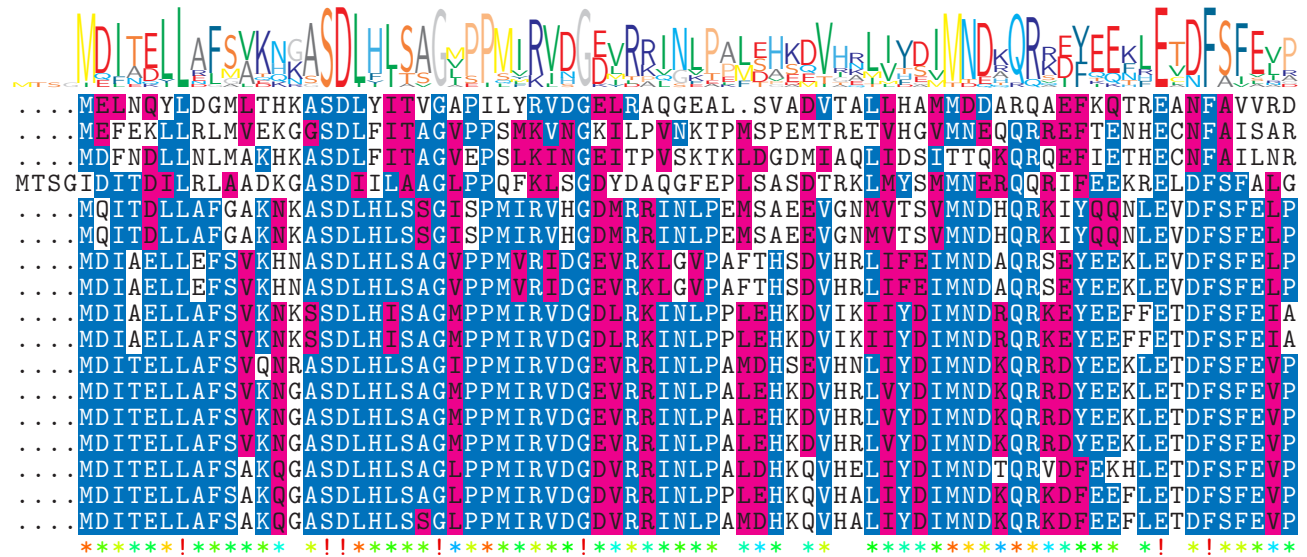


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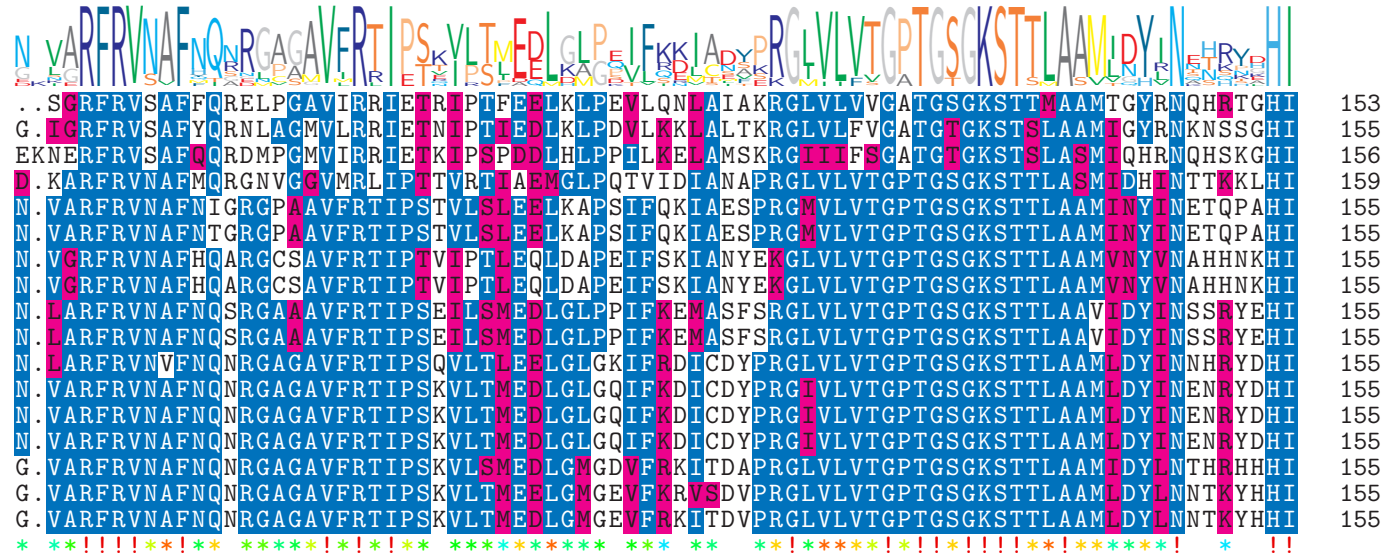
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consensus



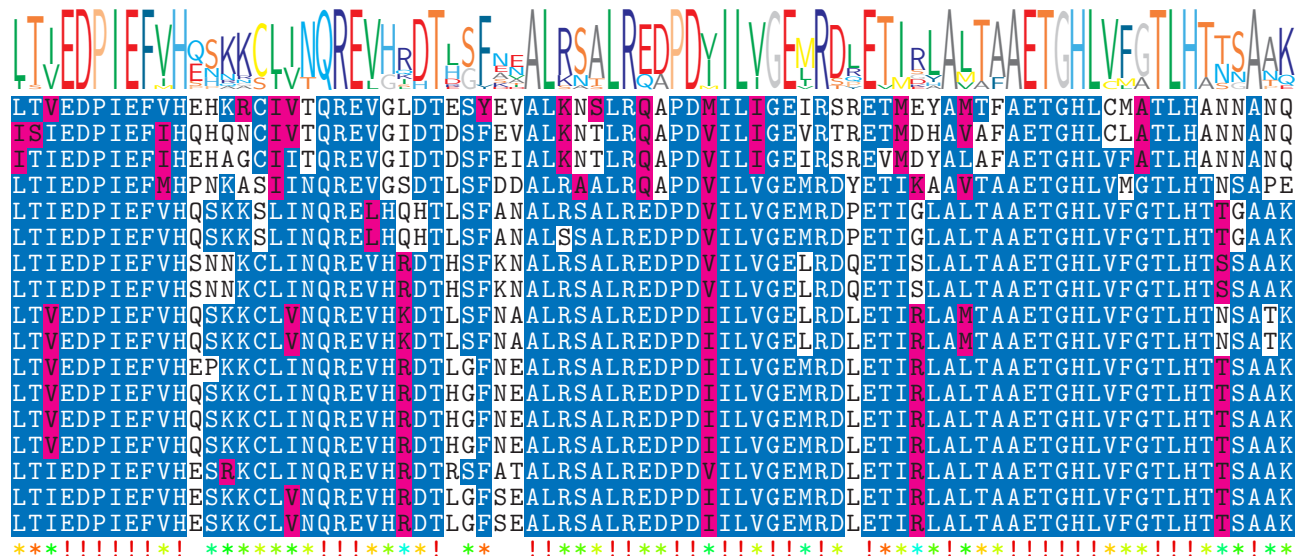
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consensus



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consensus

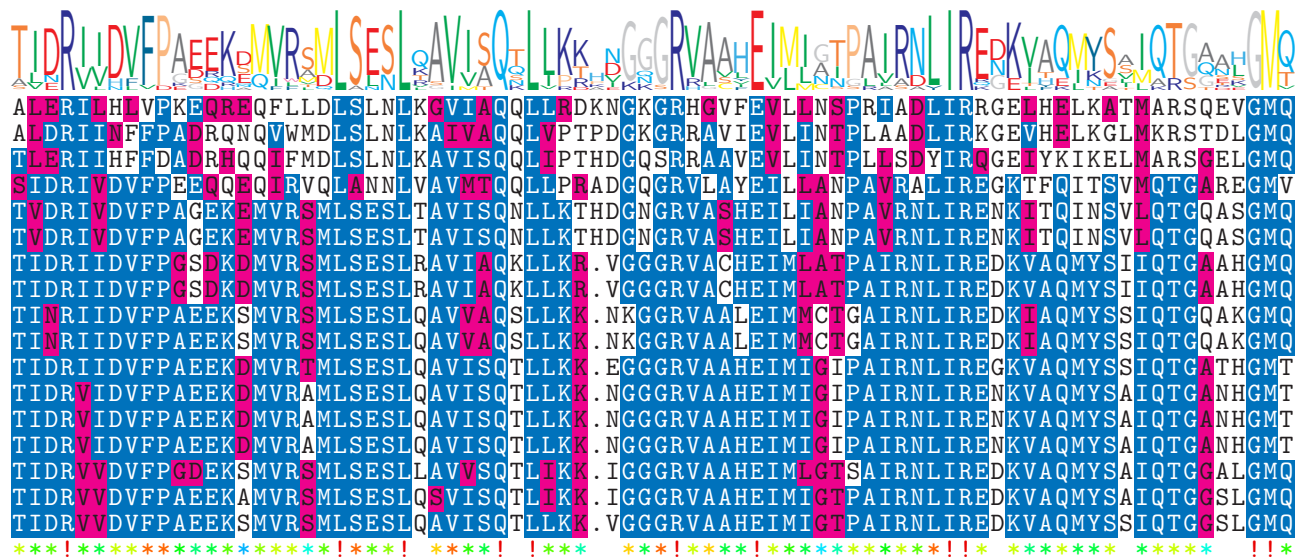


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NTDB id 1202 PSJM300 01600 AFN76400.1

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consensus



Sequence	Score
TFDQALYQLVDDKISEQDALHSADSANDLRLMLKTKRGDDY... GSGSLQNVKI..... DME.....	368
TFDQALYNLYVQGETIYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS. QGLTL..... EMSDDDPGRSFR.....	381
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TMDQSLQSLVRQGLIAPEVARRRRAQNSSESMSF.....	347
TMDQSLQSLVRQGLIAPEAARRRAQNSSESMSF.....	347
TMEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345
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TLDMCLKDLVTKGLISREHAREKARTPDNF.....	344
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NTDB	id	1081	FDQ49	RS05335	WP	006581387.1		345
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NTDB	id	75697	KI231	RS27610	WP	213026844.1		344
NTDB	id	1194	PAKAF	RS01995	WP	003084552.1		344
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consensus								

-  non conserved
-  similar
-  $\geq 50\%$ conserved