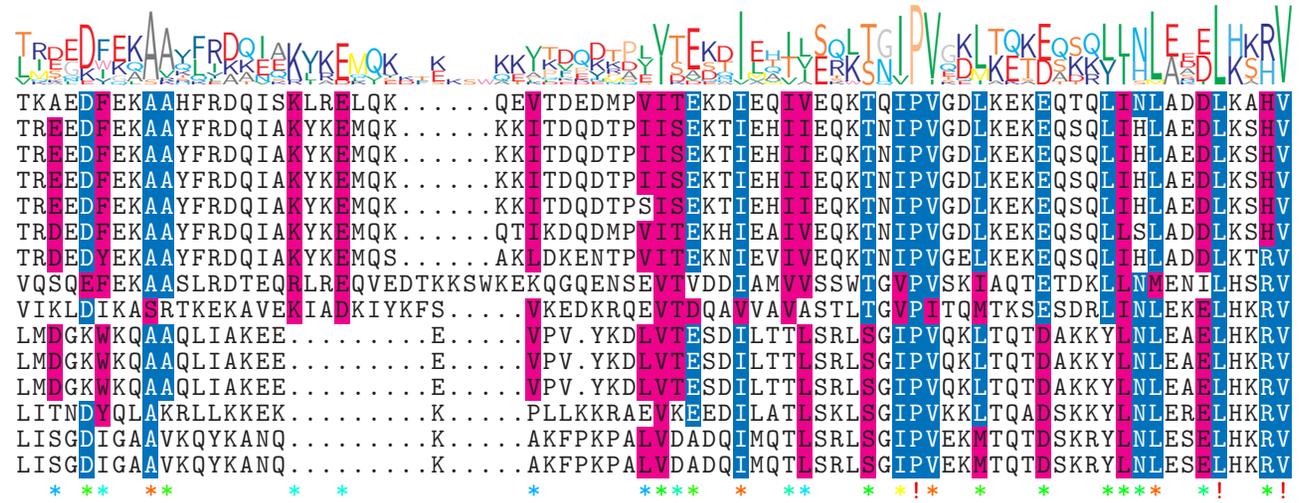


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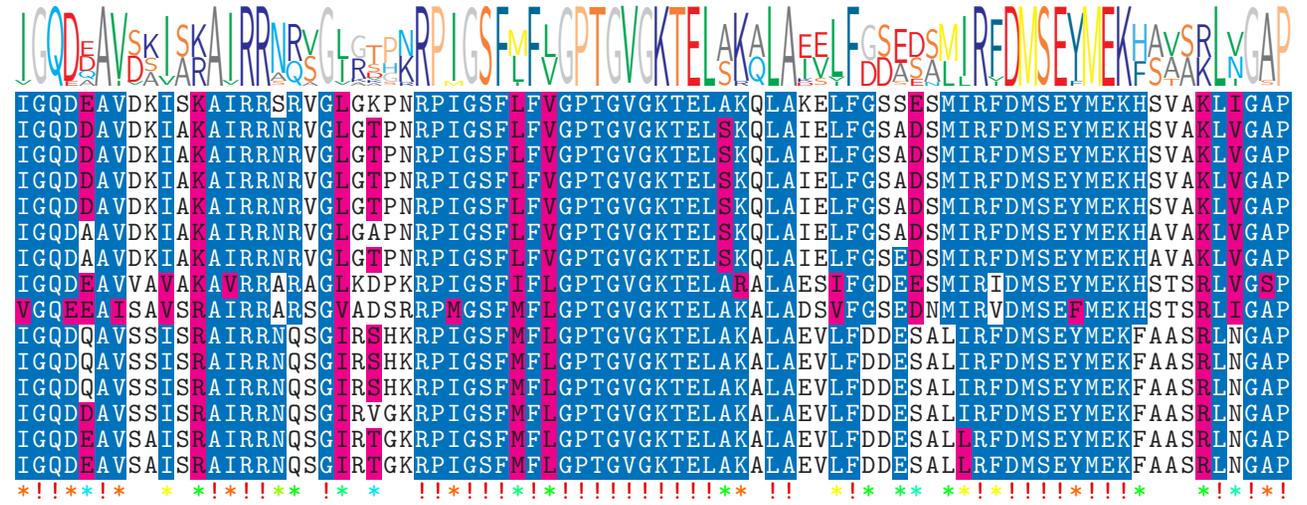
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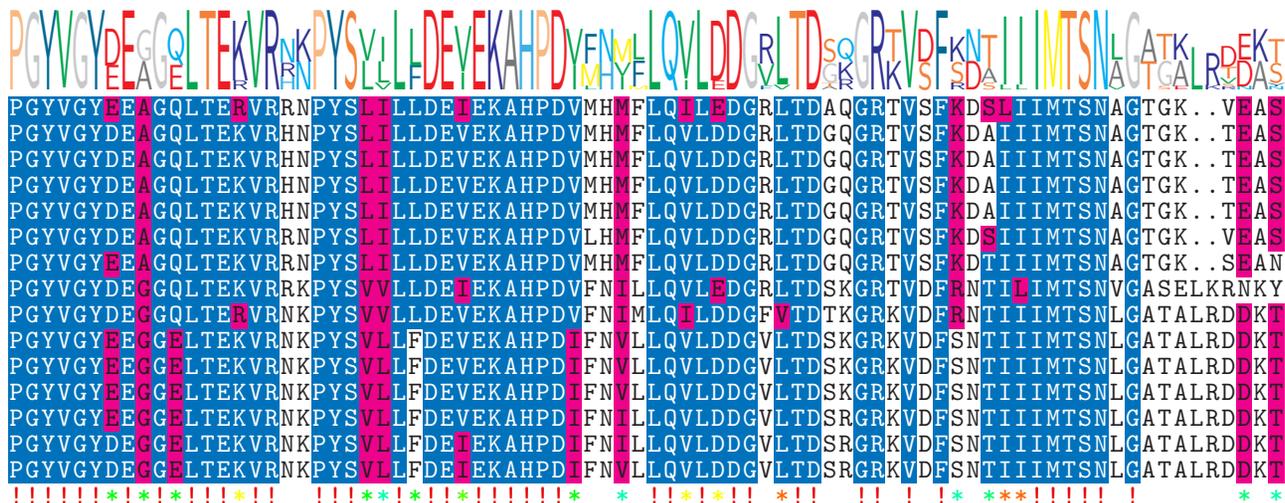


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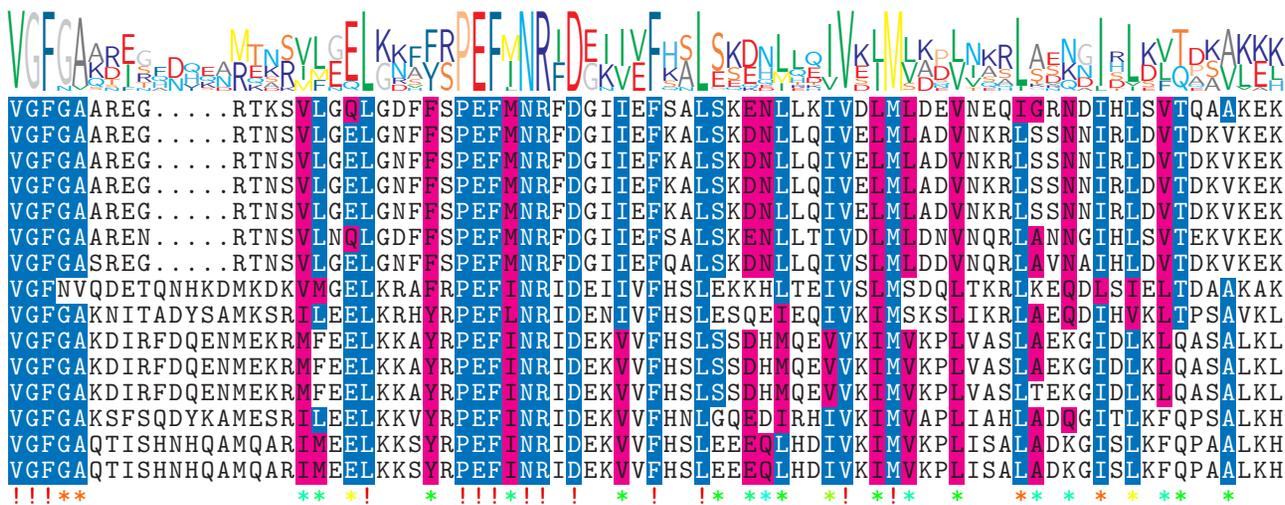


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consensus



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NTDB id 245 SPD RS03860 WP 000882517.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDL.KAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 243 SPR RS03635 WP 000882517.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDL.KAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 241 SP RS04015 WP 000882523.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKDL.KAVMTSKGNIQIKSAKKAEVKSSEKEK.....	752
NTDB id 986332 NST80 RS07365 WP 045768968.1	LVDLGYDPKMGARPLRRTIQDHIEDAITDFYLENPNEKDL.KAVMTSKGHIITIKSAKKAETTTQAETSKETD	756
NTDB id 378 SMU RS02690 WP 002263569.1	LVDLGYDPKMGARPLRRTIQEHIEDAITDFYLEHPEQKEL.KAIMTSNGKIVIKASNKVETVTKATSD.....	753
NTDB id 85 BSU 00860 NP 387967.1	VAEEGVLDLEYGARPLRRAIQKHVEDRLSEELLRGNIHKGQHTVLDVDEGEFVVKTTAKTN.....	810
NTDB id 610 V4T04 RS10165 WP 012897346.1	IAEVGFDPPEYGARPLRKALQKEVEDLSEQLLSGEIKAGNHVSIIGASNKKTKIAQIV.....	816
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NTDB id 287 SP RS11210 WP 001109712.1	LANQGYDPEMGARPLRRTLQTEVEDKLAELLKGDVAGSTLKIIGVKAGQLKFDIA.....	810
NTDB id 377 SMU RS09275 WP 002262344.1	LALAGYDAEMGARPLRRTLQTEVEDKLAELISGKLASGQTLKIIGIQKEKIKFDIV.....	813
NTDB id 297 STER RS00545 WP 011680614.1	LAKDGYDIEMGARPLRRTIQTQVEDKLSPELLGGQVVSQTLLKIGCSKDKLTFTVV.....	816
NTDB id 329 STU RS10020 WP 011225298.1	LAKDGYDIEMGARPLRRTIQTQVEDKLSPELLGGQVVSQTLLKIGCSKDKLTFTVV.....	816
consensus	* * !*** *!!!!!!***! *!! *** *	

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