 LCA RS05340 WP 231952379.1	KVGGIGPKTFEQLQSQICVN.....	235
NTDB id 1014 ACIAD RS13855 WP 004924483.1	KVKGIGPKFLEKNRDRLAT.....	128
NTDB id 36 MW RS08230 WP 001793467.1	KVKGFGSKTFDKLKSYPFTI.....	228
NTDB id 20 SA RS08005 WP 001802022.1	KVKGFGSKTFDKLKSYPFTI.....	228
NTDB id 300 STER RS07485 WP 011681496.1	NVSGIGDKTLDKIRESLYVA.....	231
NTDB id 595 KW2 RS08725 WP 021037687.1	KVSGFGDKTLEKIKDEISID.....	218
NTDB id 1116097 E7X43 RS09315 WP 243114866.1	NTSGFGDKTVEKLNQYVFIK.....	38
NTDB id 521 SMSK321 RS04960 WP 000443804.1	KVSGIGAKTIEKIKDYVTVD.....	216
NTDB id 493 SM12261 RS04300 WP 000387351.1	KVSGIGAKTIEKIKDYVTVD.....	216
NTDB id 263 KZH43 RS04195 WP 000387330.1	KVSGIGGKTIEKIKDYVTVD.....	216
NTDB id 222 SPD RS04535 WP 000387330.1	KVSGIGGKTIEKIKDYVTVD.....	216
NTDB id 188 SPR RS04295 WP 000387330.1	KVSGIGGKTIEKIKDYVTVD.....	216
NTDB id 153 SP RS04730 WP 000387344.1	KVSGIGGKTIEKIKDYVTVD.....	216
consensus	** !*! ***** * *	

non conserved
 similar
 ≥ 50% conserved