

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TIDRVIDVFPAEEKMVRSMSESQAIVISQTLKKNGGGRVAAHEIMIGTPAIRNLIENKYAQMYSAIQTGQALGMQ	313
NTDB id 1203 PSJM300 01605 AFN76401.1	ALERTIHLPKEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLINSPTIADLIRRGELHELKATMARSQEVGMQ	315
NTDB id 1039366 ACDI99 RS11870 WP 005018931.1	ALDRIINFFPADRQNNQVWMDLSLNLKATVAQQLVPTPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ	316
NTDB id 1019 ACIAD RS04205 WP 004922051.1	ALDRIVHFFEADRHSQLYMDLSLNLKAVIAQQLIPTLDGNGRRAAIEVLINTPLIADYIRKGEVHEIKDLMKRSRELGMQ	316
NTDB id 1307 DR RS10055 WP 010888596.1	TLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDGQSRRAAEVLINTPLLSDYIRQGEITYKIKELMARSQELGMQ	319
NTDB id 1116 NGFG RS10615 WP 003690200.1	SIDRIVDVFPPEEQEQEIRVQLANNLVAVMTQQLLRADGQGRVLAYEILLANPAVRALIREGKTFTQITSVMQTGAREGMV	315
NTDB id 1086 NMV RS00245 WP 002218487.1	TVDRIVDVFPAGEKEMVRSMLSESLTAVISQNLKTHDGNNGRVASHEILIANPAVRNLIRENKITQINSVLQTGGQASGMQ	315
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TVDRIVDVFPAGEKEMVRSMLSESLTAVISQNLKTHDGNNGRVASHEILIANPAVRNLIRENKITQINSVLQTGGQASGMQ	314
NTDB id 1259 GC085 RS10305 WP 010947729.1	TINRIIDVFPAEEKSMVRSMLSESLQAVVAQSLLKK.NKGGRVAAAEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGMQ	314
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TINRIIDVFPAEEKSMVRSMLSESLQAVVAQSLLKK.NKGGRVAAAEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGMQ	314
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TIDRVIDVFPAEEKDMVRAMLSESLQAVISQTLKK.NKGGRVAAAEIMIGIPAIRNLIRENKVAQMYSAIQTGANHGMT	314
NTDB id 1040 HON27 RS13360 WP 000355489.1	TIDRVIDVFPAEEKDMVRAMLSESLQAVISQTLKK.NKGGRVAAAEIMIGIPAIRNLIRENKVAQMYSAIQTGANHGMT	314
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TIDRVIDVFPAEEKDMVRAMLSESLQAVISQTLKK.NKGGRVAAAEIMIGIPAIRNLIRENKVAQMYSAIQTGANHGMT	314
NTDB id 1202 PSJM300 01600 AFN76400.1	TIDRVVDVFPAEEKAMVRSMLSESLQSVISQTLTKK.IGGGRVAAAEIMIGTPAIRNLIREDKVAQMYSAIQTGGSLSGMQ	314
consensus	***!***** ** *!***! *****! **** * *!***! *****!*****! * ** * * *****! ! *	

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKISEQDALHSSANDLRLMLKTKRGDDYGS.G...SLQNV...KIDME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEH.LTSVSQ.GL...TLEMSDDDPGRSFR.....	381
NTDB id 1039366 ACDI99 RS11870 WP 005018931.1	TFDQALYDLYKAGQISYKDALKHADAPNDLRLQIKLSEEGASQL.M.....AA...GRNITFDGQ.....	372
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPL.DNRERMSEM...ERKMTFDGQRRNLK.....	383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHLITYDKGLERAIDPKEFARLANDASAGLGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAPATP	399
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLIAPEAARRRAQNSSEMSF.....	347
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLIAPEVARRRRAQNSSEMSF.....	347
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF.....	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF.....	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 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLISRENAREKAKIPENF.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCCLKGLIAKGLISRDSAKEKAKQIPENF.....	344
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NTDB id 1177 A1552VC RS01165 WP 000422572.1		368
NTDB id 1203 PSJM300 01605 AFN76401.1		381
NTDB id 1039366 ACDI99 RS11870 WP 005018931.1		372
NTDB id 1019 ACIAD RS04205 WP 004922051.1		383
NTDB id 1307 DR RS10055 WP 010888596.1	AAPAAGRSEPGRAGGFGRR	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 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 1194 PAKAF RS01995 WP 003084552.1		344
NTDB id 1202 PSJM300 01600 AFN76400.1		344
consensus		

 non conserved
 similar
 $\geq 50\%$ conserved