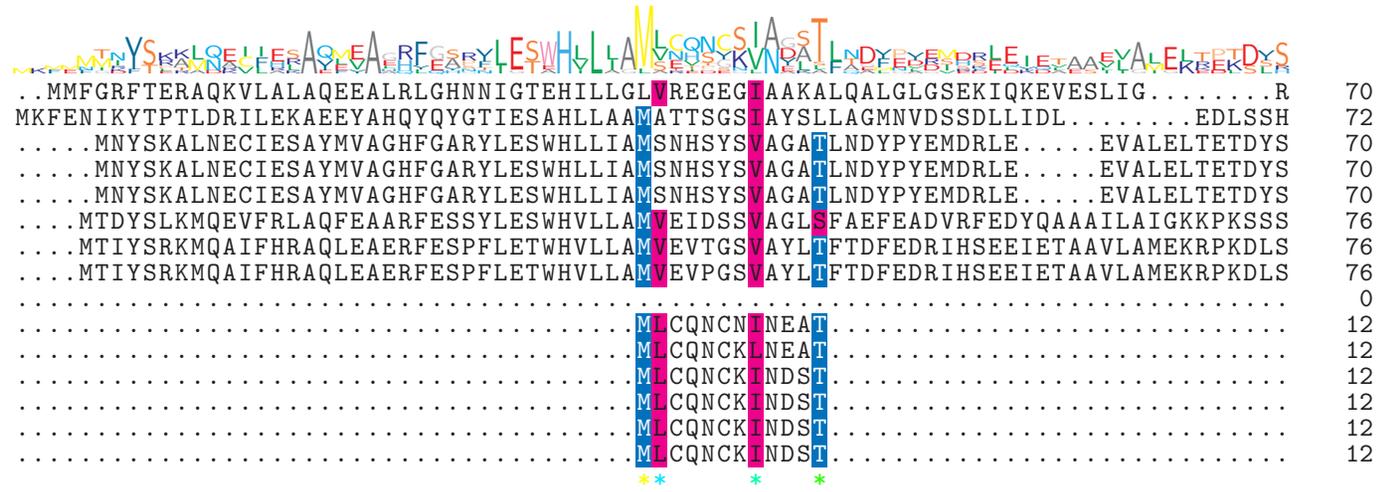


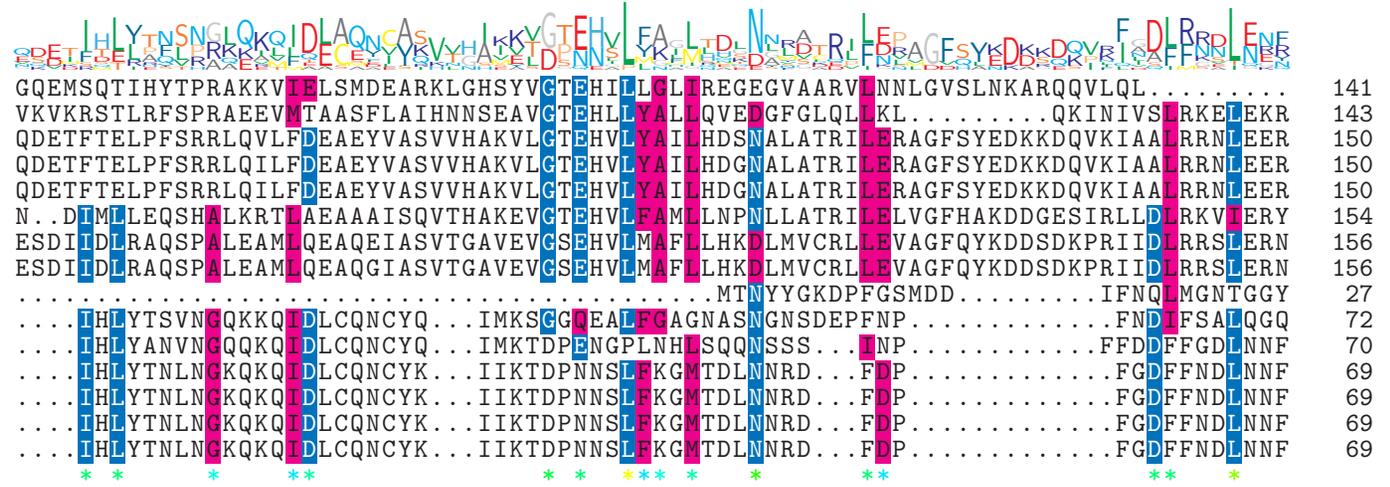
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 consensus



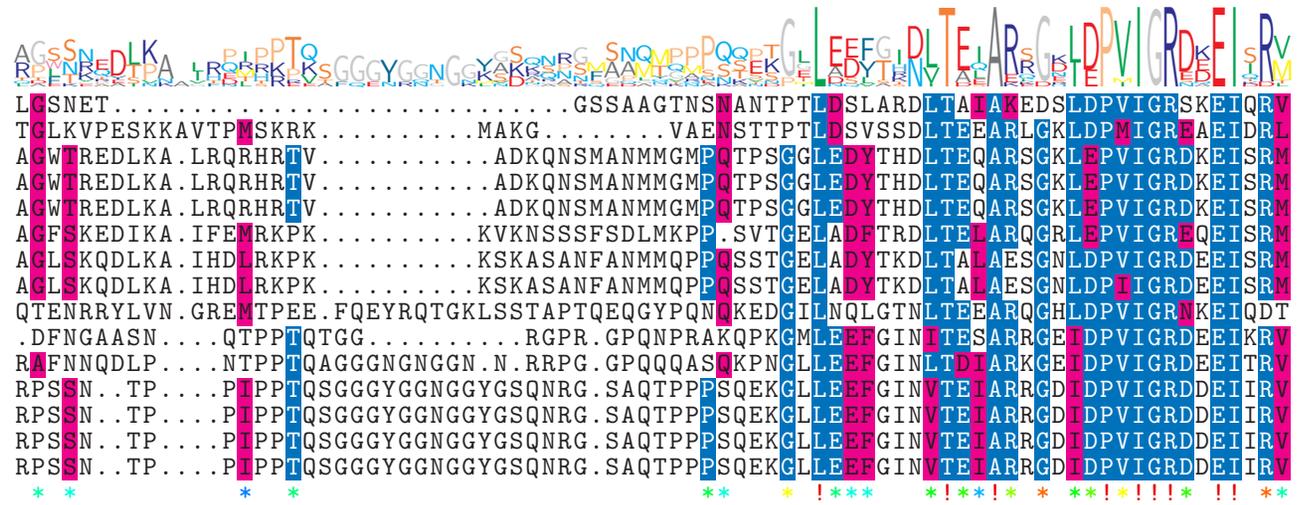
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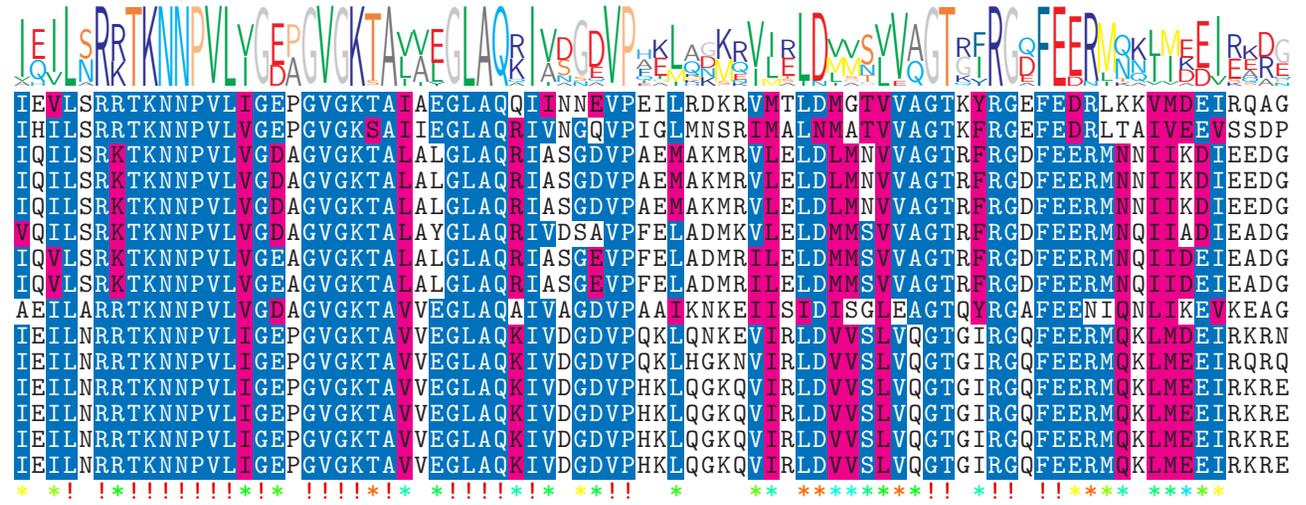
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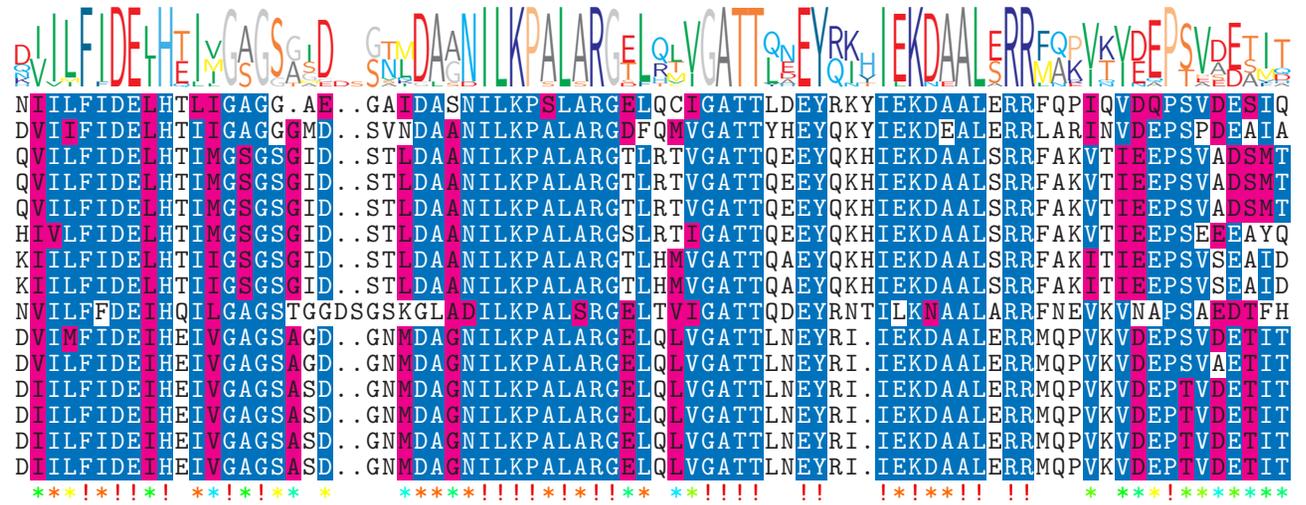
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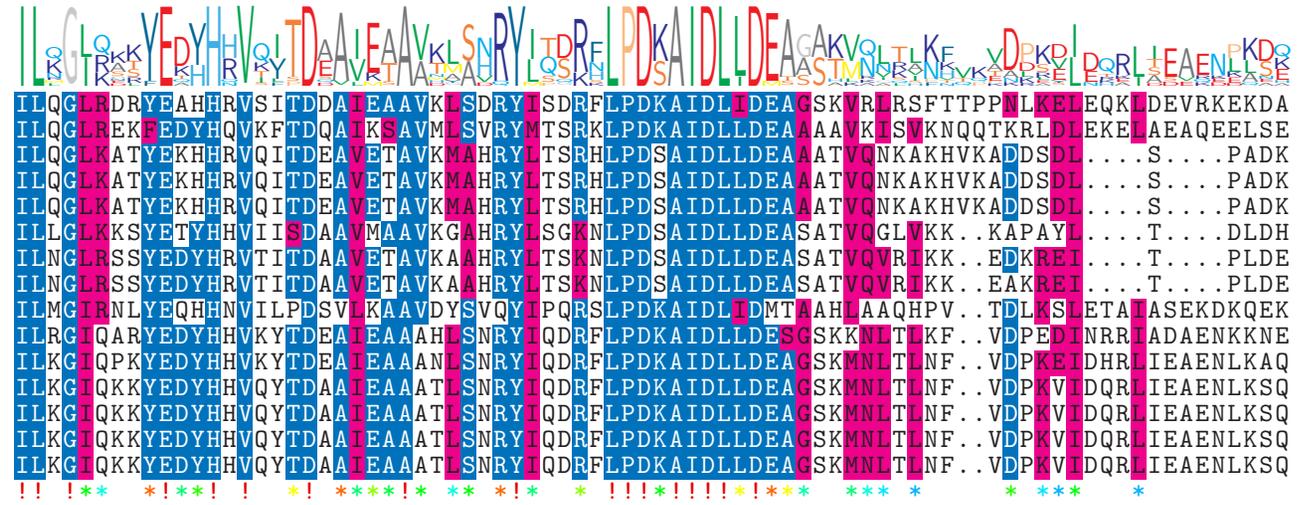
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NTDB id 610 V4T04 RS10165 WP 012897346.1	KLIAEVGFDPYEGARPLRKAQKEVEDLLSEQLLSG.....EIKAGNHVSTIGASNKKIKIAQIV....	816
NTDB id 287 SP RS11210 WP 001109712.1	KLLANQGYDPEMGARPLRRTLQTEVEDKLAELLKLG.....DLVAGSTLKIGVKAGQLKFDIA.....	810
NTDB id 290 KZH43 RS10025 WP 001109677.1	KLLANQGYDPEMGARPLRRTLQTEVEDKLAELLKLG.....DLVAGSILKIGVKAGQLKFDIA.....	810
NTDB id 289 SPD RS10700 WP 001109677.1	KLLANQGYDPEMGARPLRRTLQTEVEDKLAELLKLG.....DLVAGSILKIGVKAGQLKFDIA.....	810
NTDB id 377 SMU RS09275 WP 002262344.1	KHLALAGYDAEMGARPLRRTLQTEVEDKLAELLISG.....KLASGQTLKIGIQEKLKFDIV.....	813
NTDB id 297 STER RS00545 WP 011680614.1	KHLAKDGYDIEMGARPLRRTIQTVEDKLSSELLGG.....QVVSQTLKIGCSKDKLTFTVV.....	816
NTDB id 329 STU RS10020 WP 011225298.1	KHLAKDGYDIEMGARPLRRTIQTVEDKLSSELLGG.....QVVSQTLKIGCSKDKLTFTVV.....	816
NTDB id 1063758 ACG3JF RS00730 WP 013793858.1	EELIELGYDKAMGVRLRRVIEQEIRDRITDYYLEHPDDKHLKANLADGQIVISKQDK.....	702
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NTDB id 378 SMU RS02690 WP 002263569.1	EKLVDLGYDPKMGARPLRRTIQEHIETDAITDFYLEHPEQKELKAIMTSNGKIVIKASNK.VETVTKATSD.	753
NTDB id 241 SP RS04015 WP 000882523.1	EKLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSACK.AEVKSSEKEK.	752
NTDB id 281 KZH43 RS03550 WP 000882517.1	EKLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSACK.AEVKSSEKEK.	752
NTDB id 245 SPD RS03860 WP 000882517.1	EKLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSACK.AEVKSSEKEK.	752
NTDB id 243 SPR RS03635 WP 000882517.1	EKLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSACK.AEVKSSEKEK.	752
consensus	* * !*** *!!!!***** **! *** *	

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