





logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVDDKISEQDALHSA	DSANDLRLMLKTKRGDDYGGSLQ	...NVKIDME	368
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHISYADALKHA	ADSPNNLRLQIKLSEEANKPLDNRERMSEMERKMTFD	.....GQRRNLK	383
NTDB id 1146834 CJZ87 RS03880 WP 001055657.1	TFAQSAQWRQAQGR			326
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHLITYDKGLERA	ID.PKEFARL	..ANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAP	396
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLIAPEVARRRA	QNSESMSF		347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLIAPEAARRRA	QNSESMSF		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARGVVD	DAQEVQSKIELDLKAF		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLIARGVVD	DAQEVQSKIELDLKAF		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALN	.KSLF		344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALN	.KSLF		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TIDMCLKGLVAKGLISRENAREKAKI	.PENF		344
NTDB id 1202 PSJM300 01600 AFN76400.1	TIDSCLKGLIAKGLISRDSAKEKAKQ	.PENF		344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLLAKGILNLP	TARAVAKS.PESF		344
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQ	.PESFL		345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQ	.PESFL		345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSGVISPQTARTAAKQ	.PESFL		345
consensus	!*** * * * **** * *			

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NTDB id 1019 ACIAD RS04205 WP 004922051.1	.....	383
NTDB id 1146834 CJZ87 RS03880 WP 001055657.1	.....	326
NTDB id 1307 DR RS10055 WP 010888596.1	ATPAAPAAGRSEPGRAGGFGR	418
NTDB id 1086 NMV RS00245 WP 002218487.1	.....	347
NTDB id 1116 NGFG RS10615 WP 003690200.1	.....	347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	.....	345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	.....	345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	.....	344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	.....	344
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consensus		

non conserved  
 similar  
 ≥ 50% conserved