

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	VGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	368
NTDB id 25790 RMET RS14710 WP 029307174.1	QGMISFDQALFELYETEKITYEDALKNADSLNDLRLMIKLHSKRHRDNDLGAGTEHLNV.....I.....	379
NTDB id 1203 PSJM300 01605 AFN76401.1	LGMQTFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	LGMQTFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSEMER.....KMTFDGQRRNLK.	383
NTDB id 1307 DR RS10055 WP 010888596.1	EGMVTMDAYLAGLFRHLITYDKGLERAIDP.KEFA..RLANDASAGLG.GASMTPAAPPVPAYNSSASSSTGRGGDFGR	391
NTDB id 1185 GTF74 RS11940 WP 000350195.1	HGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	HGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	KGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	KGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	LGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	LGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	HGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	HGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	HGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	HGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVKI.....DME.....	345
consensus	!!***** * !* ***** * * * * * * *	



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NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
NTDB id 1307 DR RS10055 WP 010888596.1	GAAAPATPAAPAAGRSEPGRAGGFGR	418
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consensus	

CAAAAPATPAAPAAGRSEPGRAGGFGR

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