

logo

NTDB id 601 KW2 RS02615 WP 021036805.1	MTCQNCMINEAT.....	12
NTDB id 281 KZH43 RS03550 WP 000882517.1	MTCQNCKINDST.....	12
NTDB id 245 SPD RS03860 WP 000882517.1	MTCQNCKINDST.....	12
NTDB id 243 SPR RS03635 WP 000882517.1	MTCQNCKINDST.....	12
NTDB id 241 SP RS04015 WP 000882523.1	MTCQNCKINDST.....	12
NTDB id 1140102 DQM37 RS05955 WP 111694893.1	MTCQNCNINEST.....	12
NTDB id 378 SMU RS02690 WP 002263569.1	MTCQNCKINEAT.....	12
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NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYTPTLDRILEKAEYYAHQYQYGTIESAHLLEAMATTSGSLAYSLLAGMNVDSDDLIDL.....EDLSSH	72
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYS	70
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYS	70
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYS	70
NTDB id 377 SMU RS09275 WP 002262344.1	MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFEADVRFEDYQAAAAILAIGKKPKSSS	76
NTDB id 297 STER RS00545 WP 011680614.1	...MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVTGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
NTDB id 329 STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVPGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
consensus	**.....*	

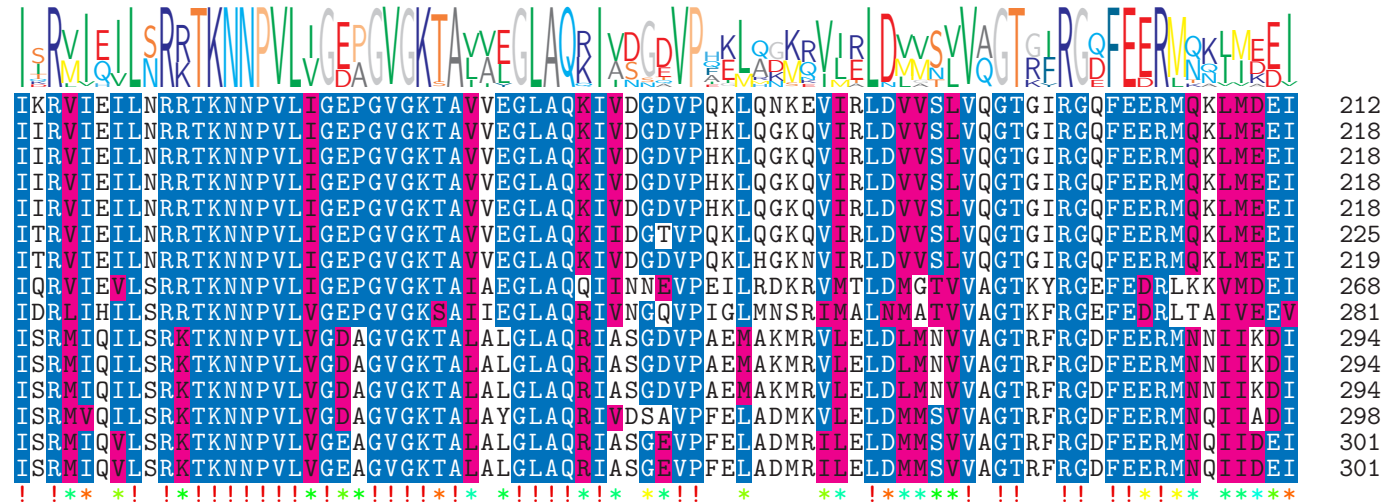
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NTDB id 281 KZH43 RS03550 WP 000882517.1	IHL YTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDLN.....NRDF.DPFGDFFNDLNNF	69
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NTDB id 378 SMU RS02690 WP 002263569.1	IHL YANVNGQKQKQIDLCQNCYQI...MKTDPENGPLNHLSSQQ...N.....SSSINPFFDDFFGD LNNF	70
NTDB id 85 BSU 00860 NP 387967.1	GQEMSQTIHYTPRAKKVIELSMDEARKLGHSYVCTEHLILGLIREGEGVAARVLNNLGVSLNKARQQVLQL.....	141
NTDB id 610 V4T04 RS10165 WP 012897346.1	VKVKRSTLRFSPRAEVMTAASFLAIHNNSEAVCTEHLIYALLQVEDGFGQLQLLKL.....QKINIVSLRKETEKR	143
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consensus	* * * * * * * * * * * * * *	

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NTDB id 329 STU RS10020 WP 011225298.1
consensus

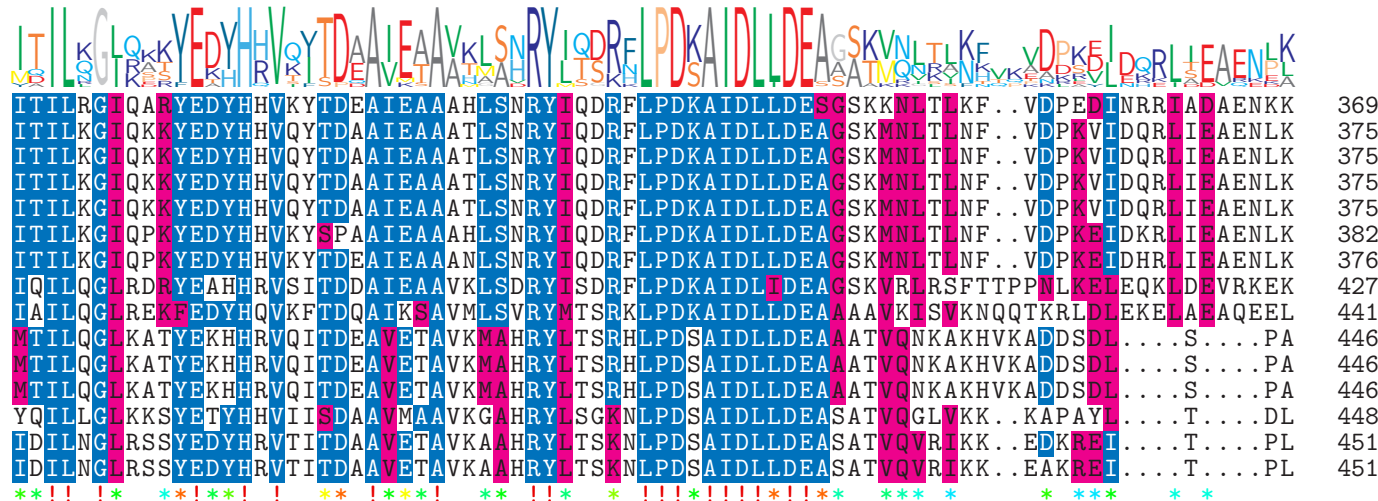
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consensus

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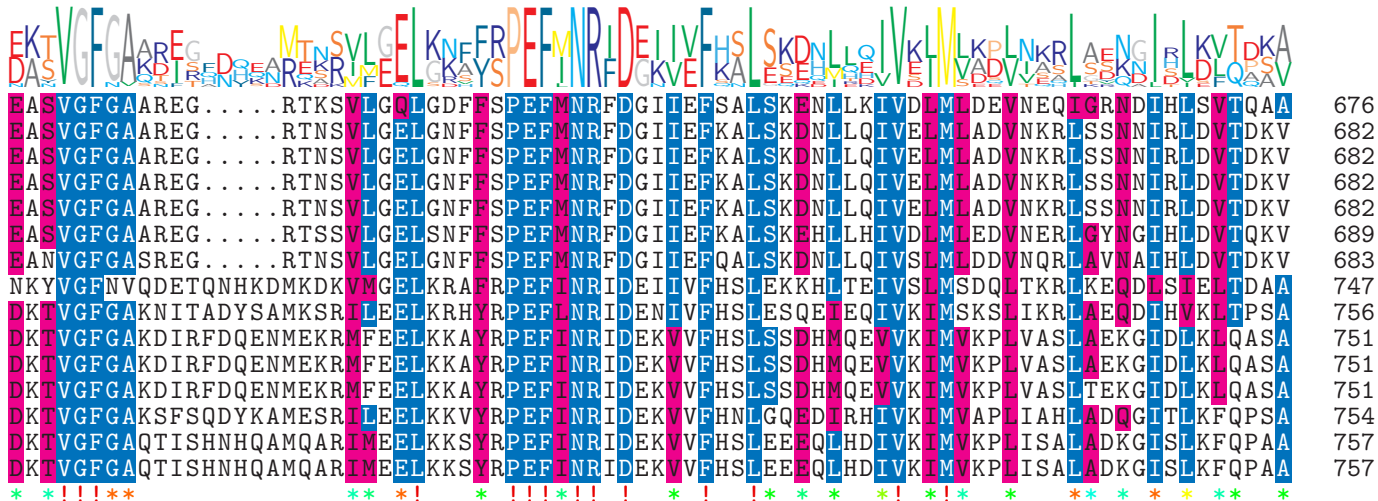
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NTDB	id	245	SPD	RS03860	WP	000882517.1	SQATREEDFEKAAAYFRDQIAKYKEMQK.....KKITDQDTPIISEKTIEHIIIEQKTNIPVGD LKEKEEQQSLIHLAEDLK	449
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NTDB	id	610	V4T04	RS10165	WP	012897346.1	SEAVIKLDIKASRTKEKAVEKAIADKIYKFS....VKEDKRQEVTDAQVA VASTLTGVPITQMTKS ESDRLINLEKELH	516
NTDB	id	290	KZH43	RS10025	WP	001109677.1	DKALMDGKW KQAAQLIAKEE.....E....VPV.YKDLVTESD ILTTLSRLSGIPVQKLTQTD AKKYLNLEAEHL	511
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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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NTDB id 241 SP RS04015 WP 000882523.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDTTIDYYLENPSEKDLKAVMTSKCNIIQIKSAKKAIEVKSSEKEK	752
NTDB id 1140102 DQM37 RS05955 WP 111694893.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDAITDYYLEHPTEKQLRALMTNSENITIKAVKEGDSFLNEESL	760
NTDB id 378 SMU RS02690 WP 002263569.1	KEKLVLDLGYDPKMGARPLRRTIQEHTIEDAITDFYLEHPEQKELKAIMTSNCKIVIKASNKVETVTKATSD	753
NTDB id 85 BSU 00860 NP 387967.1	KAKVAEEGVLDLEYGARPLRRRIQKHVEDRLSEELLRGNIHKGQHVLVDVEDGEFVVVKTAKTN	810
NTDB id 610 V4T04 RS10165 WP 012897346.1	VKLIAEVCFDPEYGARPLRKALQKEVEDLLSEQLLSGEIKAGNHVSTIGASNKKIKIAQIV	816
NTDB id 290 KZH43 RS10025 WP 001109677.1	LKLLANQGYDPEMGARPLRRTIQTEVEDKLAELLKKGDLVAGSILKIGVKACQLKFDIA	810
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consensus	* * !*** *!!!!!!***! *!! *** *	

- non conserved
- similar
- ≥ 50% conserved