

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKIS	EQDALHSA	DSANDLRLMLKTKRGDDY...	GSGSLQNVKI.....	DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEIT	YEDALLHA	DSSNDLRLMIKLGSETDGEHLTSVS.	QGLTL.....	EMSDDDPGRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHSYAD	ALKHADSPNNLRLQIKL	SEEANKPLDNRRERMSEMER.....	KMTFDGQRRLK.....		383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRBHLIT	YDKGLERAI	.DPKE..	FARLANDASAGL	.GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLI	APEVARRRA	QNSESM..	SF.....		347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLI	APEAARRRA	QNSESM..	SF.....		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARGVV	DAQEVQSKI	ELDLKA..	F.....		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLIARGVV	DAQEVQSKI	ELDLKA..	F.....		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIIS	SKHTAHEAAL	.NKSL..	F.....		344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIIS	SKHTAHEAAL	.NKSL..	F.....		344
NTDB id 1146425 NBC2815 RS13950 WP 002802057.1	TLDQHLQDLIKRSLI	TRNQAREYAK	.DKRI..	FE.....		345
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLLAKGLIS	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	TLDQSLKQLLAKGIIN	LPTARAVAK	.SPES..	F.....		344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVIS	PQTARTA	AAK.QPES..	FL.....		345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTA	AAK.QPES..	FL.....		345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTA	AAK.QPES..	FL.....		345
consensus	!*** * !* **** * *			*		



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NTDB id 1307 DR RS10055 WP 010888596.1	PATPAAPAAGRSEPGRAGGFRR	418
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consensus		



non conserved
 similar
 ≥ 50% conserved