

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	QTFDQALYQLVDDKISEQDALHSA	DSANDLRLMLKTKR.GDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	QTFDQALYNLYVQGEITYEDALLHA	DSSNDLRLMIKLGSE.ETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR...	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	QTFDQALFDLYDHQHISYADALKHA	DSPNNLRLQIKLSE.EANKPLDNRERMSEMER.....KMTFDGQRRNLK...	383
NTDB id 107286 AACH44 RS16435 WP 338659328.1	QTFEQSYQQRCRDCLISHSCALAV.....	338
NTDB id 1307 DR RS10055 WP 010888596.1	VTMDAYLAGLFRHLITYDKGLERA	IDPKEFARLANDASAGLGG.....ASMPAAPPVPAYNSASSSSTGRGGDFGRGA	393
NTDB id 1086 NMV RS00245 WP 002218487.1	QTMDQSLQSLVRQCLIAPEVARRRA	QNSEMSF.....	347
NTDB id 1116 NGFG RS10615 WP 003690200.1	QTMDQSLQSLVRQCLIAPEAARRRA	QNSEMSF.....	347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	QTMEQNAKQLIARCVVDAQEVQSKI	ELDLKAF.....	345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	QTMEQNAKQLIARCVVDAQEVQSKI	ELDLKAF.....	345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	QTLDQHLTELVNKNIISKHTAHEEA	LNK.SLF.....	344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	QTLDQHLTELVNKNIISKHTAHEEA	LNK.SLF.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	QTLDQSLKGLLAKGLISRDSA	KEKAKQP.ENF.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	QTLDQSLKGLLAKGLISRRENARE	KAKIP.ENF.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TTLDDQSLKQLLAKGIINLPTARA	VAKSP.ESF.....	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TTLDDQSLKGLVSRGVISPQTAR	TAAKQP.ESFL.....	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TTLDDQSLKGLVARGVISPQTAR	TAAKQP.ESFL.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TTLDDQSLKGLVARGVISPQTAR	TAAKQP.ESFL.....	345
consensus	*!*** * * * * * * * *		

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NTDB id 1307 DR RS10055 WP 010888596.1	AAPATPAAPAAAGRSEPGRAGGFGRR	418
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NTDB id 1116 NGFG RS10615 WP 003690200.1	347
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consensus	

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