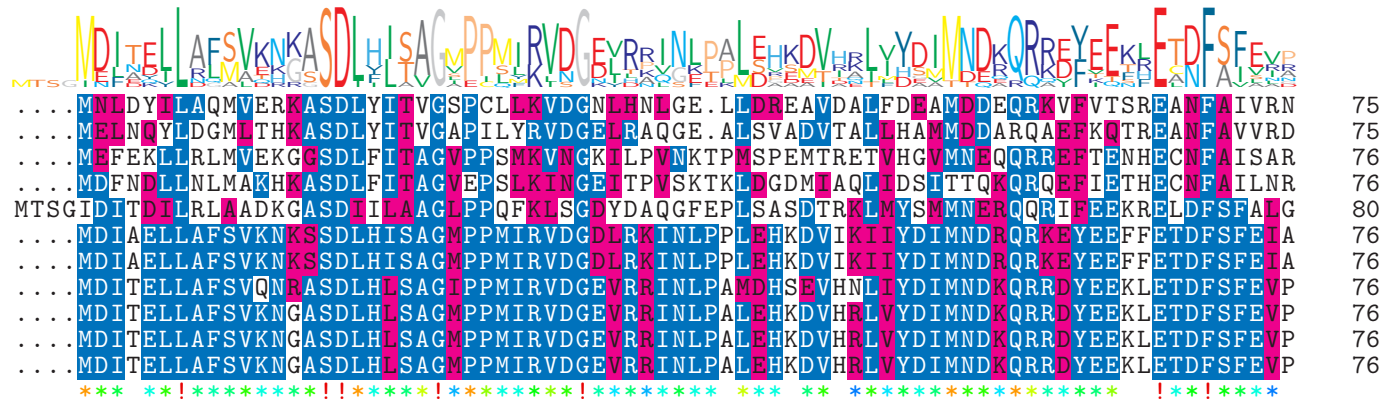


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NTDB id 1177 A1552VC RS01165 WP 000422572.1
NTDB id 1203 PSJM300 01605 AFN76401.1
NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 1307 DR RS10055 WP 01088596.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GC085 RS10305 WP 010947729.1
NTDB 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

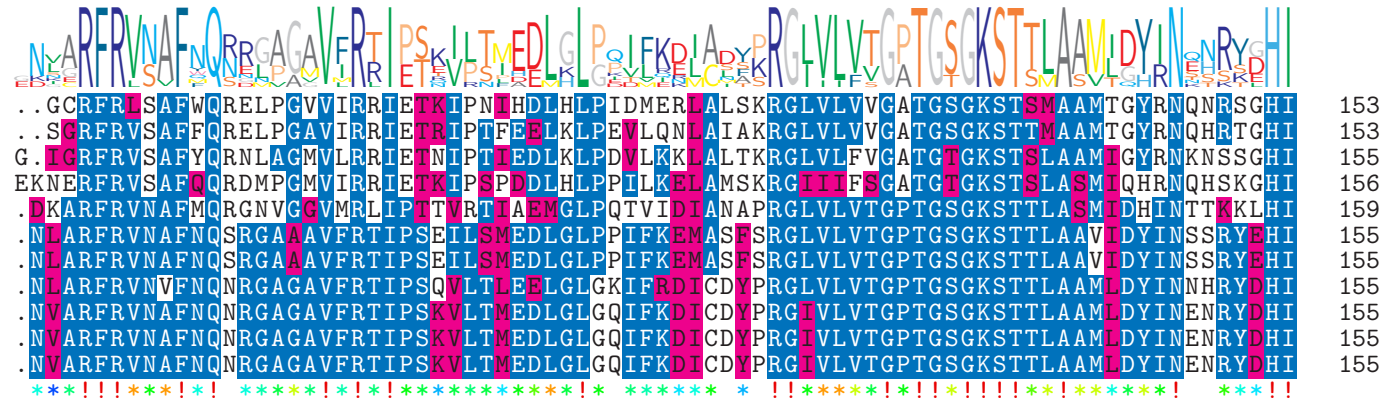
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NTDB id 1307 DR RS10055 WP 01088596.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GC085 RS10305 WP 010947729.1
NTDB id 1020 ACIAD RS04210 WP 011182164.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
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NTDB id 1040 HON27 RS13360 WP 000355489.1
consensus

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NTDB id 1104994 ACN08L RS02870 WP 008987965.1
NTDB id 1177 A1552VC RS01165 WP 000422572.1
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NTDB id 1040 HON27 RS13360 WP 000355489.1
consensus

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logo

NTDB id 1104994 ACN08L RS02870 WP 008987965.1	ALERTIHIVPKERRDQFLFDLSLNLKCILAQQLIPDANGIGRHCYVYELLINTPRVADLIRRGELHEIKDVMAKSNQAGMV	313
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NTDB id 1203 PSJM300 01605 AFN76401.1	ALDRIINFFPADRQNNQVWMDLSLNLKAIVAQQLVPTPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLMQ	315
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NTDB id 1307 DR RS10055 WP 010888596.1	SIDRIVDVFPEEQEQEIRVQLANNLVAVMTQQLPRADGQGRVLAYEILLANPAVRALIREGKTFFQITSVMQTGAREGMV	319
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TINRIIDVFPAEEKSMVRSMLSESLQAVVAQSLLK.KNKGGRVAALEIMMCTGAIRNLIREDKIAQMYSSIQTGQAKGMQ	314
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consensus	***!***** *!***!*****!*** *****!*** !*** *****!***** ***** !!	

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NTDB id 1104994 ACN08L RS02870 WP 008987965.1	TFDQSLYALFRDCKITEQDALHHADS PNDLRLMIKMGDPNNKT...ASSLSGVKL.....DL.....	367
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NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPTARTAAKQPESFL.....	345
consensus	!***! !* ****!* ** ! * *	

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NTDB id 1104994 ACN08L RS02870 WP 008987965.1	PATPAAPAAGRSEPGRAGGFGRR	367
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NTDB id 1060 ABD1 RS04380 WP 000355489.1		345
NTDB id 1040 HON27 RS13360 WP 000355489.1		345
consensus		

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