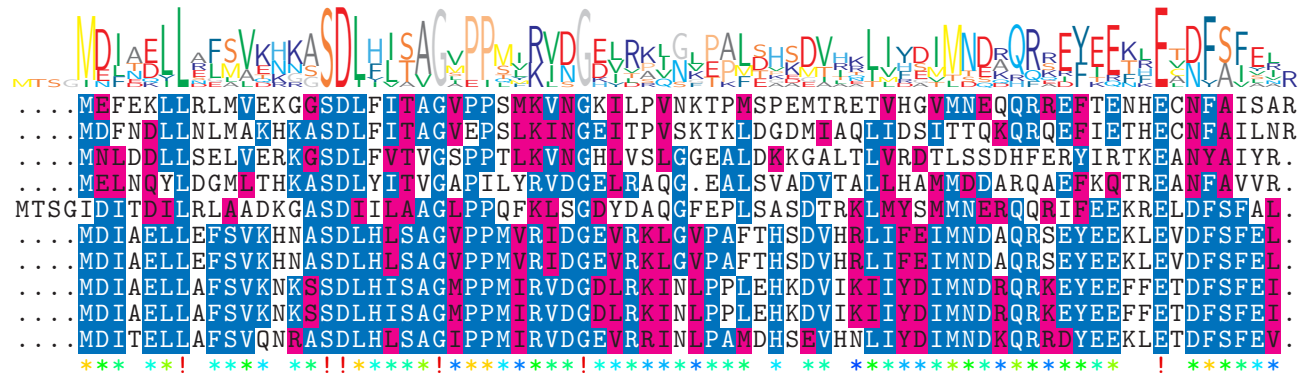


logo

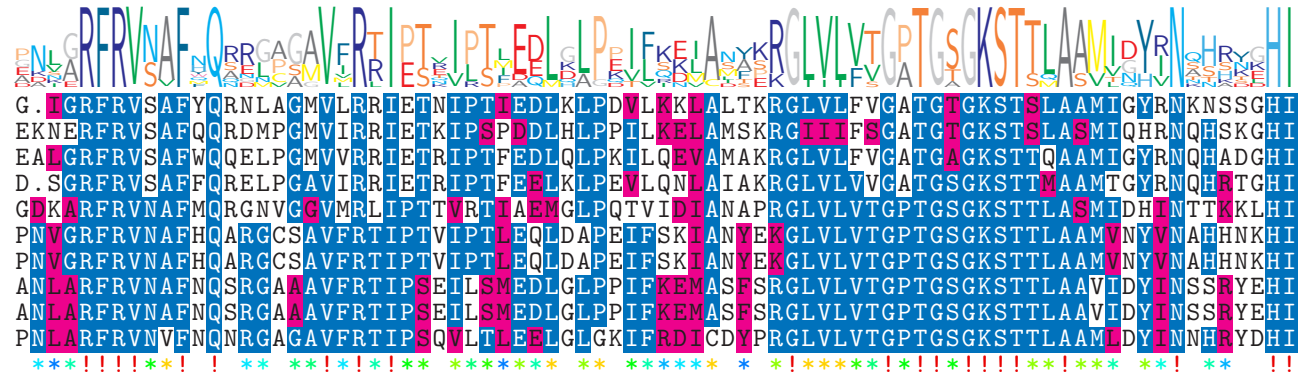
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 NTDB id 366764 FGM04 RS10735 WP 005340356.1  
 NTDB id 1177 A1552VC RS01165 WP 000422572.1  
 NTDB id 1307 DR RS10055 WP 010888596.1  
 NTDB id 1185 GTF74 RS11940 WP 000350195.1  
 NTDB id 1176 A1552VC RS01160 WP 000350195.1  
 NTDB id 1283 ACE17W RS02630 WP 010947729.1  
 NTDB id 1259 GC085 RS10305 WP 010947729.1  
 NTDB id 1020 ACIAD RS04210 WP 011182164.1  
 consensus



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NTDB id 1203 PSJM300 01605 AFN76401.1  
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 NTDB id 1177 A1552VC RS01165 WP 000422572.1  
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 NTDB id 1283 ACE17W RS02630 WP 010947729.1  
 NTDB id 1259 GC085 RS10305 WP 010947729.1  
 NTDB id 1020 ACIAD RS04210 WP 011182164.1  
 consensus



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NTDB id 1203 PSJM300 01605 AFN76401.1  
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 NTDB id 1283 ACE17W RS02630 WP 010947729.1  
 NTDB id 1259 GC085 RS10305 WP 010947729.1  
 NTDB id 1020 ACIAD RS04210 WP 011182164.1  
 consensus



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NTDB id 1019 ACIAD RS04205 WP 004922051.1	ALDRRIINFFPADRQNVVWMDLSLNLKAIVAQQLVPTPDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ	316
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NTDB id 1177 A1552VC RS01165 WP 000422572.1	ALDRILHLVLPQDKHRQFLFDLSFNLKAIVAQQLVPSMDGKRRCAAFEILLNTPLITDIIRKGEHRLKEVMTKSTELGMQ	313
NTDB id 1307 DR RS10055 WP 010888596.1	ALERILHLVLPKEQREQFLDLNLSLNLKGVIAQQLLRDKNGKGRHGVFEVLLNSPRIADLIRRGELHELKATMARSQEVGMQ	319
NTDB id 1185 GTF74 RS11940 WP 000350195.1	SIDRIVDVFPPEEQEQIRVQLANNLVAVMTQQLLRADGQGRVLAYEILLANPAVRALIREGKTFQITSVMQTGAAREGMV	314
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TIDRIIDVFPDSDKDMVRSMLSESLRAVIAQKLLKR.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAHGMQ	314
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TIDRIIDVFPDSDKDMVRSMLSESLRAVIAQKLLKR.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAHGMQ	314
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NTDB id 1020 ACIAD RS04210 WP 011182164.1	TINRIIDVFPAAEAKSMVRSMLSESLQAVVAQSLLKK.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ	314
consensus	TIDRIIDVFPAAEKDMVRTMLSESLQAVISQTLLKK.EGGGRVAAHEIMIGIPAIRNLIREGKVAQMYSIIQTGATHGMT	314

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NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALKQYORGISYTALEHADNDLRLFLSSEADGLLSSMONVTLDMSEDCRSEFR	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	LEAATFLEKNEVPEADVIAEAFKSAAMKLSSEADGLLSSMONVTLDMSEDCRSEFR	383
NTDB id 366764 FGM04 RS10735 WP 005340356.1	TFDQALYNYLVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLT.....LEMSDDDPGRSFR.....	369
NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALFSLFCAGQIGYSEALAHADSANDLRLLIKLSGRERLG...TGTLDNVT.....LDE.....	368
NTDB id 1307 DR RS10055 WP 010888596.1	TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDYG...SGSLQNVK.....IDME.....	395
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMDAYLAGLFRHLLITYDKGLER.A..IDPKEFARLANDASAGLGGA.SMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	345
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NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEA.A..LNKSLF.....	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEA.A..LNKSLF.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKQLIAKGIINLPTARAV.A..KSPESF.....	344
consensus	!*** *!* ***** ** * ***** *	

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NTDB id 1203 PSJM300 01605 AFN76401.1	PATPAAPAAGRSEPCGRAGGFGRR	381
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NTDB id 1259 GC085 RS10305 WP 010947729.1	.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	.....	344
consensus	.....	

- X non conserved
- X similar
- X ≥ 50% conserved