

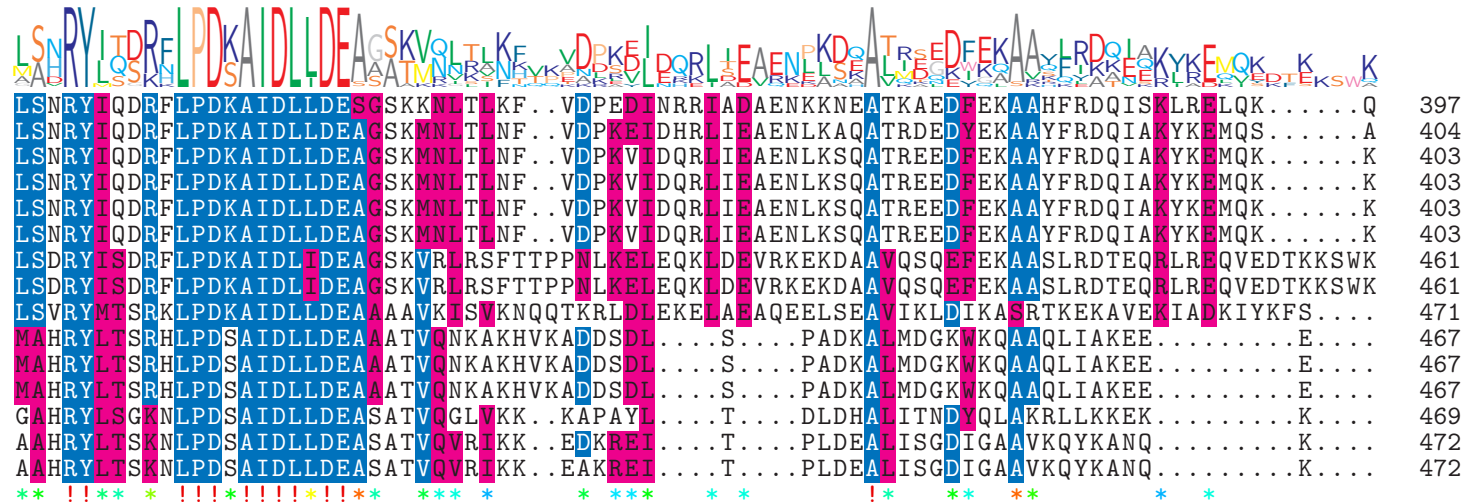
Accession	Species	Gene	Protein	Consensus
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NTDB id 243 SPR RS03635 WP 000882517.1	...	...	MLCQNCNINDS	TI 13
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NTDB id 85 BSU 00860 NP 387967.1	..MMFGR	FTIERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGLAAKALQALGLGSEKIQKEVESLIG.....RGQEMS		75
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NTDB id 289 SPD RS10700 WP 001109677.1	MN	YSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	F	75
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NTDB id 327 STU RS10020 WP 011225298.1	...MTI	YSRKMQAIFHRAQLEAERFESPFLTEWHVLLAMVEVPGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLSES	DI	81
consensus	***	* * *	* * * * *	**

NTDB	id	601	KW2	RS02615	WP	021036805.1	HLYTSVNGQKKQIDLCQNCYQI...MKS	GGQEA	FGAGNAS	NGNSDE		PFNP	FND	DIFSAL	QGGQ	DFNGAASN.	80
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NTDB	id	281	KZH43	RS03550	WP	000882517.1	HLYTNLNGQKKQIDLCQNCYKI...IKTDPNNS	LFKGMTDL	NNRD		FDP	FGDFFND	LNNFRPS	N..TP.			76
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NTDB	id	329	STER	RS00545	WP	011680614.1	DLRAQSPAELAMLQEAQEIASVTGAVEV	GSEHVLMAFL	LHKDLMVCRLL	EVAGFQYKDDSDKPRIIDLRRSL	ERNAGLSKQDLKA						166
NTDB	id	327	STU	RS10020	WP	011225298.1	DLRAQSPAELAMLQEAQGIASVTGAVEV	GSEHVLMAFL	LHKDLMVCRLL	EVAGFQYKDDSDKPRIIDLRRSL	ERNAGLSKQDLKA						166
consensus							* * * * *	*	*****	* * *	*	*		*	*	*	


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NTDB id 610 V4T04 RS10165 WP 012897346.1
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NTDB id 329 STU RS10020 WP 011225298.1
consensus

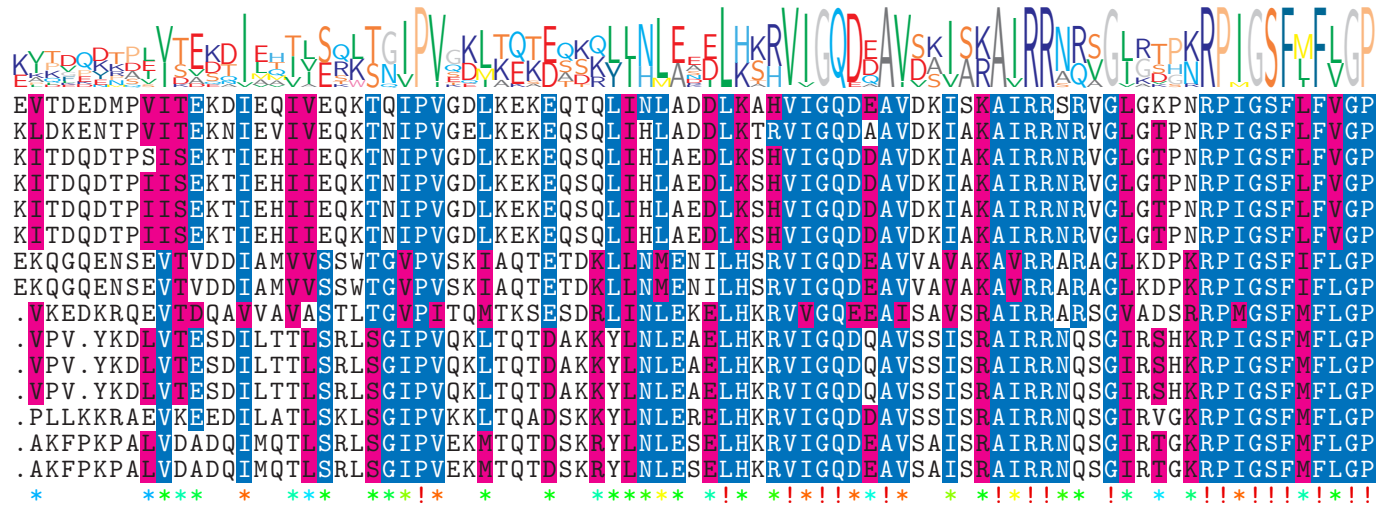
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NTDB id 329 STU RS10020 WP 011225298.1
consensus

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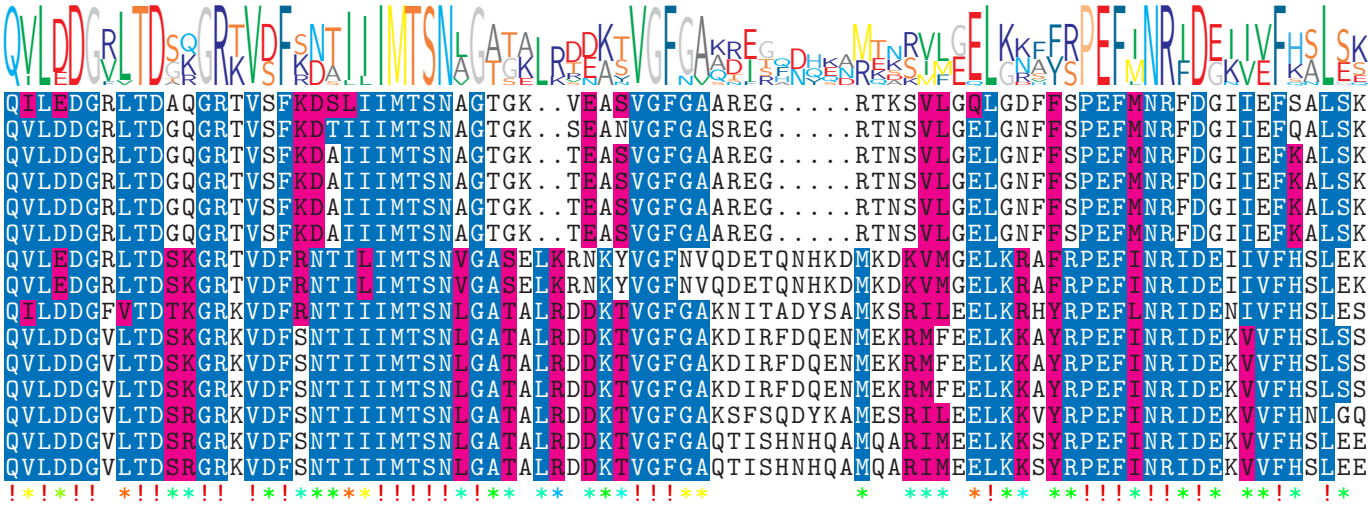
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consensus

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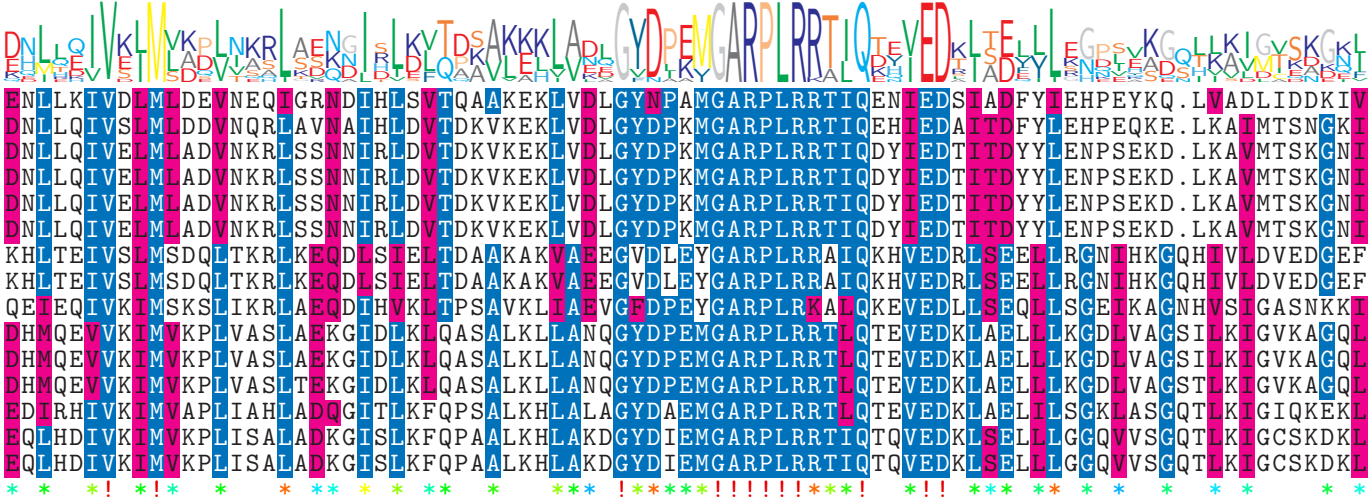
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NTDB	id	297	STER	RS00545	WP	011680614.1
NTDB	id	329	STU	RS10020	WP	011225298.1
consensus						



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NTDB	id	601	KW2	RS02615	WP	021036805.1
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NTDB	id	329	STU	RS10020	WP	011225298.1
consensus						



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consensus		



-  non conserved
-  similar
-  ≥ 50% conserved