

Logo



NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK....KTNFDDITKKYGA ERD KALADALAL IEK DFGKGSIMRLGER.AANQKVSVVSSGSLALDIALGAGGY PKGR	74
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK....PKKLEEISKKFGA ERE KALNDALKL IEK DFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGY PKGR	73
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKK....PKKLEEISKKFGA ERE KALNDALKL IEK DFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGY PKGR	73
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKK....PKKLEEISKKFGA ERE KALNDALKL IEK DFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGY PKGR	73
NTDB id 159 SP RS09750 WP 001085462.1	.MAKK....PKKLEEISKKFGA ERE KALNDALKL IEK DFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGY PKGR	73
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKK....PKKLDEISKKFGA ERE KALNDALKL IEK DFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGY PKGR	73
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKK....PKKLDEISKKFGA ERE KALNDALKL IEK DFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGY PKGR	73
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR....IKKTEEITKKFGD ERR KALDDALKN IEK DFGKGA V MRLGER.AEQKVQVMSSGSLALDIALGAGGY PKGR	73
NTDB id 453 Spy49 1753c ACI62003.1	.MAKK....LKKNEEITKKFGD ERR KALDDALKN IEK DFGKGA V MRLGER.AEQKVQVMSSGSLALDIALGAGGY PKGR	73
NTDB id 625 LCA RS02525 WP 011374196.1MAKD ER QAALDAAL KK IEKN FG KGSIM RM GE K .VDTQV ST VSSGSLALDEALGVGGY PRGR	60
NTDB id 42884 GTCBUS3UF5 RS06910 WP 021322290.1MKR...	3
NTDB id 115 BSU 16940 NP 389576.2MS DR QAALDMAL KQ IEK Q FGKGSIM KL GE K .TDTR IT ST VP SGSLALDIALG IG GY PRGR	58
NTDB id 1313 DR RS12030 WP 010888966.1MSKDATKEISAP TD AK ERS KA IT ET AM S Q IEKA FG KGSIM KL GA E .SKLDV Q V ST GSLSLDIALGVGGY IPRGR	72
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NTDB id 1211 C694 RS00795 WP 000963128.1MAID ED K Q KA IS LA IK Q ID K V FGK GA L V RL GDK .QVE K IDS IS TGS LGL DLALG IG GV PKGR	61
NTDB id 1214 NCTC11637 00359 SQJ03206.1MAID ED K Q KA IS LA IK Q ID K V FGK GA L V RL GDK .QVE K IDA IS TGS LGL DLALG IG GV PKGR	61
NTDB id 1334 RAOC RS04870 WP 004918236.1MAKT..ETTSPND DK KA AA V LEK LD KT Y G K G T VM KL G D GT VD HN IE VI PS GS L GL DLALG V GGY PKGR	68
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENK KSS KN NTV ..VK Q VE PE Q KK Q A LE AA LA Q IEK Q FGKGSIM KL G QT .EQLD IE SV ST GS LGL DLALG IG GL PMGR	76
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NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....K..KAAS MSA EK Q K AL AA LA AA LA Q IEK Q FGKGSIM KM G DA .EVE P V Q V ST GS LGL D V ALG V G GL PR GR	68
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NTDB id 1142 GTF74 RS11550 WP 000344154.1MDEN KQ KA AA LA AA L GQ IEK Q FGKGSIM RL G DN .RAMD VET IS TGS LSLDIALGAG GL PM GR	59
NTDB id 1205 PSJM300 05545 AFN77183.1MDEN KK R AL AA AA L GQ IEK Q FGK GA V MR M GDH .DR QA IP SI STGS LGL DIALG IG GL PKGR	59
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDEN KS K AL QA AA LS Q IEK Q FGK NT V M RL GDN .TV Q AVE AV STGS LTL DIALG IG GL PKGR	59
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consensus

- X non conserved
- X similar
- X ≥ 50% conserved