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NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 281 KZH43 RS03550 WP 000882517.1
NTDB id 245 SPD RS03860 WP 000882517.1
NTDB id 243 SPR RS03635 WP 000882517.1
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NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

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[illegible]

logo

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NTDB id 329 STU RS10020 WP 011225298.1
consensus

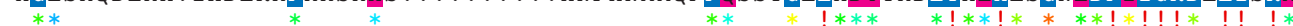
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.....IHLYTNLNGKQKKQIDLCQNCYK...IIKTDPNNSLFKGMTD...LNNRD.FDP.....FGDFFNDL...	66
.....IHLYTNLNGKQKKQIDLCQNCYK...IIKTDPNNSLFKGMTD...LNNRD.FDP.....FGDFFNDL...	66
.....IHLYTNLNGKQKKQIDLCQNCYK...IIKTDPNNSLFKGMTD...LNNRD.FDP.....FGDFFNDL...	66
.....IHLYNNLNGKQKKQIDLCQNCYQ...IMKTDPNNAILRGLGD...LTNPNNMDP.....FSEFFNHL...	67
.....IHLYANVNGKQKKQIDLCQNCYQ...IMKTDPENGPLNHLG...QNSSSTINPF.....FDFFFGDL...	67
GQEMSQTIHYTPRAKKVIELSMDEARKLGHSYVGTGTHIILGLIREGEGVAARVLNNLGVSLNKARQQVLQL.....	141
VKVKRSTLRFSPRAFEVMTAASFLAIHNNSEAVGTEHLLYALLQVEDGFGQLKL.....QKINIVSLRKLEKR	143
QDETFTFLPFSRRLLQVLFDAAEYVASVHVAKVLGTEHVLVYAILHDGNALATRIERAGFSYEDKKDQVKIAALRRNLEER	150
QDETFTFLPFSRRLLQILFDAAEYVASVHVAKVLGTEHVLVYAILHDGNALATRIERAGFSYEDKKDQVKIAALRRNLEER	150
QDETFTFLPFSRRLLQILFDAAEYVASVHVAKVLGTEHVLVYAILHDGNALATRIERAGFSYEDKKDQVKIAALRRNLEER	150
N..DIMILEQSHALKRTLAEEAAISQVTHAKEVGTGTHVLVYAILHDGNALATRIELVGFHAKDDGESIRLLDLRKVITERY	154
ESDIIDLRAQSPALEAMLEAQEIASVTGAVEVGSHEVLMAFLLHKDLMVCRLLLEVAGFYQKDDSDKPRIIDLRRSLERN	156
ESDIIDLRAQSPALEAMLEAQGIASVTGAVEVGSHEVLMAFLLHKDLMVCRLLLEVAGFYQKDDSDKPRIIDLRRSLERN	156

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NTDB id 289 SPD RS10700 WP 001109677.1
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NTDB id 329 STU RS10020 WP 011225298.1
consensus

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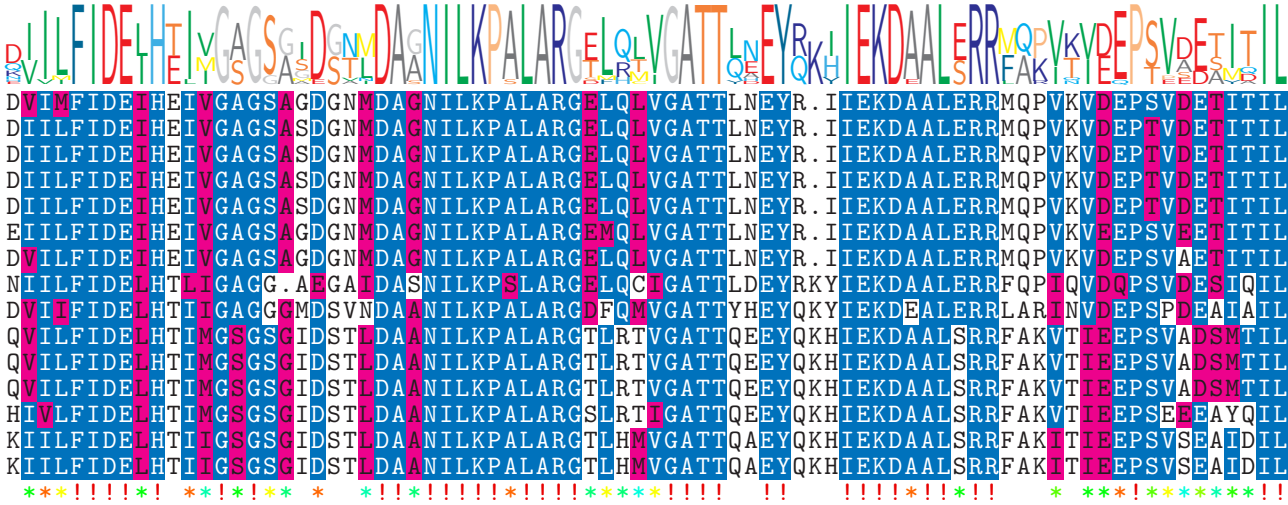
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consensus

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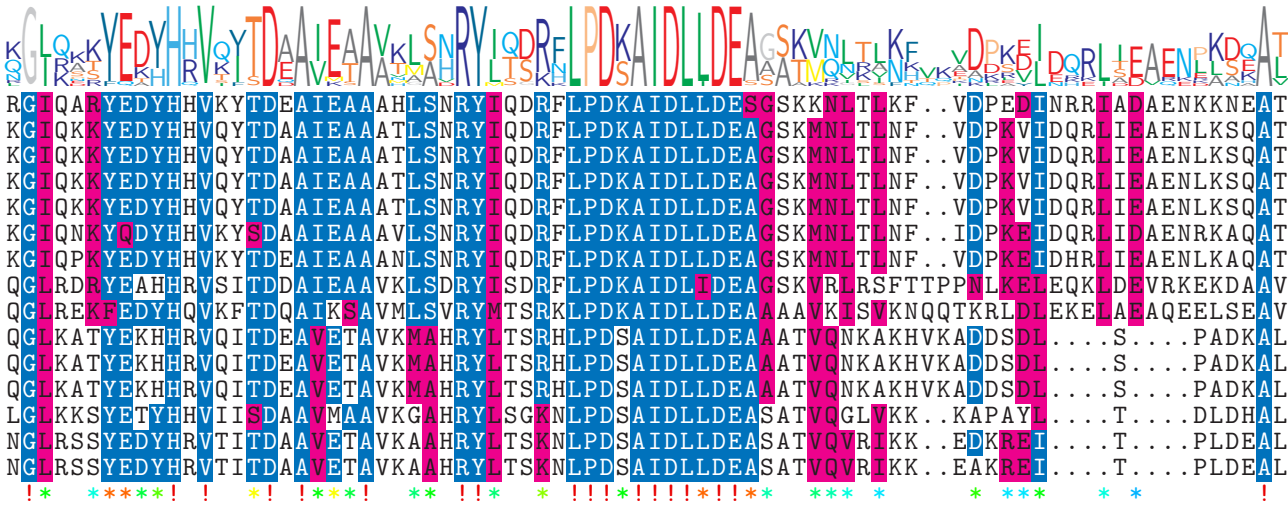
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NTDB id	111597	HAS68	RS03505	WP	009909464.1
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consensus					



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logo

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



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consensus

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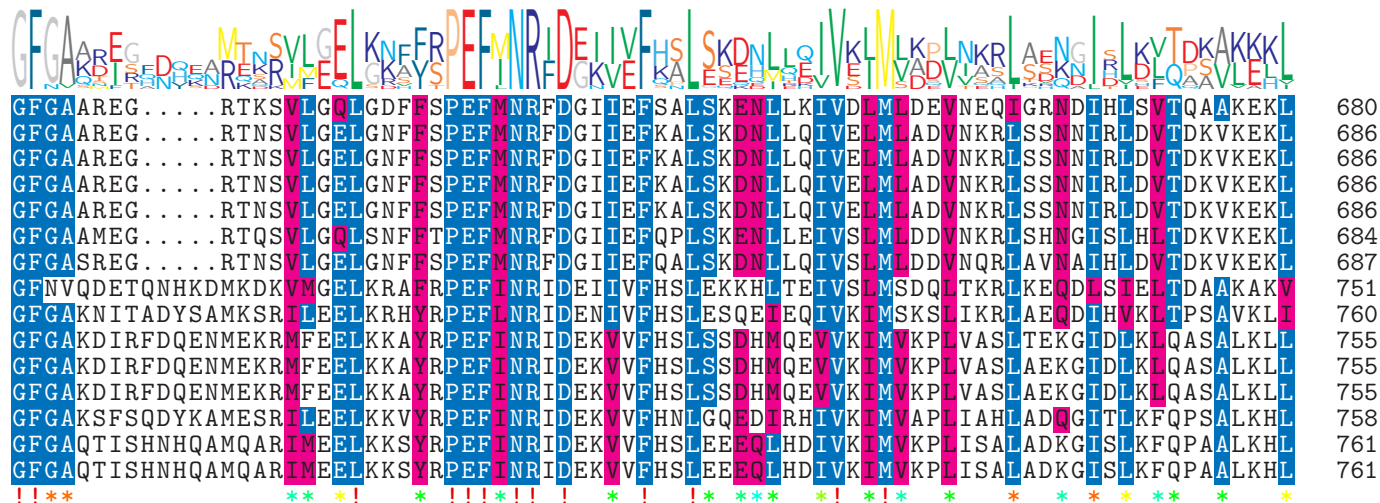
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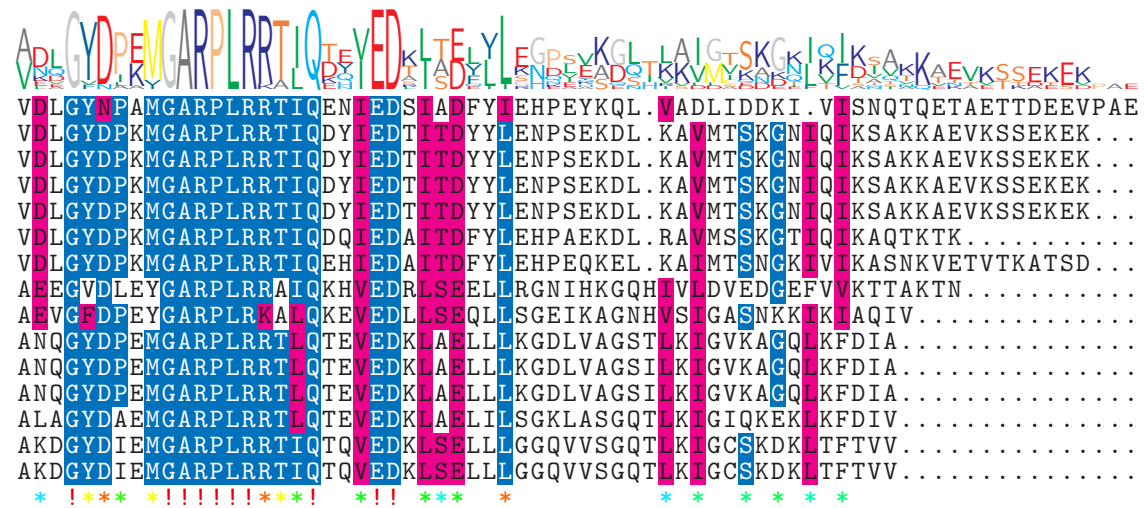
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	non conserved
	similar
	$\geq 50\%$ conserved