

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKTI	SEQDALHSADSANDLRLMLKTKRGDDY...	GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQCEIT	YEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....	EMSDDDPGRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHIS	YADALKHADSPNNLRLQIKLSEEANKPLDNRRERMSEMER.....	KMTFDGQRRNLK.....	383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHILIT	YDKGLERAI.DPKE..FARLANDASAGL.GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA		395
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLIA	PEVARRRAQNSESM..SF.....		347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLIA	PEAARRRAQNSESM..SF.....		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARGVVD	DAQEVQSKIELDLKA..F.....		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLIARGVVD	DAQEVQSKIELDLKA..F.....		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISK	HTAHEAAL.NKSL..F.....		344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	TLDQHLTELVNKNIISK	HTAHEAAL.NKSL..F.....		344
NTDB id 1101885 Q9286 RS06895 WP 011258220.1	TLDQHLQDLVKRSLIT	RNQAREYAK.DKRI..FE.....		345
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCCLKGLIAKGLIS	RDSAKEKAK.QPEN..F.....		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLIS	RENAREKAK.IPEN..F.....		344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLIAKGIINL	PTARAVAK.SPES..F.....		344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVISP	QTARTAAK.QPES..FL.....		345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTAAK.QPES..FL.....		345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTAAK.QPES..FL.....		345
consensus	!*** * !* **** * *			

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	PATPAAPAAGRSEPGRAGGFRR	368
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	PATPAAPAAGRSEPGRAGGFRR	418
NTDB id 1086 NMV RS00245 WP 002218487.1		347
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consensus		

- ⓧ non conserved
- ✖ similar
- ⓧ ≥ 50% conserved