

logo

NTDB id 601 KW2 RS02615 WP 021036805.1	M.CQNCNINEAT.....I	13
NTDB id 281 KZH43 RS03550 WP 000882517.1	M.CQNCKINDST.....I	13
NTDB id 245 SPD RS03860 WP 000882517.1	M.CQNCKINDST.....I	13
NTDB id 243 SPR RS03635 WP 000882517.1	M.CQNCKINDST.....I	13
NTDB id 241 SP RS04015 WP 000882523.1	M.CQNCKINDST.....I	13
NTDB id 54312 SDSE RS18105 WP 015057878.1	M.CQNCNINEST.....I	13
NTDB id 378 SMU RS02690 WP 002263569.1	M.CQNCKINEAT.....I	13
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTEAQQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLIG.....RGQEMS	75
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYPTLDRILEKAEYYAHQYQYGTIESAHLLEAAATTSGSLAYSLLAGMNVDSDDLIDL.....EDLSSHVKVKR	77
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	75
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	75
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	75
NTDB id 377 SMU RS09275 WP 002262344.1	...MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFADVRVFEDYQAAAILAIGKKPKSSSN..DI	79
NTDB id 297 STER RS00545 WP 011680614.1	...MTIYSRKMQAIFHRAQLEAERFESPFLFETWHVLLAMVEVTGSAVYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLSESDI	81
NTDB id 329 STU RS10020 WP 011225298.1	...MTIYSRKMQAIFHRAQLEAERFESPFLFETWHVLLAMVEVPGSAVYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLSESDI	81
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NTDB id 601 KW2 RS02615 WP 021036805.1	HLIYTSVNGCQKQIDLCQNCYQI...MKS...EALFGAGNASNGNSD.....EPFNP.FNDIFSALQGGQ.DFNGAASNQ	81
NTDB id 281 KZH43 RS03550 WP 000882517.1	HLIYTNLNGKQKQIDLCQNCYKI...KTDPNNSLEKGMTDL...NN.....RDFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 245 SPD RS03860 WP 000882517.1	HLIYTNLNGKQKQIDLCQNCYKI...KTDPNNSLEKGMTDL...NN.....RDFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 243 SPR RS03635 WP 000882517.1	HLIYTNLNGKQKQIDLCQNCYKI...KTDPNNSLEKGMTDL...NN.....RDFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 241 SP RS04015 WP 000882523.1	HLIYTNLNGKQKQIDLCQNCYKI...KTDPNNSLEKGMTDL...NN.....RDFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 54312 SDSE RS18105 WP 015057878.1	HLIYTNVNGKQKQIDLCQNCYQI...MKTDPNPLGLGNGGQRAQE.....RSSNPFFDDFFGD LNNFRAFGN..LPN	81
NTDB id 378 SMU RS02690 WP 002263569.1	HLIYANVNGCQKQIDLCQNCYQI...MKTDPENGPLNHL...SQQNS.....SSINPFFDDFFGD LNNFRAFNNQDLPN	80
NTDB id 85 BSU 00860 NP 387967.1	QTIHYTPRAKKVIELSMDEARKLGHSYVCTEHLILGLIREGEGVAARVLNNLGVSLNKARQQVLQL.....LCSNET....	147
NTDB id 610 V4T04 RS10165 WP 012897346.1	STLRFSPRAEIVMTAASF LAIHNNSEAVCTEHLIYALIQVEDGFGLQLLKL.....QKINIVSLRKELEKRTGLKVPESSK	153
NTDB id 290 KZH43 RS10025 WP 001109677.1	TELPFSRRLQILFDEAEYVASV VHAKVLGTEHVLYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 289 SPD RS10700 WP 001109677.1	TELPFSRRLQILFDEAEYVASV VHAKVLGTEHVLYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 287 SP RS11210 WP 001109712.1	TELPFSRRLQVLFDFAEYVASV VHAKVLGTEHVLYAILHDSNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 377 SMU RS09275 WP 002262344.1	MILEQSHALKITLAEAAAISQVTHAKEVCTEHLVLFAMLLNPNLLATRILELVGFHAKDDGESIRLLDLRKVTERYACFSKEDIKA	164
NTDB id 297 STER RS00545 WP 011680614.1	DIRAQSPALEAMIQEAQGIASVTGAVEVGEHVLMAFLLHKDLMVCRLLLEVAGFYKDDSDKPRIIDLRRSLERNAGLSKQDLKA	166
NTDB id 329 STU RS10020 WP 011225298.1	DIRAQSPALEAMIQEAQGIASVTGAVEVGEHVLMAFLLHKDLMVCRLLLEVAGFYKDDSDKPRIIDLRRSLERNAGLSKQDLKA	166
consensus	* * * * * * * * * * * * * * *	

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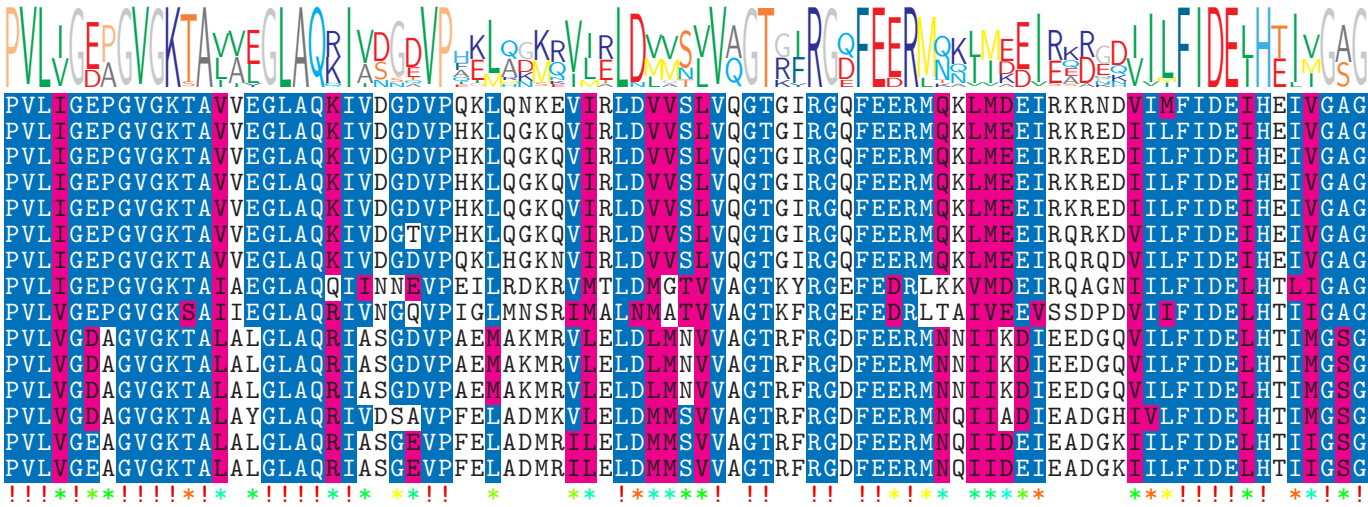
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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus



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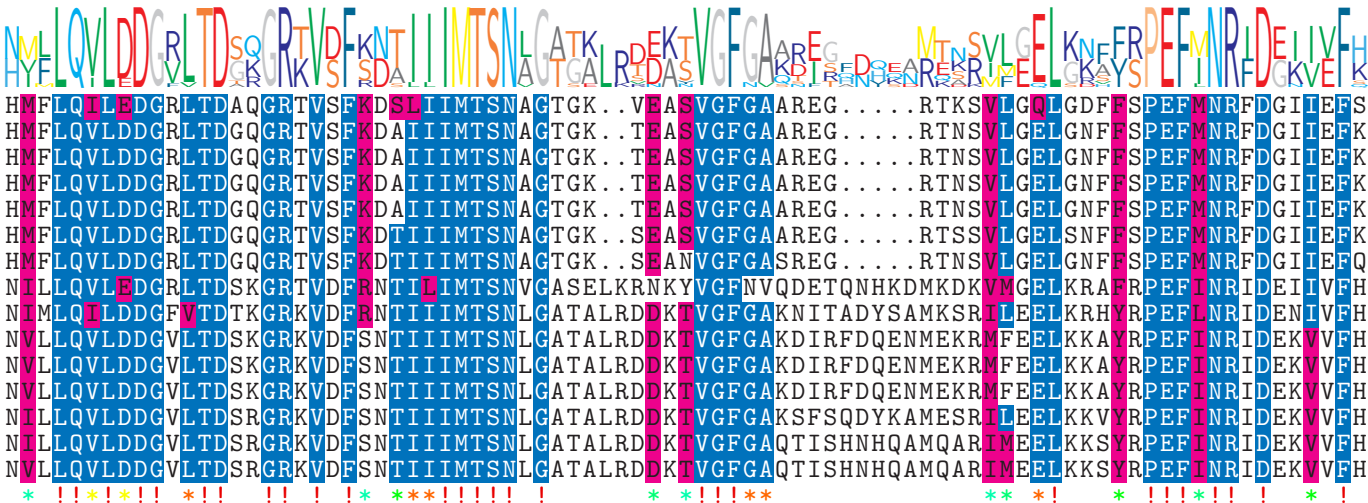
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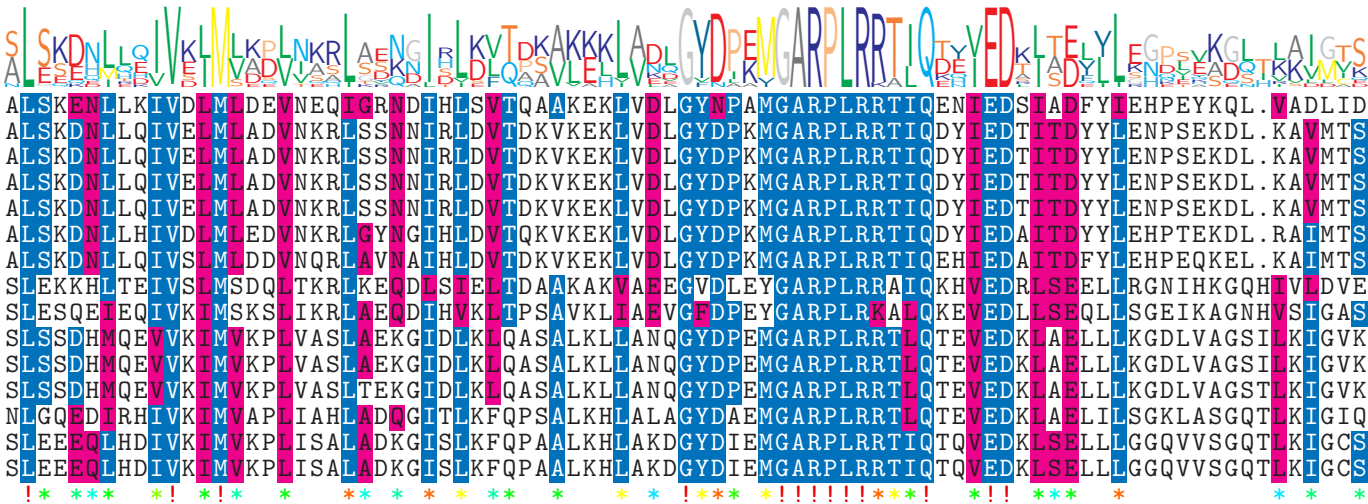
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NTDB id 601 KW2 RS02615 WP 021036805.1
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consensus



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logo

NTDB id 601 KW2 RS02615 WP 021036805.1	DKI	VISNQTQE.TAETTDDEEVP	AE.	748
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NTDB id 243 SPR RS03635 WP 000882517.1	K	NIQIKSAKKA	AEVKSSEKEK....	752
NTDB id 241 SP RS04015 WP 000882523.1	K	NIQIKSAKKA	AEVKSSEKEK....	752
NTDB id 54312 SDSE RS18105 WP 015057878.1	NSN	IMIKAAKKDET	VTAKQDKTDEN	763
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NTDB id 85 BSU 00860 NP 387967.1	D	GEFVVKTTAKTN		810
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NTDB id 287 SP RS11210 WP 001109712.1	A	QIKFDIA		810
NTDB id 377 SMU RS09275 WP 002262344.1	K	EIKFDIV		813
NTDB id 297 STER RS00545 WP 011680614.1	K	DKLTFTVV		816
NTDB id 329 STU RS10020 WP 011225298.1	K	DKLTFTVV		816
consensus	*	*		

- non conserved
- similar
- ≥ 50% conserved