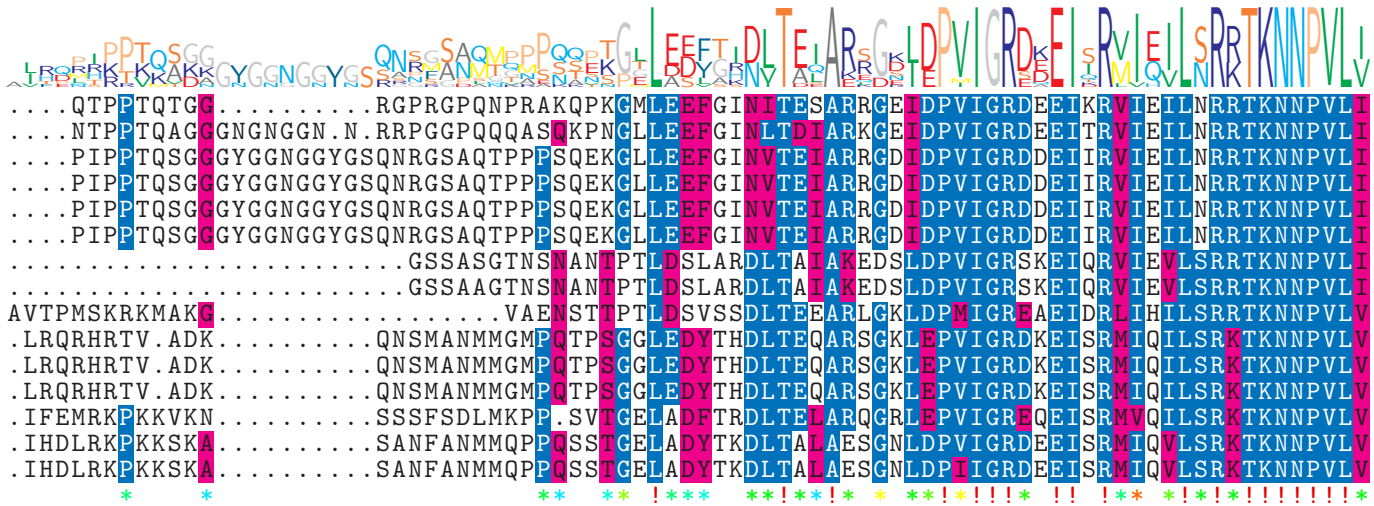


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NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGLAAKALQALGLGSEKIQKEVESLIG.....RGQEMS	75
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NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
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NTDB id 329 STU RS10020 WP 011225298.1	...MTIYSRKMQAIFHRAQLEAERFESPFLETWHVLLAMVEVPGSAVYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLSESDI	81
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NTDB	id	297	STU	RS10020	WP	011225298.1	DlRaQSpaLEAmLQEAQGiasvtGAveVgSEhVlMAFlLHKDLMVCRLLeVAGfQYKDDSDKPRIIdLRRSLERNAGLSKQDLKA	166
consensus							* * * * *	

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NTDB id 329 STU RS10020 WP 011225298.1
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consensus



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NTDB id 85 BSU 00860 NP 387967.1	GAIDASNILKPSLARGEELQCIGATTTLDEYRKYIEKDAALERRFQPIQVDQPSVDESIIQLGLRDRYEAAHHRVSITDDAIEAAVK	376
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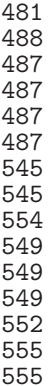
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NTDB id 377 SMU RS09275 WP 002262344.1	GAHRYLSGNLPDSAIDLLDEASATVQGLVKK..KAPAYL....T....DLDHALITNDYQLAKRLLKKEK.....KP...	470
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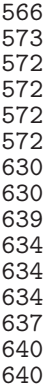
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consensus

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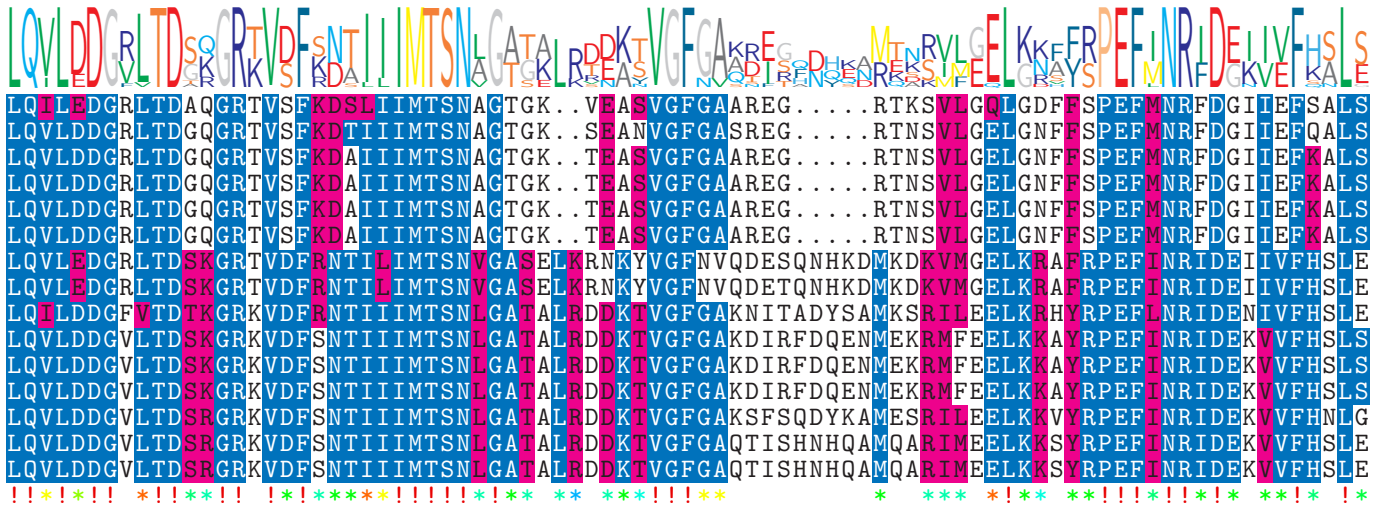


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consensus



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consensus	*																				

- ⓧ non conserved
- ✖ similar
- ⓧ ≥ 50% conserved