

logo

NTDB id 1177	A1552VC	RS01165	WP 000422572.1	TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203	PSJM300	01605	AFN76401.1	TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.....	381
NTDB id 1019	ACIAD	RS04205	WP 004922051.1	TFDQALFDLYDHQHSYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSEMER.....KMTFDGQRRNLK.....	383
NTDB id 288180	DBN47	RS12805	WP 003386380.1	TMDMSLAELYKREITISYDTALNYSVD.KDLITKIINL.....	350
NTDB id 1307	DR	RS10055	WP 010888596.1	TMDAYLAGLFRHLLITYDKGLERAID.PKEFARLAND..ASAGL.GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1116	NGFG	RS10615	WP 003690200.1	TMDQSLQSLVVRQGLIAPAAARRRAQNSESMSF.....	347
NTDB id 1086	NMV	RS00245	WP 002218487.1	TMDQSLQSLVVRQGLIAPAVARRRAQNSESMSF.....	347
NTDB id 1185	GTF74	RS11940	WP 000350195.1	TMEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345
NTDB id 1176	A1552VC	RS01160	WP 000350195.1	TMEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	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	TLDMCLKGLVAKGLISRENAREKAKI.PENF.....	344
NTDB id 1202	PSJM300	01600	AFN76400.1	TLDSCLKGLLAKGLISRDSAKEKAKQ.PENF.....	344
NTDB id 1020	ACIAD	RS04210	WP 011182164.1	TLDQSLKQLLAKGIINLPTARAVAKS.PESF.....	344
NTDB id 1081	FDQ49	RS05335	WP 006581387.1	TLDQSLKGLVSRGVISPQTARTAAKQ.PESFL.....	345
NTDB id 1060	ABD1	RS04380	WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQ.PESFL.....	345
NTDB id 1040	HON27	RS13360	WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQ.PESFL.....	345
consensus				!*** * !* ***** * *	

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NTDB id 1307	DR	RS10055	WP 010888596.1	PATPAAPAAGRSEPRAGGFGR	418
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NTDB id 1086	NMV	RS00245	WP 002218487.1	347
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consensus				

- X non conserved
- X similar
- X ≥ 50% conserved