

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKISEQDALHSA	DSANDLRLMLKTKRGDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEITYEDALLHA	DSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHSYADALKHAD	SPNNLRLQIKLSEEANKPLDNRRERMSEMER.....KMTFDGQRRLK.....	383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRRHLYTYDKGLERA	ID.PKEFARLAND..ASAGL.GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLIAPAAARRRA	QNSEMSF.....	347
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLIAPVARRRA	QNSEMSF.....	347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARGVVDAQEVQSKI	ELDLKAF.....	345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLIARGVVDAQEVQSKI	ELDLKAF.....	345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEA	ALN.KSLF.....	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEA	ALN.KSLF.....	344
NTDB id 160918 Cater282 RS03365 WP 061532286.1	TLDSNLTTELVKRNVISMAAARA	AKS.PDNFPG.....	347
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLIAKGLISRDSA	KEKAKQ.PENF.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMLKGLVAKGLISRENA	REKAKI.PENF.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLIAKGIINLPTA	RAVAKS.PESF.....	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVISPQT	ARTAAKQ.PESFL.....	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVISPQT	ARTAAKQ.PESFL.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQT	ARTAAKQ.PESFL.....	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	PATPAAPAAGRSEPGRAGGFRR	418
NTDB id 1116 NGFG RS10615 WP 003690200.1	347
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consensus	

non conserved
 similar
 ≥ 50% conserved