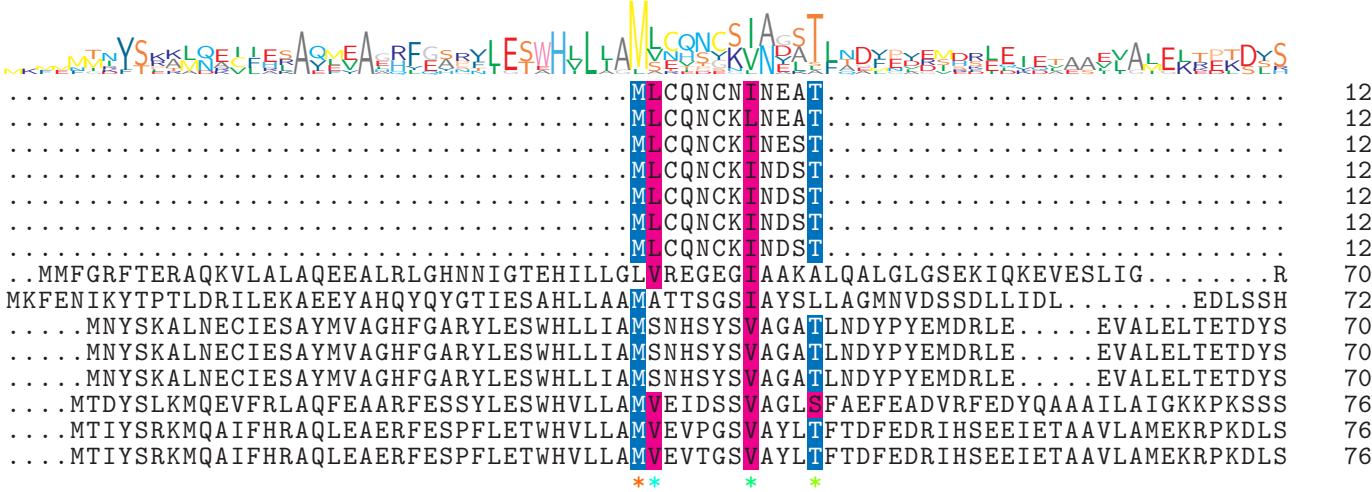


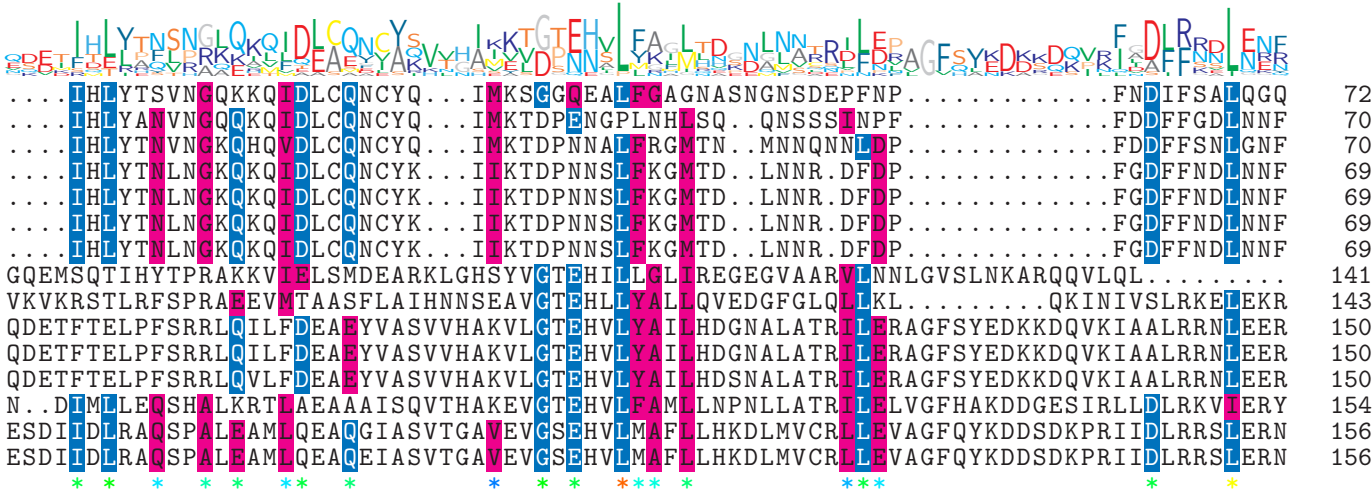
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NTDB id 329 STU RS10020 WP 011225298.1
NTDB id 297 STER RS00545 WP 011680614.1
consensus



logo

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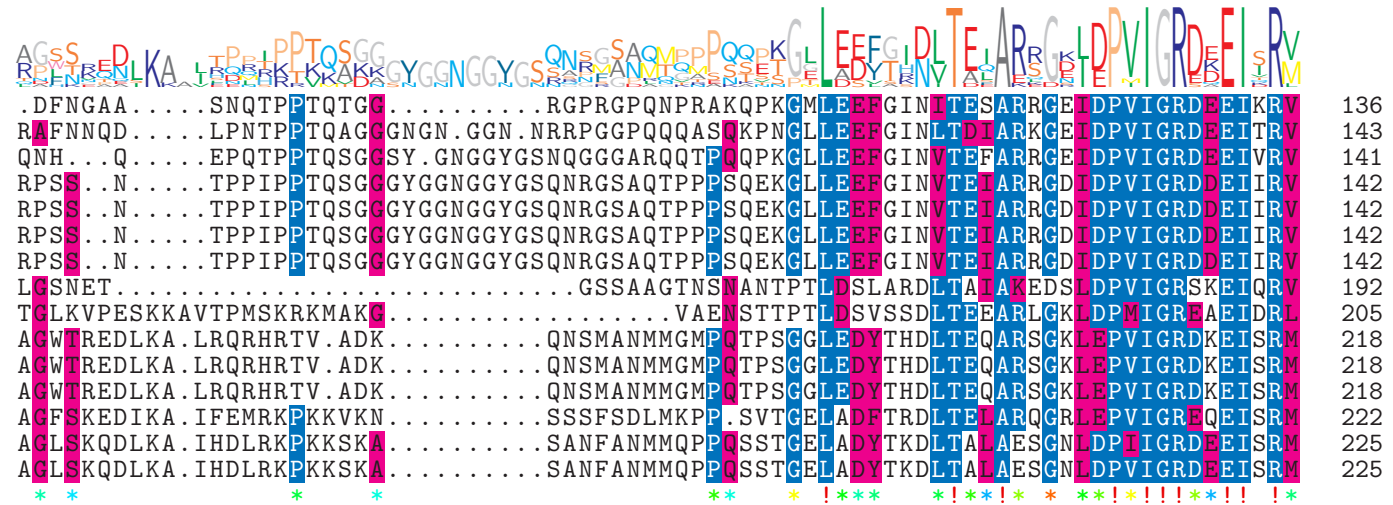


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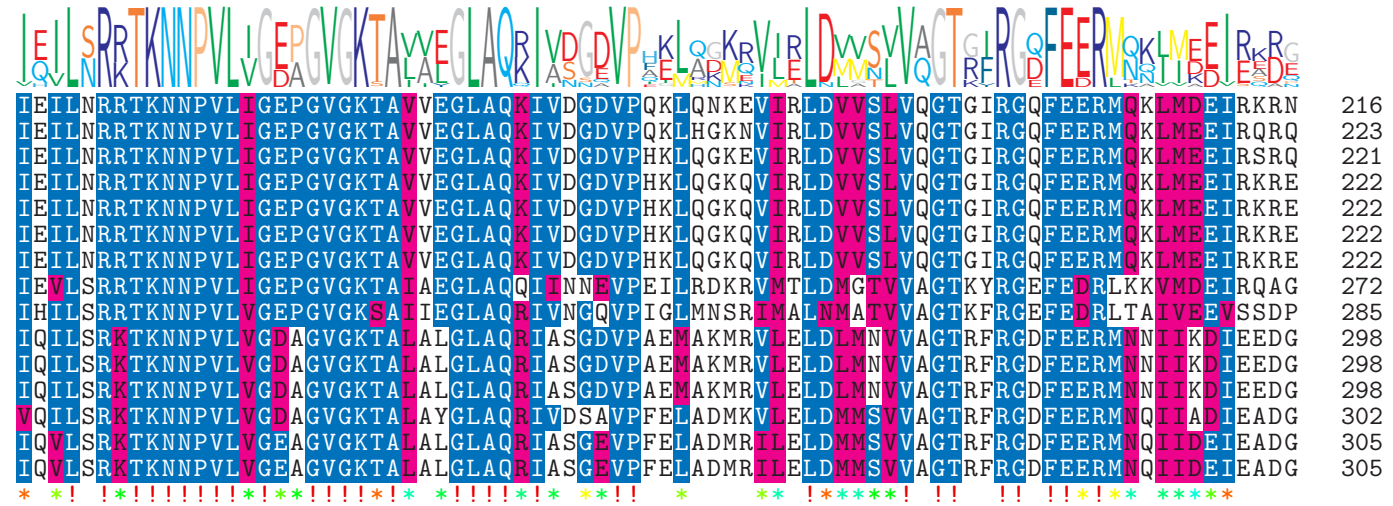


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consensus

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NTDB id	601	KW2	RS02615	WP	021036805.1	DVILFIDELTHEIVGAGSAGDGNMDAGNILKPALARGE	295
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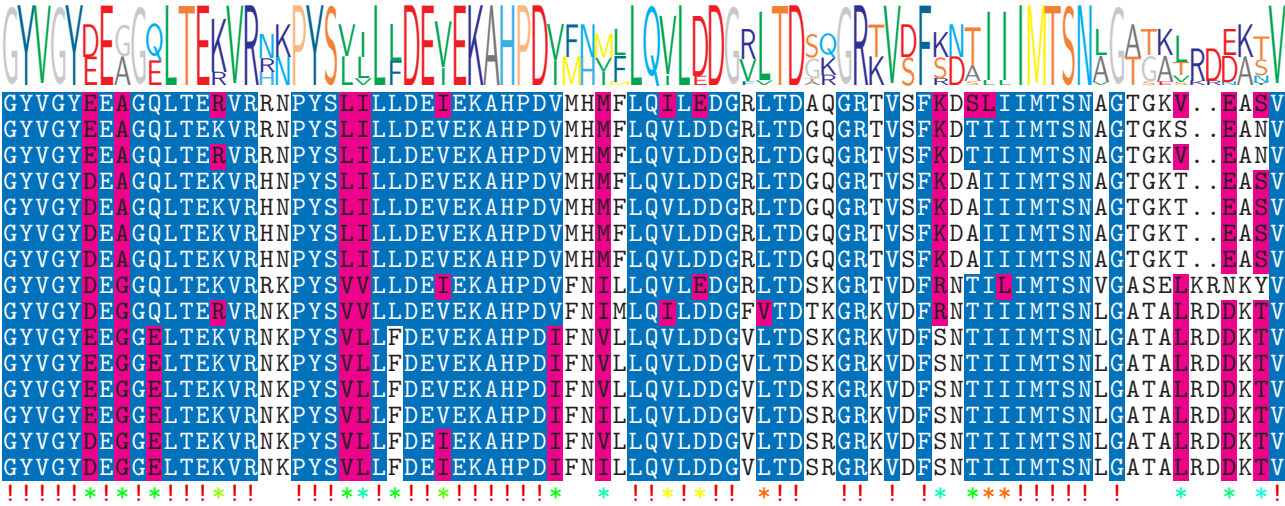
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consensus



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NTDB id 601 KW2 RS02615 WP 021036805.1	VDLGYNPAMGARPLRRTIQENTIEDSTADFYIEHPEYKQLVADLIDDKIVISNQTQ..ETAETTTDEEVPAE.....	748
NTDB id 378 SMU RS02690 WP 002263569.1	VDLGYDPKMGARPLRRTIQEHIEDAITDFYLEHPEQKELKATMTSNGKIVIKASNKVETVTKATSD.....	753
NTDB id 156787 SanJ4206 RS06965 WP 037582766.1	VNLGYDPKMGARPLRRTIQDHIEDAITDYYLEHPSEKELKAVMTSNGKIVIKSTKKAENEANKEIQTPAADDNAN	760
NTDB id 281 KZH43 RS03550 WP 000882517.1	VDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 245 SPD RS03860 WP 000882517.1	VDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 243 SPR RS03635 WP 000882517.1	VDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 241 SP RS04015 WP 000882523.1	VDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDLKAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 85 BSU 00860 NP 387967.1	AEEGVLDLEYGARPLRRRAIQKHVEDRLSEELLRGNIHKGQHIVLDVEDGEFVVVKTAKTN.....	810
NTDB id 610 V4T04 RS10165 WP 012897346.1	AEVGFDPPEYGARPLRKALQKEVEDLLSEQLLSGEIKAGNHVSTIGASNKKTKIAQIV.....	816
NTDB id 290 KZH43 RS10025 WP 001109677.1	ANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDLVAGSILKIGVKAGQLKFDIA.....	810
NTDB id 289 SPD RS10700 WP 001109677.1	ANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDLVAGSILKIGVKAGQLKFDIA.....	810
NTDB id 287 SP RS11210 WP 001109712.1	ANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDLVAGSTLKIGVKAGQLKFDIA.....	810
NTDB id 377 SMU RS09275 WP 002262344.1	ALAGYDAEMGARPLRRTLQTEVEDKLAELLLSGKLASGQTLKIGIQKEKLKFDIV.....	813
NTDB id 329 STU RS10020 WP 011225298.1	AKDGYDIEMGARPLRRTIQTQVEDKLSELLLGGQVVSGQTLKIGCSKDKLTFTVV.....	816
NTDB id 297 STER RS00545 WP 011680614.1	AKDGYDIEMGARPLRRTIQTQVEDKLSELLLGGQVVSGQTLKIGCSKDKLTFTVV.....	816
consensus	!*** *!!!!!!***! *!! *** *	

- non conserved
- similar
- ≥ 50% conserved