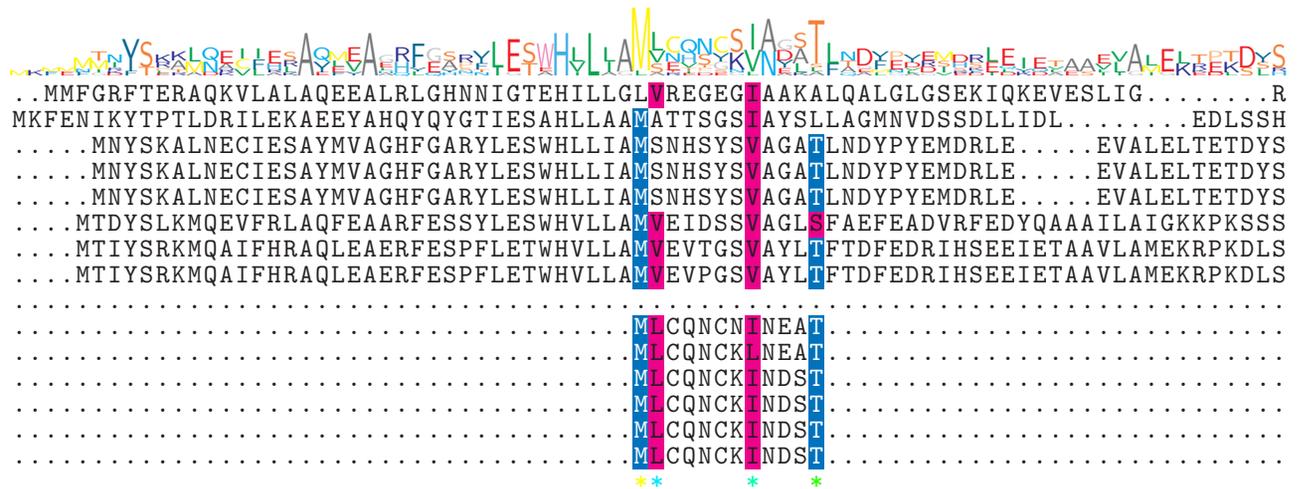


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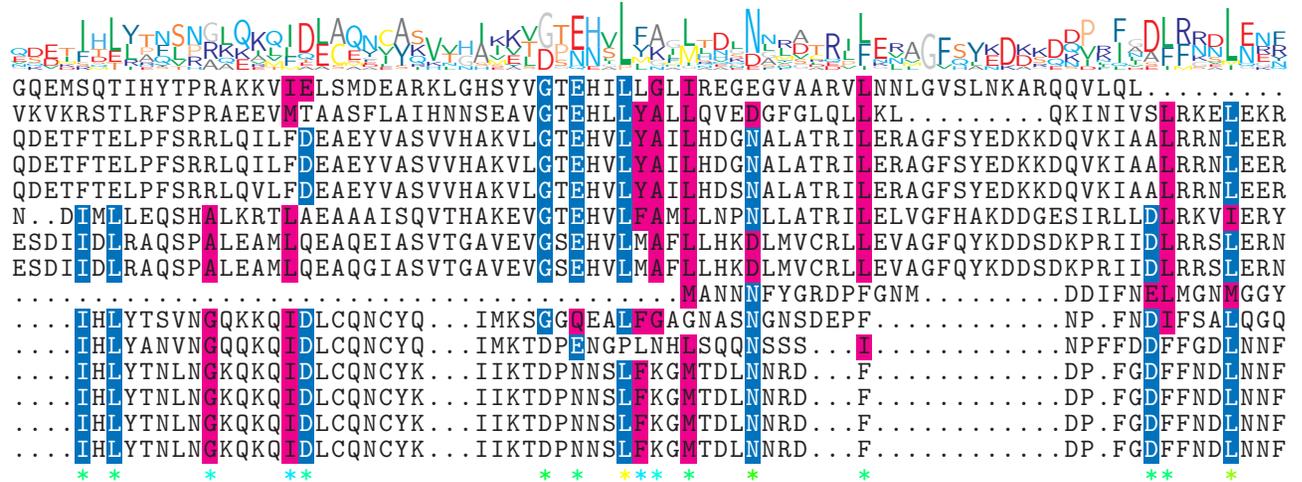
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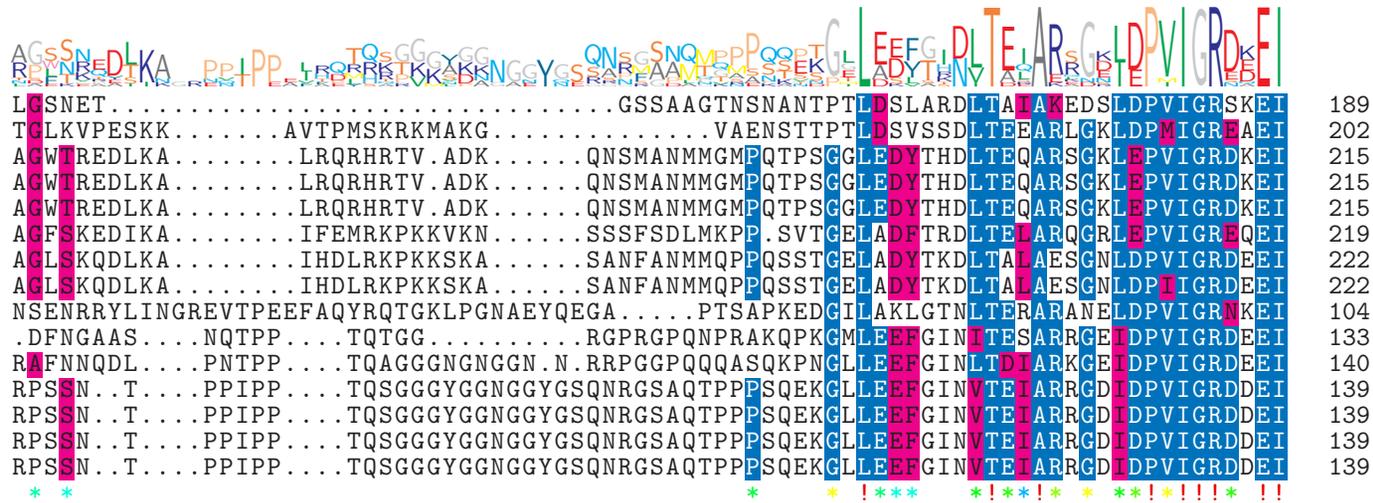
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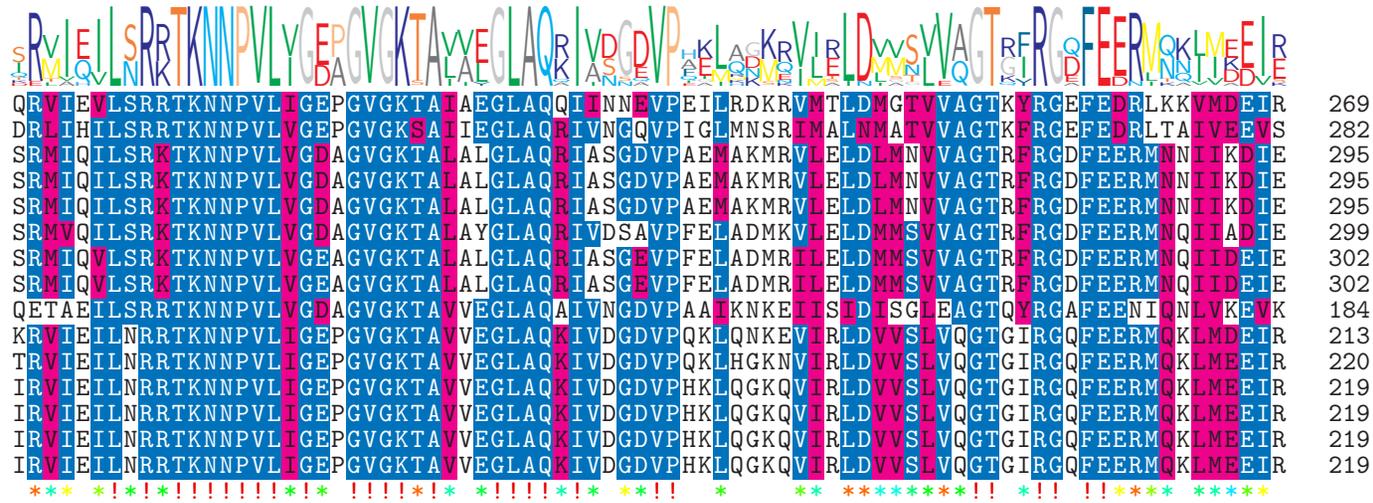
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NTDB id 290 KZH43 RS10025 WP 001109677.1	SALKLLANQ	GYDPEMGARPLRR	TIQTEVEDKLAELL	LKGDVAGSILKIGV	KAGQLKFDIA.....	810					
NTDB id 289 SPD RS10700 WP 001109677.1	SALKLLANQ	GYDPEMGARPLRR	TIQTEVEDKLAELL	LKGDVAGSILKIGV	KAGQLKFDIA.....	810					
NTDB id 287 SP RS11210 WP 001109712.1	SALKLLANQ	GYDPEMGARPLRR	TIQTEVEDKLAELL	LKGDVAGSILKIGV	KAGQLKFDIA.....	810					
NTDB id 377 SMU RS09275 WP 002262344.1	SALKHLALA	GYDAEMGARPLRR	TIQTEVEDKLAELI	LSGKLASGQTLKIGI	QKEKLFKFDIV.....	813					
NTDB id 297 STER RS00545 WP 011680614.1	AALKHLAKD	GYDIEMGARPLRR	TIQTQVEDKLSSELL	GGQVVSQTLKIGCS	KDKLTFTVV.....	816					
NTDB id 329 STU RS10020 WP 011225298.1	AALKHLAKD	GYDIEMGARPLRR	TIQTQVEDKLSSELL	GGQVVSQTLKIGCS	KDKLTFTVV.....	816					
NTDB id 40053 SGBAA2069 RS06475 WP 012962041.1	AAKQHLLI	EGYDEAMGVRPLRR	VIEQEI	RDKITDFYLDHTDVK	.HLKADMVDGELVISEK	702				
NTDB id 601 KW2 RS02615 WP 021036805.1	AAKEKLV	DLGYNPAMGARPLRR	TIQENIEDSIA	DFYIEHPEYK..QLVAD	LIDDKIVISNQTQETAETTDEEVP	AE	748				
NTDB id 378 SMU RS02690 WP 002263569.1	KVKEKLV	DLGYDPKMGARPLRR	TIQEHI	EADITDFYIEHPEQK..ELKAIM	TSNGKIVIKASNK.VETVTKATSD	.	753				
NTDB id 241 SP RS04015 WP 000882523.1	KVKEKLV	DLGYDPKMGARPLRR	TIQDYIEDTITD	YYLENPSEK..DLKAVMT	SKGNIQIKSACK.AEVKSSEKEK	.	752				
NTDB id 281 KZH43 RS03550 WP 000882517.1	KVKEKLV	DLGYDPKMGARPLRR	TIQDYIEDTITD	YYLENPSEK..DLKAVMT	SKGNIQIKSACK.AEVKSSEKEK	.	752				
NTDB id 245 SPD RS03860 WP 000882517.1	KVKEKLV	DLGYDPKMGARPLRR	TIQDYIEDTITD	YYLENPSEK..DLKAVMT	SKGNIQIKSACK.AEVKSSEKEK	.	752				
NTDB id 243 SPR RS03635 WP 000882517.1	KVKEKLV	DLGYDPKMGARPLRR	TIQDYIEDTITD	YYLENPSEK..DLKAVMT	SKGNIQIKSACK.AEVKSSEKEK	.	752				
consensus	*	*	*	!***	*!***	!***	***	*	**	*	*

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