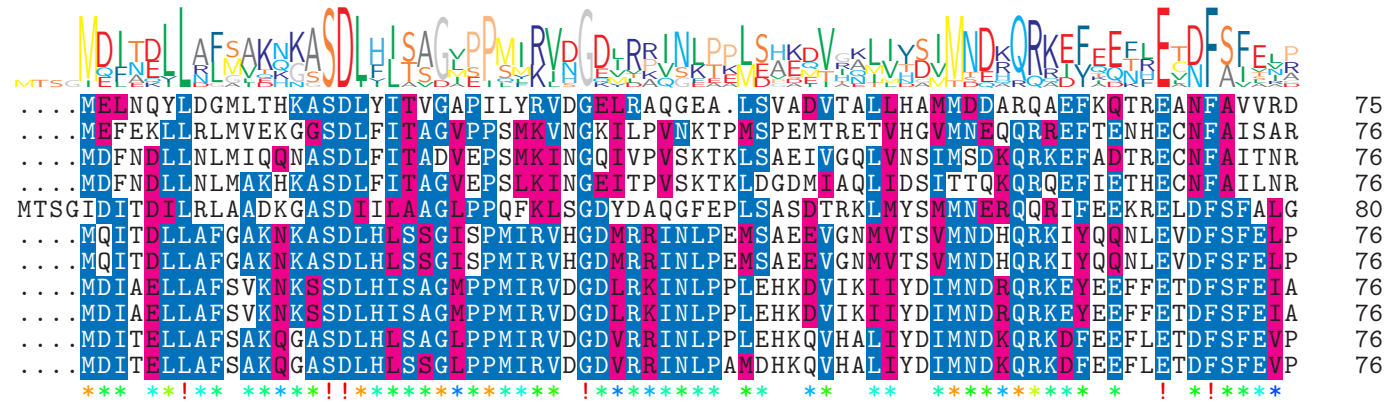


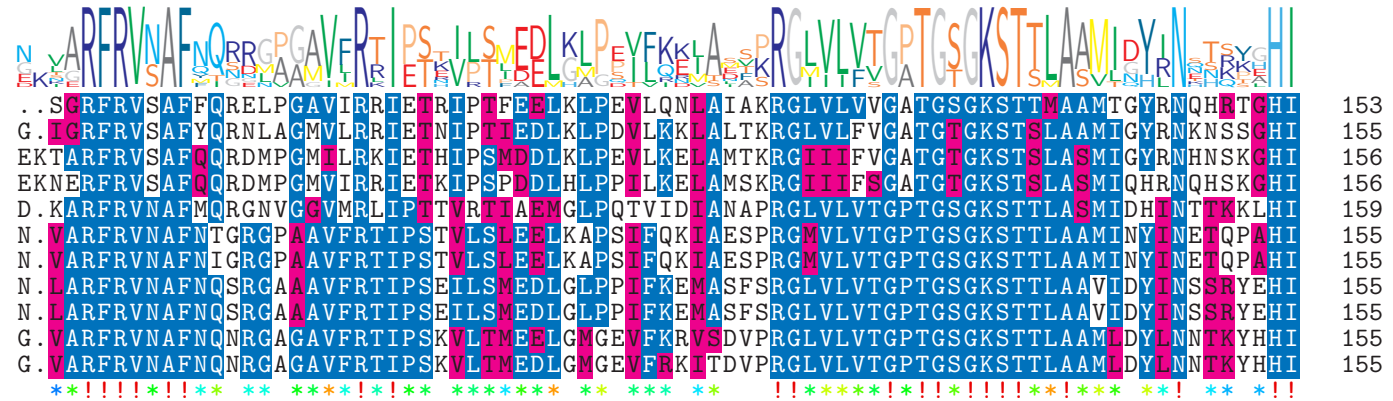
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NTDB id 1203 PSJM300 01605 AFN76401.1
NTDB id 370686 AhaeAN54 RS13375 WP 005081584.1
NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 1307 DR RS10055 WP 010888596.1
NTDB id 1116 NGFG RS10615 WP 003690200.1
NTDB id 1086 NMV RS00245 WP 002218487.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GC085 RS10305 WP 010947729.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1202 PSJM300 01600 AFN76400.1
consensus
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NTDB id 1177 A1552VC RS01165 WP 000422572.1
NTDB id 1203 PSJM300 01605 AFN76401.1
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NTDB id 1019 ACIAD RS04205 WP 004922051.1
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NTDB id 1202 PSJM300 01600 AFN76400.1
consensus

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consensus

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logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TIDRILYDVFPAAEEKSMVRSMLSESRAVISQQLKTDGGRVAAHEILLNTPAURNLRECKLAQIYSSNOTGQAL	313
NTDB id 1203 PSJM300 01605 AFN76401.1	ALERILHIVPKEQREQFLLDLSLNLKGVIAQQLLRDKNGKGRHGVFEVLLNSPRIADLIRRGELHEIKATMARSQEVGMQ	315
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NTDB id 1202 PSJM300 01600 AFN76400.1	TIDRVVDVFPAAEEKAMVRSMLSESLQSVISQTLTKKI.GGGRVAAHEIMIGTPAIRNLIREDKVAQMYSAITQTGGSILGMQ	314
consensus	TIDRVVDVFPAAEEKSMVRSMLSESLQAVISQTLTKKI.GGGRVAAHEIMIGTPAIRNLIREDKVAQMYSSITQTGGSILGMQ	314

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVDDKISEQDALHSAADSANDIRMLKTKRGDDYGS...GSLQNV....KIDME.....	368
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NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIIISKHTAHEAALNKSLF.....	344
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NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCCLKGLIAKGLISRDSAKEKAKQPENF.....	344
consensus	!!*! !* **!* * * ! *	

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	AAPAAGRSEPCGRAGGFCRR	368
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NTDB id 1307 DR RS10055 WP 010888596.1	AAPAAGRSEPGRAGGFGR	418
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NTDB id 1194 PAKAF RS01995 WP 003084552.1		344
NTDB id 1202 PSJM300 01600 AFN76400.1		344
consensus		

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved