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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKIS	EQDALHSA	DSANDLRLMLKTKRGDDY...	GSGSLQNVKI.....	DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEI	TYEDALLHA	DSSNDLRLMIKLGSETDGEHLTSVS.	QGLTL.....	EMSDDDPGRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHS	YADALKHAD	SPNNLRLQIKLSEEANKPLDNRERMSEMER.....	KMTFDGQRRLK.....		383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFR	RHLITYDK	GLERA	IDPKEFARLANDASAGL....	GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLI	APEVARRRA	QNSESMSF.....			347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLI	APEAARRRA	QNSESMSF.....			347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLI	ARGVVDAQEV	QSKIELDLKAF.....			345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLI	ARGVVDAQEV	QSKIELDLKAF.....			345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTEL	VNKNIISKHTA	HEAALNKSLF.....			344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTEL	VNKNIISKHTA	HEAALNKSLF.....			344
NTDB id 27175 SPUTW3181 RS06885 WP 011788674.1	TLEQCLQNLVNRGLI	TREDAMSKS	SNKQATF.....			345
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLIS	RENAREKAK	KIPENF.....			344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLLAKGLIS	RDSAKEKAK	KQPENF.....			344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLLAKGI	INLPTARAVAK	SPESEF.....			344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVIS	PQTARTAAK	QPESFL.....			345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTAAK	QPESFL.....			345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTAAK	QPESFL.....			345
consensus	!*** * !* **** * *					



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NTDB id 1203 PSJM300 01605 AFN76401.1	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
NTDB id 1307 DR RS10055 WP 010888596.1	PATPAAPAAGRSEPGRAGGFGRR	418
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consensus		



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