

logo

NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHILITYDKCLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSAS	398
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVLRQGLIAPEAARRRAQNSESMF	347
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVLRQGLIAPEVARRRAQNSESMF	347
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLISRENAREKAKIPENF	344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLAKGLISRDSAKEKAKQPENF	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLIAKGIINLPTARAVAKSPESF	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVISPQTARTAAKQPESFL	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQPESFL	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQPESFL	345
NTDB id 669225 IEQ11 RS20200 WP 036104874.1	SFDQCLFRLYKEGRIEMEEALKAADSRDGLALKFRLSEGGETGEH	371
NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDY	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEH	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPL	383
consensus	*!*!* ! !* ***!* * ! * *	

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NTDB id 1307 DR RS10055 WP 010888596.1	PAAPAAGRSEPGRAGGFRR	418
NTDB id 1116 NGFG RS10615 WP 003690200.1	347
NTDB id 1086 NMV RS00245 WP 002218487.1	347
NTDB id 1283 ACE17W RS02630 WP 010947729.1	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	344
NTDB id 1202 PSJM300 01600 AFN76400.1	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	345
NTDB id 669225 IEQ11 RS20200 WP 036104874.1	371
NTDB id 1177 A1552VC RS01165 WP 000422572.1	368
NTDB id 1203 PSJM300 01605 AFN76401.1	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	383
consensus		

non conserved
 similar
 ≥ 50% conserved