

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVDDDKISEQDALHSA	DSANDLRLMLKTKRGDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQCEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.....	381	
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPLDNRRERMSEMER.....KMTFDGQRRNLK.....	383	
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHLITTYDKGLERAI	IDPKEFARLANDASAGL...GGASMPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLI	APEVARRRAQNSSESMSF.....	347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLI	APEAARRRAQNSSESMSF.....	347
NTDB id 1054801 BCU15 RS13140 WP 010435660.1	TMEQNAKQLIAQCLV	DSDEVEKKIEIQASMF.....	345
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARCVV	DAQEVQSKIELDLKAF.....	345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLIARCVV	DAQEVQSKIELDLKAF.....	345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAA	LNKSLF.....	344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAA	LNKSLF.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCCLKGLIAKGLISR	DSAKEKAKQPENF.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLIAKGLISRENAREKAKIPENF.....	344	
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLIAKGIINLPTARAVAKSPESF.....	344	
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTAAKQPESFL.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTAAKQPESFL.....	345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVIS	PQTARTAAKQPESFL.....	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	PATPAAPAAGRSEPGRAGGFGR	418
NTDB id 1086 NMV RS00245 WP 002218487.1	347
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consensus	

non conserved
 similar
 ≥ 50% conserved