

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVDDDKISEQDALHSA	DSANDLRLMLKTKR.GDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVYVQGEITYEDALLHADSSNDLRLMIKLG	S.ETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSE	.EANKPLDNRERMSEMER.....KMTFDGQRRNLK....	383
NTDB id 1171373 FDG16 RS03955 WP 001055649.1	TFAQSAQWRQAQCRL		326
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHLITVDKGLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAA	394
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLIAPPEVARRRAQNSESMSF		347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLIAPPEVARRRAQNSESMSF		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARCVVDAQEVQSKIELDLKAF		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLIARCVVDAQEVQSKIELDLKAF		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNK.SLF		344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNK.SLF		344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDQHLTELVNKNIISKHTAHEAALNK.SLF		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDQSLKGLLAKGLISRENAREKAKIP.ENF		344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKGLLAKGLISRENAREKAKIP.ENF		344
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQP.ESFL		345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQP.ESFL		345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVARGVISPQTARTAAKQP.ESFL		345
consensus	!*** * * * **** * * *		

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	APATPAAPAAAGRSEPGRAGGFGRR	368
NTDB id 1203 PSJM300 01605 AFN76401.1	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
NTDB id 1171373 FDG16 RS03955 WP 001055649.1	326
NTDB id 1307 DR RS10055 WP 010888596.1	APATPAAPAAAGRSEPGRAGGFGRR	418
NTDB id 1086 NMV RS00245 WP 002218487.1	347
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NTDB id 1185 GTF74 RS11940 WP 000350195.1	345
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1	345
consensus		

non conserved
 similar
 ≥ 50% conserved