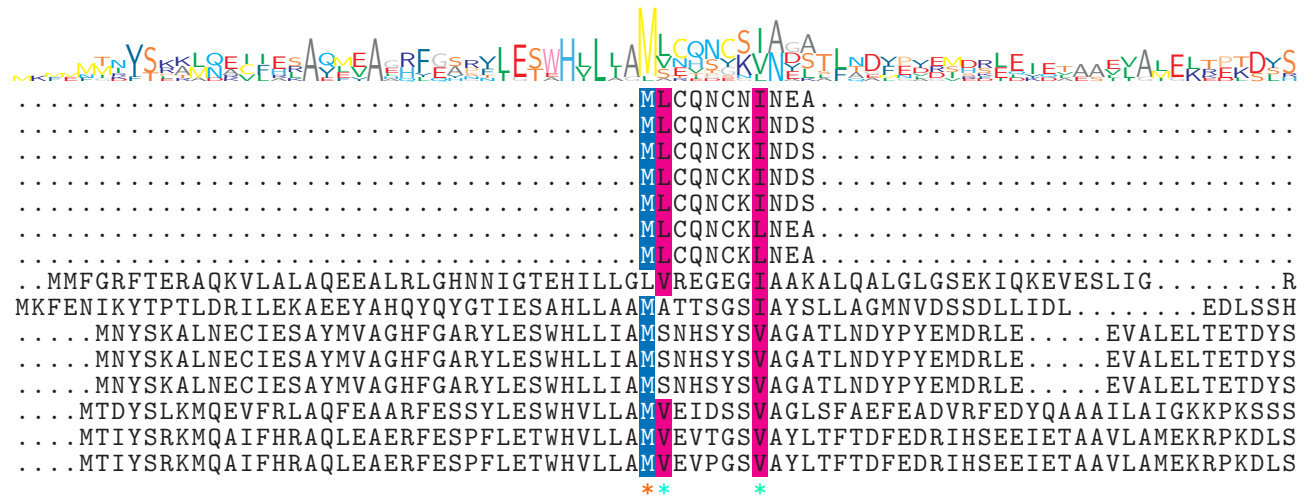


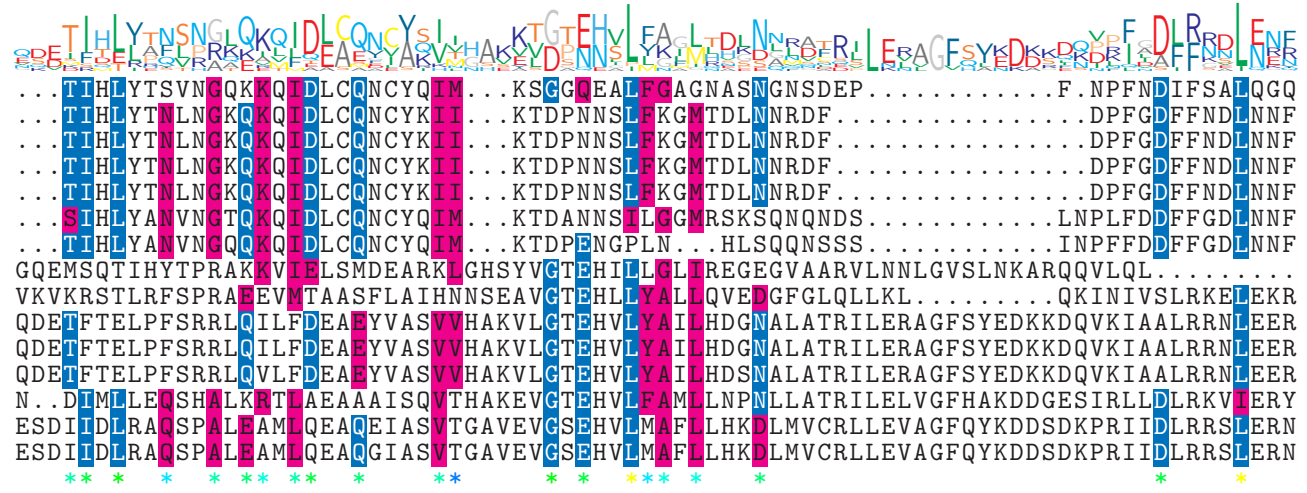
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 NTDB id 243 SPR RS03635 WP 000882517.1
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 NTDB id 289 SPD RS10700 WP 001109677.1
 NTDB id 287 SP RS11210 WP 001109712.1
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 consensus



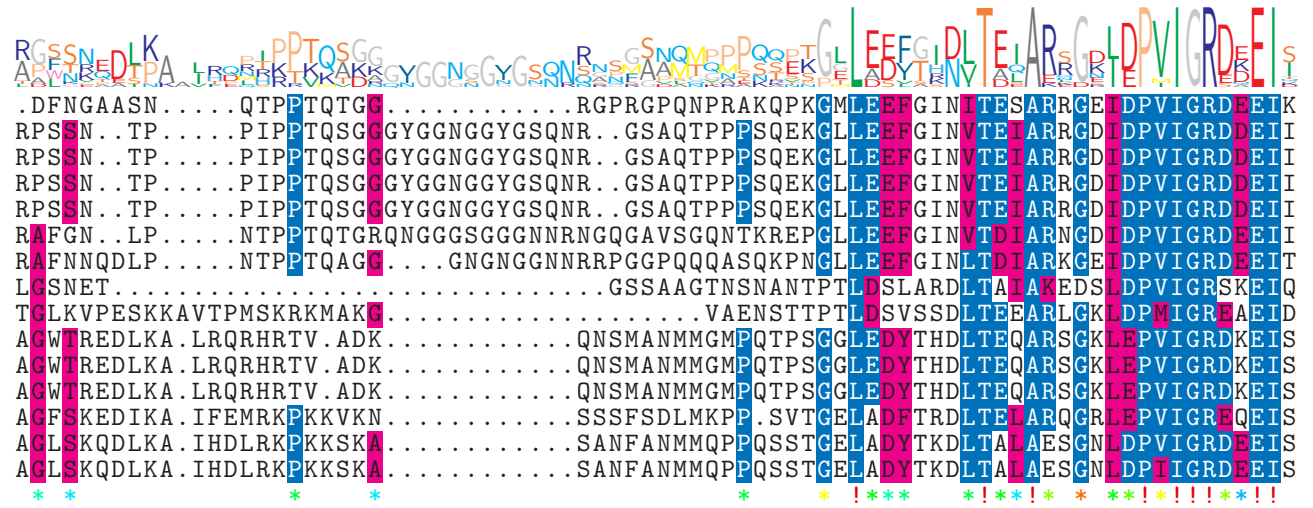
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 consensus



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NTDB id 601 KW2 RS02615 WP 021036805.1
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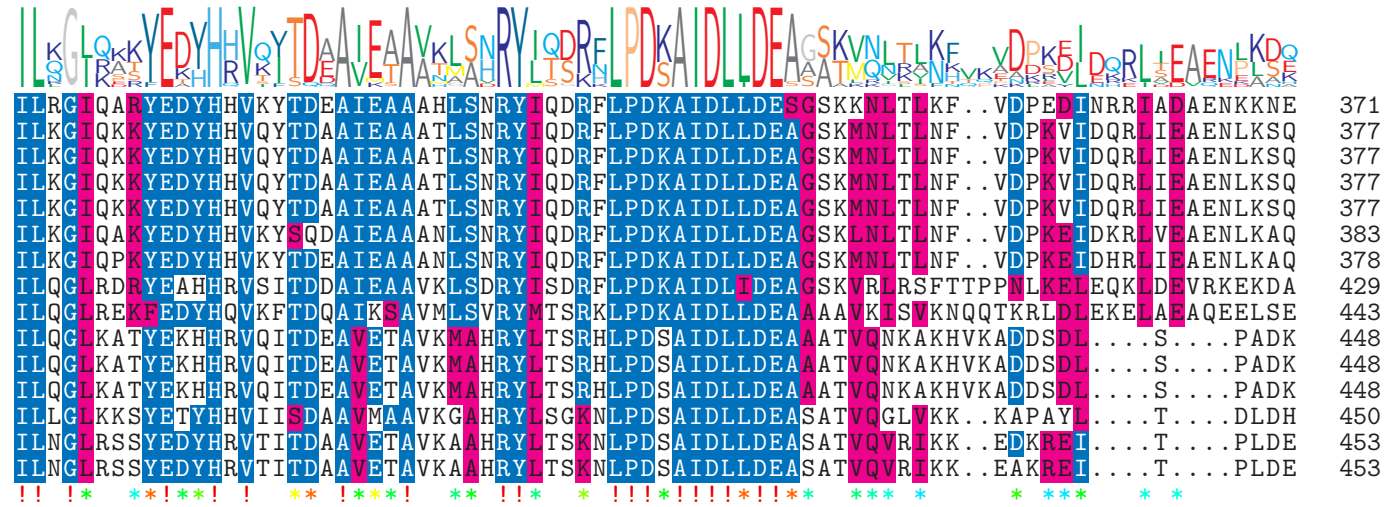
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 consensus



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NTDB id 601 KW2 RS02615 WP 021036805.1	KLVDLGYNPAMGARPLRRTIQENTEDSADFYEHPEYKQLVADLIDDKIVISNQTQETAETTDEEVPAE	748
NTDB id 281 KZH43 RS03550 WP 000882517.1	KLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAIEVKSSEKEK..	752
NTDB id 245 SPD RS03860 WP 000882517.1	KLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAIEVKSSEKEK..	752
NTDB id 243 SPR RS03635 WP 000882517.1	KLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAIEVKSSEKEK..	752
NTDB id 241 SP RS04015 WP 000882523.1	KLVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAIEVKSSEKEK..	752
NTDB id 1121994 EL147 RS03475 WP 007894778.1	KLVDLGYDAKMGARPLRRTIQDHIEDAITDYYLENPTEKNL.KAMLSNCGNIVIKSAKKLDKAQSVSQQKD	760
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NTDB id 85 BSU 00860 NP 387967.1	KVAEEGVLDLEYGARPLRRAIKHVEDRLSEELLRGNIHKGQHVLVDVEDGEFVVKTTAKTN.....	810
NTDB id 610 V4T04 RS10165 WP 012897346.1	LIAEVCFDPEYGARPLRKAQKEVEDLLSEQLLSGEIKAGNHVSTGASNKKIKIAQIV.....	816
NTDB id 290 KZH43 RS10025 WP 001109677.1	LLANQGYDPEMGARPLRRTIQTEVEDKLAPELLKGDVLVAGSILKIGVKAQQLKFDIA.....	810
NTDB id 289 SPD RS10700 WP 001109677.1	LLANQGYDPEMGARPLRRTIQTEVEDKLAPELLKGDVLVAGSILKIGVKAQQLKFDIA.....	810
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NTDB id 297 STER RS00545 WP 011680614.1	HLAKDGYDIEMGARPLRRTIQTVEDKLSPELLGGQVVSGQTLKIGCSKDKLTFTVV.....	816
NTDB id 329 STU RS10020 WP 011225298.1	HLAKDGYDIEMGARPLRRTIQTVEDKLSPELLGGQVVSGQTLKIGCSKDKLTFTVV.....	816
consensus	* * !*** *!!!!!!***! *!! *** *	

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