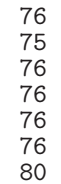
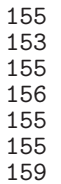


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NTDB id 1203 PSJM300 01605 AFN76401.1
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NTDB id 1307 DR RS10055 WP 010888596.1
consensus
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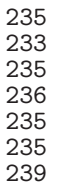
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NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1307 DR RS10055 WP 010888596.1
consensus
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NTDB id 397314 N028 RS14285 WP 003400837.1
NTDB id 1177 A1552VC RS01165 WP 000422572.
NTDB id 1203 PSJM300 01605 AFN76401.1
NTDB id 1019 ACIAD RS04205 WP 004922051.1
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NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1307 DR RS10055 WP 01088596.1
consensus

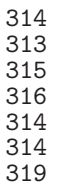
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NTDB id 397314 N028 RS14285 WP 003400837.1
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NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 1202 PSJM300 01600 AFN76400.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1307 DR RS10055 WP 010885596.1
consensus

```



logo

NTDB id 397314 N028 RS14285 WP 003400837.1	TFDSALFNLAVEGAISEEEALKNADSQNNVRLRLKLHSEGGAG.SLTT.PPPAPTGSSTASTAEWGLVDDDA.....	384
NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDDKISEQDALHSADSANDLRLMLKTKRGDDY....GS.GS.....LQNVKIDM.E.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGE.HLTS.VS.....QGLTLEM.SDDD.....	375
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKP.LDNR.ER.....MSEMERKM.TFDG.....	377
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCLKGLLAKGLISRDSAKEKAKQPENF.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLISRENAREKAKIPENF.....	344
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRRLITIDKGLERAIDPKEFARLANDASAGLGASMTPAAPPVPAYNSASS.SSTGRGGDFGRGAAAPAT	398
consensus	! * ! * ! * ! * * * * * ! * * * * *	

logo

NTDB id 397314 N028 RS14285 WP 003400837.1	PGPQA.....	389
NTDB id 1177 A1552VC RS01165 WP 000422572.1		368
NTDB id 1203 PSJM300 01605 AFN76401.1	PGRSFR....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	QRRNLK....	383
NTDB id 1202 PSJM300 01600 AFN76400.1		344
NTDB id 1194 PAKAF RS01995 WP 003084552.1		344
NTDB id 1307 DR RS10055 WP 010888596.1	PAAPAAGRSEPGRAGGFGRR	418
consensus		

- non conserved
- similar
- ≥ 50% conserved