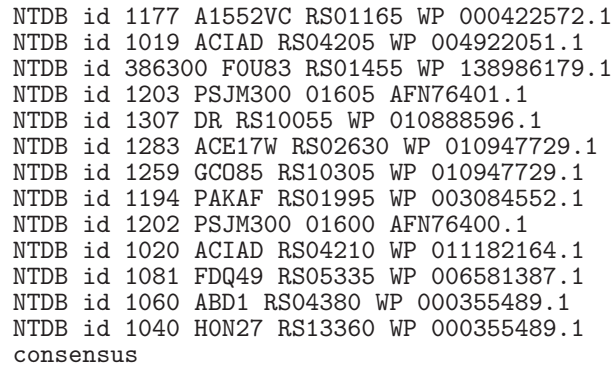
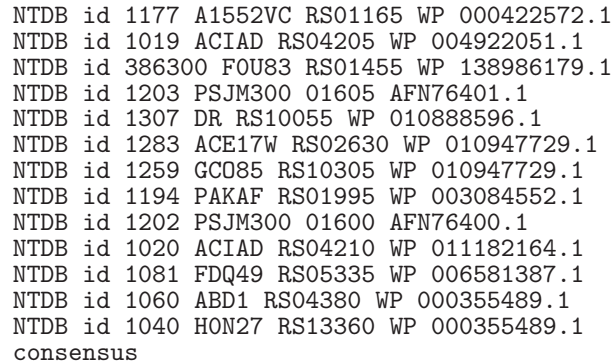


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NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 386300 FOU83 RS01455 WP 138986179.1
NTDB id 1203 PSJM300 01605 AFN76401.1
NTDB id 1307 DR RS10055 WP 010888596.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GCO85 RS10305 WP 010947729.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1202 PSJM300 01600 AFN76400.1
NTDB id 1020 ACIAD RS04210 WP 011182164.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HON27 RS13360 WP 000355489.1
consensus

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	ALERIILHIVPKEEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLINSPRIADLIRRGELHELKATMARSQEVGMQ	313
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDGQSRRAAVEVLINTPLLSDIYRQGEIYKIKELMARSQELGMQ	316
NTDB id 386300 FOU83 RS01455 WP 138986179.1	ALDRIVSFFPPEHHDLWMELSLNLKAIVAQQLIPTKDGKGRRAAIEVMICTPLIQDMIRKGEVHEIKDVMKKSTNLGMQ	315
NTDB id 1203 PSJM300 01605 AFN76401.1	ALDRIINFFPADRQNVWMDLSLNLKAIVAQQLVPTPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLMQ	315
NTDB id 1307 DR RS10055 WP 010888596.1	SIDRIVDVFPPEEQQEQRVQLANNLVAVMTQQLPRADGQGRVLAYEILANPAVRALIREGKTFQITSVMTGAREGMV	319
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NTDB id 1040 HON27 RS13360 WP 000355489.1	TIDRVIDVFPAAEKDMVRAMLSESLQAVISQTLIK.KNKGGRVAAHEIMIGIPAIRNLIREDKVAQMYSIAITGANHGMT	314
consensus	***!***** ** *!***!*****!*** * ***!*** !*** *****!*** ***** ***** !!*	

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVDDKISEQDALHSADSANDLRLMLKTRGDDYDGS...S...LQNVK.IDME.....	368
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NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELNVKNKIISKHTAHEAALNKSLF.....	344
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NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLLAKGLISRDSAKEKAKQPENF.....	344
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NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQPESFL.....	345
consensus	!*** ! !* ***** ** * !* * *	

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	ATPAAPAAAGRSEPCGRAGGFCRR	368
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NTDB id 1040 HON27 RS13360 WP 000355489.1		345
consensus		

 non conserved
 similar
 $\geq 50\%$ conserved