

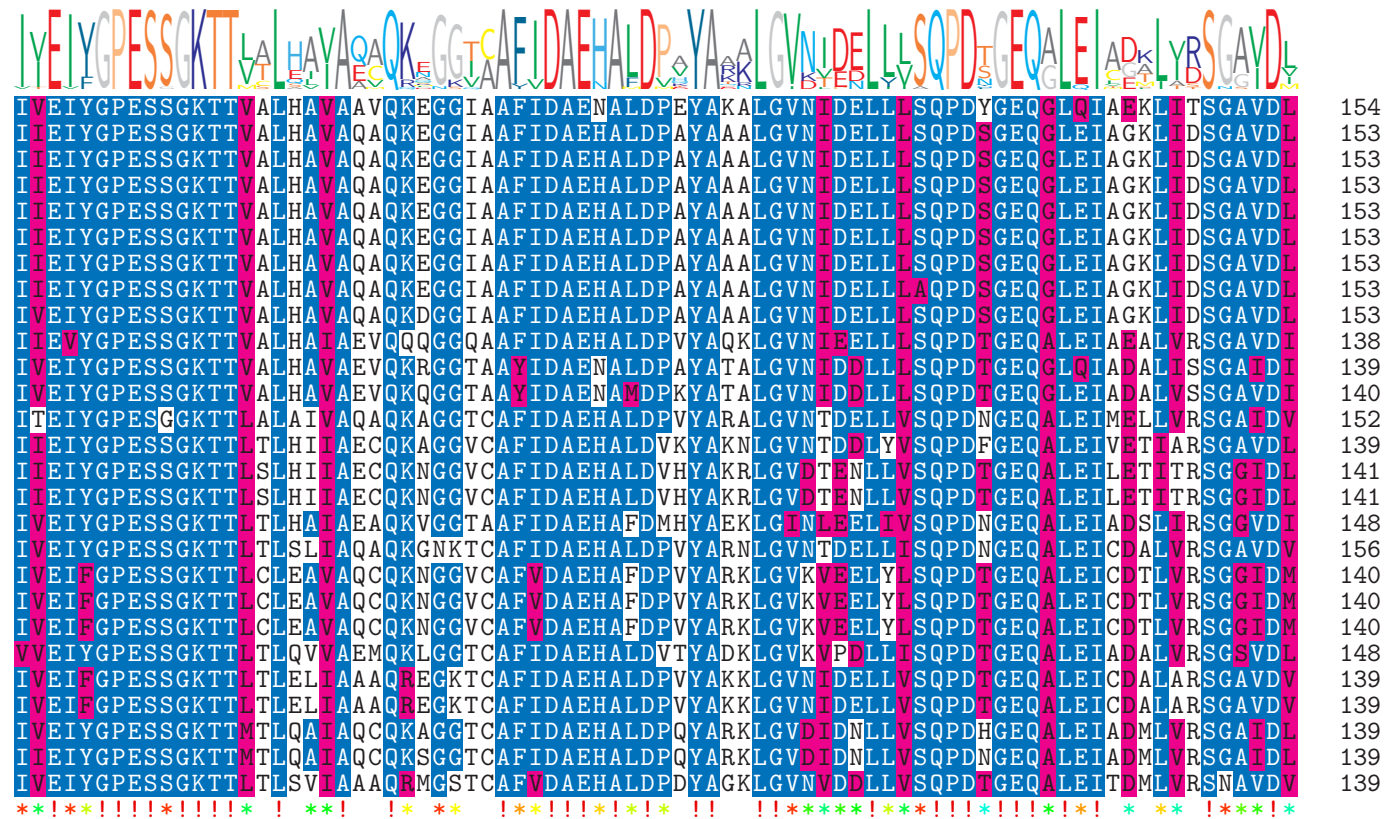
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NTDB id 228 SPD RS09265 WP 001085462.1	.MAK.....KPKKLEEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQV	MSSGSLALD	I	ALGSGGY	P	KGR	73		
NTDB id 194 SPR RS08825 WP 001085462.1	.MAK.....KPKKLEEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQV	MSSGSLALD	I	ALGSGGY	P	KGR	73		
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NTDB id 115 BSU 16940 NP 389576.2	MSDRQAALDMALKQ	IEKQFGKGSIMKLGEK	.TDTRI	STVPSGSLALD	T	ALGIGGY	P	RGR	58			
NTDB id 179158 A6F53 RS05555 WP 011667973.1	MADERQAALDKALKK	IEKD	FGKGSIMRLGDN	.SNLEVET	VPSGSLALD	V	ALGVGGY	P	RGR	59		
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NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTDAK	ERSKAT	IETAMSQIEKA	FGKGSIMKLGA	E	.SKLDVQV	VSTGSLSLD	L	ALGVGGI	P	RGR	72
NTDB id 1245 Cj1673c YP 002345041.1	MDNKRKSLDAALKS	LDKT	FGKGTILRLGDK	.EVEQ	IDSIGTGSVGLDL	ALGIGGV	P	KGR	59			
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTV..VKQVEP	EQQKQ	ALEAALAQIEKQ	FGKGSIMKLQ	QT.EQLD	IESVST	GSLGLDL	ALGIGGL	P	MGR	76	
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NTDB id 600 KW2 RS01765 WP 011675413.1
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NTDB id 499 SM12261 RS08315 WP 078228268.1
NTDB id 453 Spy49 1753c ACI62003.1
NTDB id 408 SMU RS09530 WP 002262392.1
NTDB id 115 BSU 16940 NP 389576.2
NTDB id 179158 A6F53 RS05555 WP 011667973.1
NTDB id 625 LCA RS02525 WP 011374196.1
NTDB id 1313 DR RS12030 WP 010888966.1
NTDB id 1245 Cj1673c YP 002345041.1
NTDB id 1211 C694 RS00795 WP 000963128.1
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NTDB id 1334 RA0C RS04870 WP 004918236.1
NTDB id 1389 A4U84 RS07560 WP 021112712.1
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NTDB id 1124 NGFG RS03960 WP 003688695.1
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NTDB id 1427 RS RS02760 WP 011000510.1
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consensus

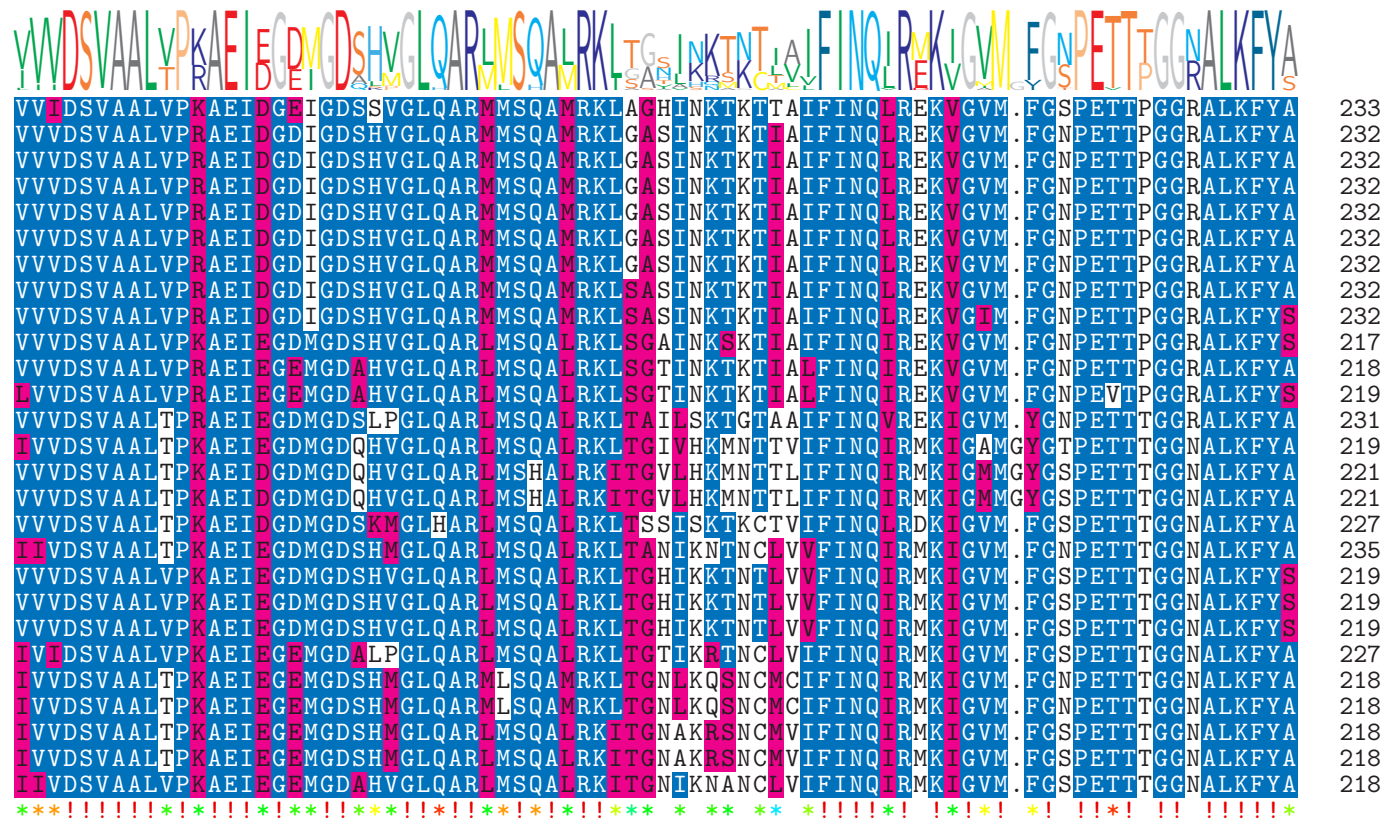
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consensus

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logo

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NTDB id 625 LCA RS02525 WP 011374196.1	TVRLEVRRAETIKNGT.....DMIGNRARIKVVKNKVAPPFKVAEVDIMYGGGISRTGELVDMAVEKDIINKSGSWYSYG	294
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NTDB id 1205 PSJM300 05545 AFN77183.1	SVRLDIRRTGA....V.KEGDEVVGSETRVVVVKNKVAPPFRQAEFQILYGKGIYRNGEITIDLGVQQGLVEKSGAWAYK	293
consensus	*!!**!* * ** *****!!*!!*!!*!!* * *****!!*!!** * ** *!!*!!*!!	

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consensus

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Sequence	Score
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NTDB id 228 SPD RS09265 WP 001085462.1	DELEIEIEE		388
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	S.....		349
NTDB id 1205 PSJM300 05545 AFN77183.1			347
consensus			

-  non conserved
-  similar
-  ≥ 50% conserved