

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	QTFDQALYQLVDDKISEQDALHSA	ADSANDLRLMLKTKRGDDYGGSLQ	...NVKIDME	368
NTDB id 1019 ACIAD RS04205 WP 004922051.1	QTFDQALFDLYDHQHISYADAL	KHADSPNNLRLQIKLSEEANKPLD	NRERMSEMERKMTFD	383
NTDB id 1145197 DXE35 RS01910 WP 114689377.1	ITMEASLQQLEQAGQIDPL			333
NTDB id 1307 DR RS10055 WP 010888596.1	VTMDAYLAGLFRHLITTYDK	CLERAIDPKEFARLAND	...ASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1086 NMV RS00245 WP 002218487.1	QTMDQSLQSLVRQGLIAPEVARRRA	QNSEMSF		347
NTDB id 1116 NGFG RS10615 WP 003690200.1	QTMDQSLQSLVRQGLIAPEAARRRA	QNSEMSF		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	QTMEQNAKQLIARGVVD	DAQEVQSKIELDLKAF		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	QTMEQNAKQLIARGVVD	DAQEVQSKIELDLKAF		345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	QTLDQHLTELVNKNIISKHTAHEAALNK	SLF		344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	QTLDQHLTELVNKNIISKHTAHEAALNK	SLF		344
NTDB id 1202 PSJM300 01600 AFN76400.1	QTLDQSLKGLIAKGLISRDSA	KEKAKQP.ENF		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	QTLDQSLKGLIAKGLISRDSA	KEKAKQP.ENF		344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TTLDDQSLKQLIAKGLINLPTARAVAKSP	ESF		344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TTLDDQSLKGLVSRGVISPQTARTAAKQP	ESFL		345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TTLDDQSLKGLVSRGVISPQTARTAAKQP	ESFL		345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TTLDDQSLKGLVSRGVISPQTARTAAKQP	ESFL		345
consensus	*!*** * !* ***** * *			

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NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
NTDB id 1145197 DXE35 RS01910 WP 114689377.1	333
NTDB id 1307 DR RS10055 WP 010888596.1	PATPAAPAAGRSEPGRAGGFGR	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
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NTDB id 1202 PSJM300 01600 AFN76400.1	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	344
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NTDB id 1040 HON27 RS13360 WP 000355489.1	345
consensus	

non conserved
 similar
 ≥ 50% conserved