

logo

NTDB id	Accession	Sequence	Position
NTDB id 601	KW2 RS02615 WP 021036805.1	MLCQNCNINEAT	13
NTDB id 281	KZH43 RS03550 WP 000882517.1	MLCQNCNKINDST	13
NTDB id 245	SPD RS03860 WP 000882517.1	MLCQNCNKINDST	13
NTDB id 243	SPR RS03635 WP 000882517.1	MLCQNCNKINDST	13
NTDB id 241	SP RS04015 WP 000882523.1	MLCQNCNKINDST	13
NTDB id 31365	SEZ RS03075 WP 012515248.1	MLCQNCNINEAT	13
NTDB id 378	SMU RS02690 WP 002263569.1	MLCQNCNINEAT	13
NTDB id 85	BSU 00860 NP 387967.1	MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGGIAAKALQALGLGSEKIQKEVESLIG.....RGQEMS	75
NTDB id 610	V4T04 RS10165 WP 012897346.1	MKFENIKYPTPTLDRILEKAEYYAHQYQYGTIESAHLAAMATTSGSIAYSLLAGMNVDSDDLIDL.....EDLSSHVKVKR	77
NTDB id 290	KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 289	SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 287	SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 377	SMU RS09275 WP 002262344.1	...MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFADVRVFEDYQAAAILAIGKKPKSSSN..DI	79
NTDB id 297	STER RS00545 WP 011680614.1	...MTIYSRKMQAIFHRAQLEAERFESPFLFETWHVLLAMVEVTGSAVYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDI	81
NTDB id 329	STU RS10020 WP 011225298.1	...MTIYSRKMQAIFHRAQLEAERFESPFLFETWHVLLAMVEVPGSAVYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDI	81
consensus		MLCQNCNINEAT	

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NTDB id 601 KW2 RS02615 WP 021036805.1	HL YTSVNGQKKQIDLCQNCYQI...MKS GQGEALFGAGNASNGNS.....DEPFNP.FNDIFSALQGGQ.DFNGAAS..	79
NTDB id 281 KZH43 RS03550 WP 000882517.1	HL YTNLNGKQKKQIDLCQNCYKI...IKTDPNNSLFGKGMTDLN.....NRDF.DPFGDFFNDLNNFRPSS..NT..	75
NTDB id 245 SPD RS03860 WP 000882517.1	HL YTNLNGKQKKQIDLCQNCYKI...IKTDPNNSLFGKGMTDLN.....NRDF.DPFGDFFNDLNNFRPSS..NT..	75
NTDB id 243 SPR RS03635 WP 000882517.1	HL YTNLNGKQKKQIDLCQNCYKI...IKTDPNNSLFGKGMTDLN.....NRDF.DPFGDFFNDLNNFRPSS..NT..	75
NTDB id 241 SP RS04015 WP 000882523.1	HL YTNLNGKQKKQIDLCQNCYKI...IKTDPNNSLFGKGMTDLN.....NRDF.DPFGDFFNDLNNFRPSS..NT..	75
NTDB id 31365 SEZ RS03075 WP 012515248.1	HL YTSINGKQRQIDLCQNCYQI...MKTDPKNPILGGYNGNQKAQ.....QPSQNPFFDDFFGDLNNFRAFG..QL..	79
NTDB id 378 SMU RS02690 WP 002263569.1	HL YANVNGQKKQIDLCQNCYQI...MKTDPENGPLNHL...SQQN.....SSSINPFFDDFFGDLNNFRAFNNQDL..	78
NTDB id 85 BSU 00860 NP 387967.1	QTIHYTPRAKKVIELSMDEARKLGHSYVGT EHI LLGLIREGEGVAARVLNNLGVSLNKARQQVLQL.....LGSNET....	147
NTDB id 610 V4T04 RS10165 WP 012897346.1	STLRFSPRAEVEVMTAASFLAIHNSEAVGT EHL L YALLQVEDGFGQLLKL.....QKINIVSLRKLEKRTGLKVPESKK	153
NTDB id 290 KZH43 RS10025 WP 001109677.1	TELPFSRRLQILFDEAEYVASVVHAKVLGT EHV L YAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 289 SPD RS10700 WP 001109677.1	TELPFSRRLQILFDEAEYVASVVHAKVLGT EHV L YAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 287 SP RS11210 WP 001109712.1	TELPFSRRLQVLFDEAEYVASVVHAKVLGT EHV L YAILHDSNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 377 SMU RS09275 WP 002262344.1	MLLEQSHALKRTLAEAAAISQVTHAKEVGT EHV L FAML LNPNLLATRILELVGFHAKDDGESIRLLDLRKV IERYAGFSKEDIKA	164
NTDB id 297 STER RS00545 WP 011680614.1	DLRAQSPALEAMLQEAQEIASVTGAVEVGEHVLMFAFLHKKDLMVCRLLLEVAGFQYKDDSDKPRIIDLRRSLERNAGLSKQDLKA	166
NTDB id 329 STU RS10020 WP 011225298.1	DLRAQSPALEAMLQEAQGIASVTGAVEVGEHVLMFAFLHKKDLMVCRLLLEVAGFQYKDDSDKPRIIDLRRSLERNAGLSKQDLKA	166
consensus	* * * * *	

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NTDB id 297 STER RS00545 WP 011680614.1
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consensus

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.IHDLRKKKSASA.....NFANMMQPQSSTGELADYTKDLTALAESGNLDPVIIGRDEEISRMIVQLSRKTNNNPVLV 240

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consensus

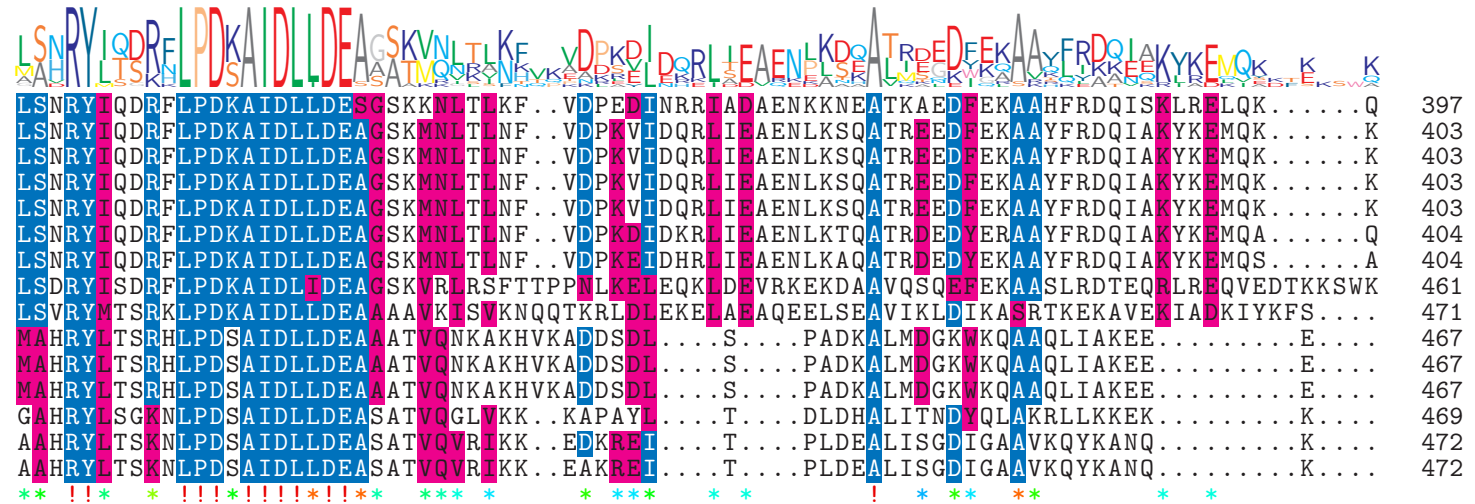
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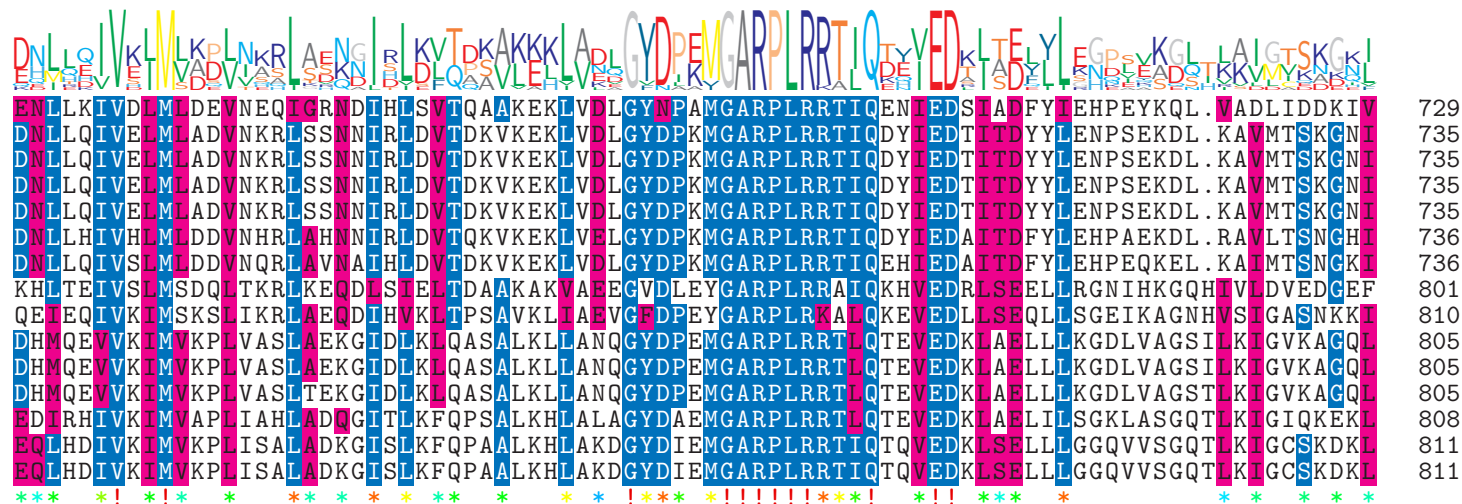
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NTDB id 329 STU RS10020 WP 011225298.1	TFTVV.....	816
consensus		

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