

logo

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NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
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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...TIHLYTSVNGQKKQIDLCQNCYQI...MKS	GGREALFGAGNASNGNSDEPF.....	NPFNDIFSALQG.	71
...TIHLYTNLNGKQKKIDLCQNCYKI...	IKTDPNNSLFKGMTDLNRR..DF.....	DPFGDFFNDLNNF	69
...TIHLYTNLNGKQKKIDLCQNCYKI...	IKTDPNNSLFKGMTDLNRR..DF.....	DPFGDFFNDLNNF	69
...TIHLYTNLNGKQKKIDLCQNCYKI...	IKTDPNNSLFKGMTDLNRR..DF.....	DPFGDFFNDLNNF	69
...TIHLYTNVNGKQKQVDLCQNCYQI...	IKTDPNNSLFSGLNHVSHAPGGIN.....	PFFDDFFGDLNNF	72
...TIHLYANVNGQKKQIDLCQNCYQI...	MKTDPENGPLNHLSSQ..QNSSSIN.....	PFFDDFFGDLNNF	70
GQEMSQTIHYTPRAKKVIELSMDEARKLGH	SYVGTEHILLGLIREGEGVAARVLNNL	GVSLNKARQQVLQL.....	141
VKVKRSTLRFSPRAEEVMTAASFLAIHNS	SEAVGTEHLLYALLQVEDGFGQLKL.....	QKINIVSLRKELEKR	143
QDETFTPELPSRRLQILFDEAEYVASV	VHAKVLGTEHVLAILHDGNALATRIL	ERAGFSYEDKKDQVKIAALRRNLEER	150
QDETFTPELPSRRLQILFDEAEYVASV	VHAKVLGTEHVLAILHDGNALATRIL	ERAGFSYEDKKDQVKIAALRRNLEER	150
QDETFTPELPSRRLQVLFDEAEYVASV	VHAKVLGTEHVLAILHDSNALATRIL	ERAGFSYEDKKDQVKIAALRRNLEER	150
N..DITMLEQSHALKRTLAEAAAI	SVTVHAKVGEHVLFAMLLNPNLLATRIL	ELVGFHAKDDGESIRLLDLRKVITERY	154
ESDIIDLRAQSPALEAMLQEAQEIAS	VTGAVEVGSHEVLMAFLLHKDLMVCR	LLEVAGFQYKDDSDKPRIIDLRRSLERN	156
ESDIIDLRAQSPALEAMLQEAQIAS	VTGAVEVGSHEVLMAFLLHKDLMVCR	LLEVAGFQYKDDSDKPRIIDLRRSLERN	156



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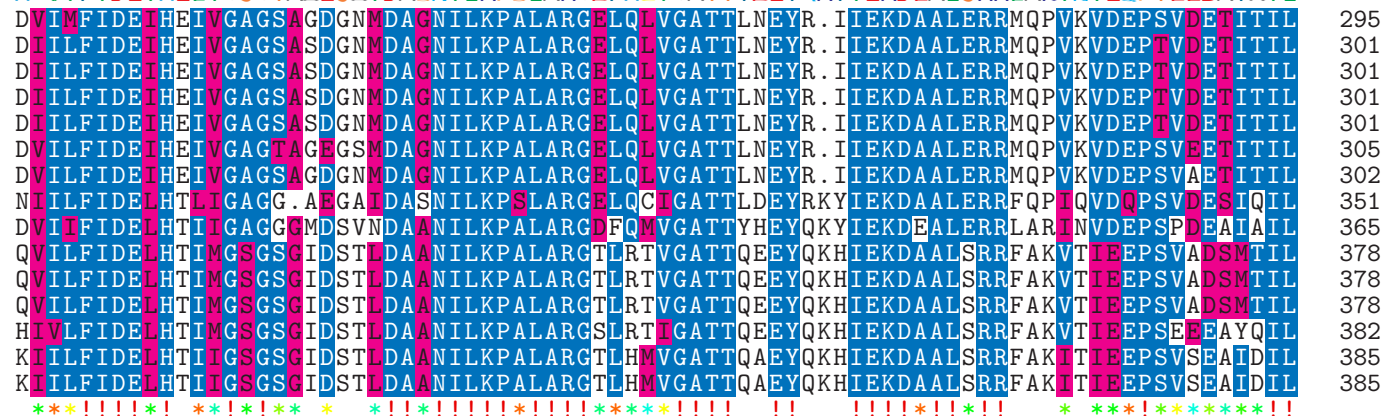
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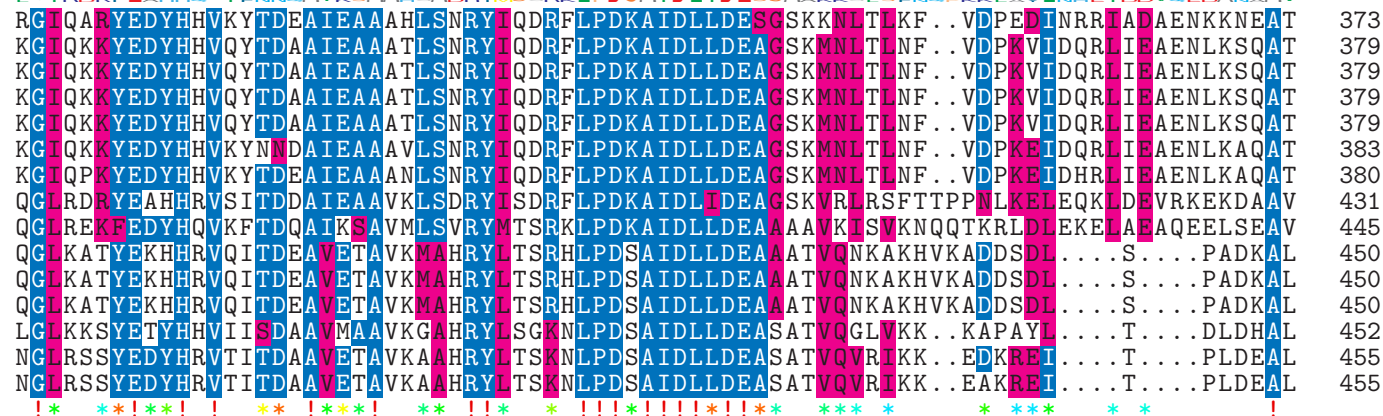
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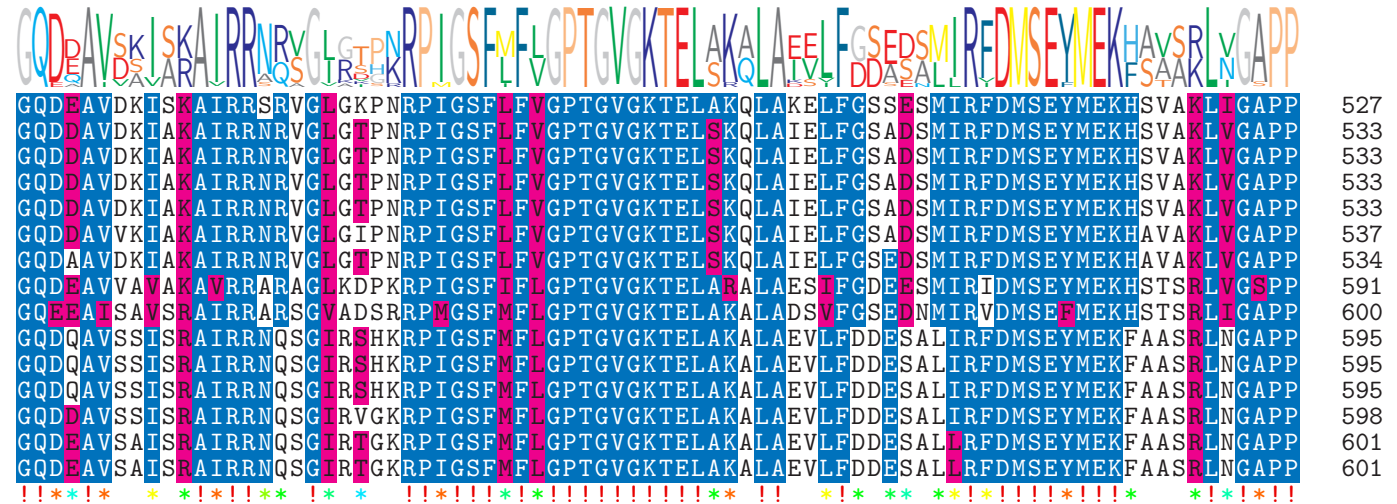
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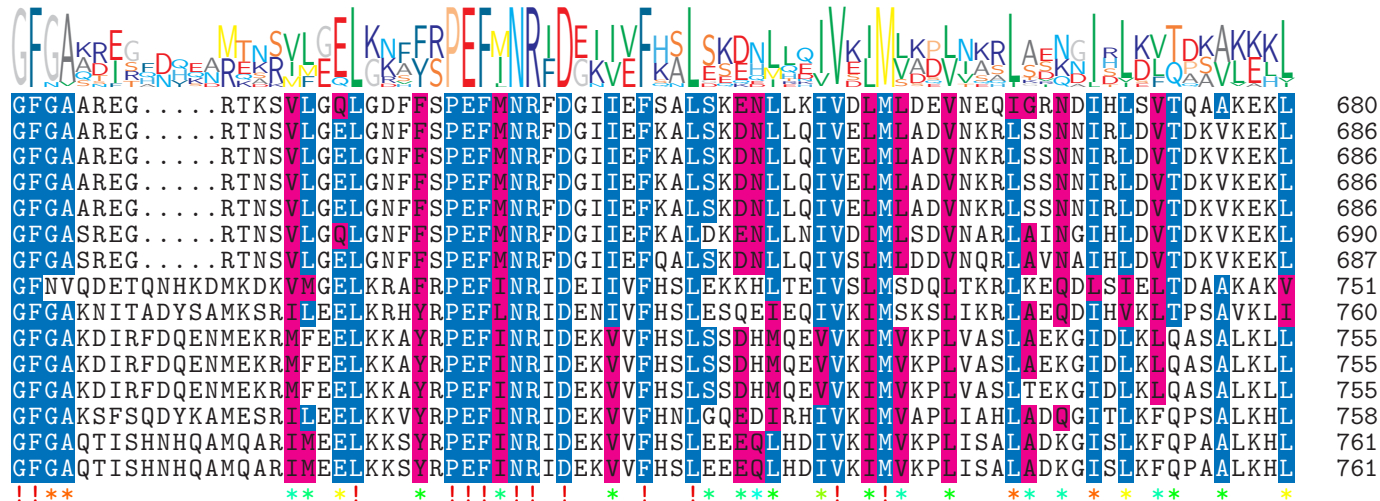
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logo



 non conserved
 similar
 ≥ 50% conserved