

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	ALKHADSPNNLRLQIKLSEEANKPLDN	RERMSEMER.....	KMTFDGQRRLK.....		383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFR	RHLITYDKGLERA	IDPKEFARLANDASAGL...	GGASMTPAAPPVPAYNSASSSSTGRGGDF	FGRGAAA	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	ARGVVD	DAQEVQSKIEL..	DLKAF.....		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLI	ARGVVD	DAQEVQSKIEL..	DLKAF.....		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTEL	VNKNII	SKHTAHEAALNKSLF.....			344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTEL	VNKNII	SKHTAHEAALNKSLF.....			344
NTDB id 1155751 ALTBGP9 RS15130 WP 156078754.1	TMDQCLQRLV	AMGVITKQDA	AAKSF	FDKQSMNSF.....		347
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLI	AKGLISR	DSAKEKAK	KQPNF.....		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLV	AKGLISR	ENAREKAK	KIPNF.....		344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLI	AKGIINLPT	ARAVAK	SPEF.....		344
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLV	ARGVIS	PQTARTAA	KQPESFL.....		345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLV	ARGVIS	PQTARTAA	KQPESFL.....		345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLV	SRGVIS	PQTARTAA	KQPESFL.....		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	

non conserved
 similar
 ≥ 50% conserved