

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	TFDQALFDLYDHQHSYADALKHADSPNNLRLQIKLSEEANKPLDNRRERMSEMER.....KMTFDGQRRNLK.....	383		
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHLITYDKGLERAI	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 214581 FORC53 RS02350 WP 026050359.1	TMEQNAARQLMAQCKV	AKEEVDAKIEIETIQF.....	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	TLDSCLKGLIAKGLISRDS	AKEKAKQPENF.....	344	
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLISRENAREKA	KIPENF.....	344	
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLIAKGIINLPT	ARAVAKSPESF.....	344	
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTA	AKQPESFL.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVIS	PQTARTA	AKQPESFL.....	345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVIS	PQTARTA	AKQPESFL.....	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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NTDB id 214581 FORC53 RS02350 WP 026050359.1	345
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consensus		

non conserved
 similar
 ≥ 50% conserved