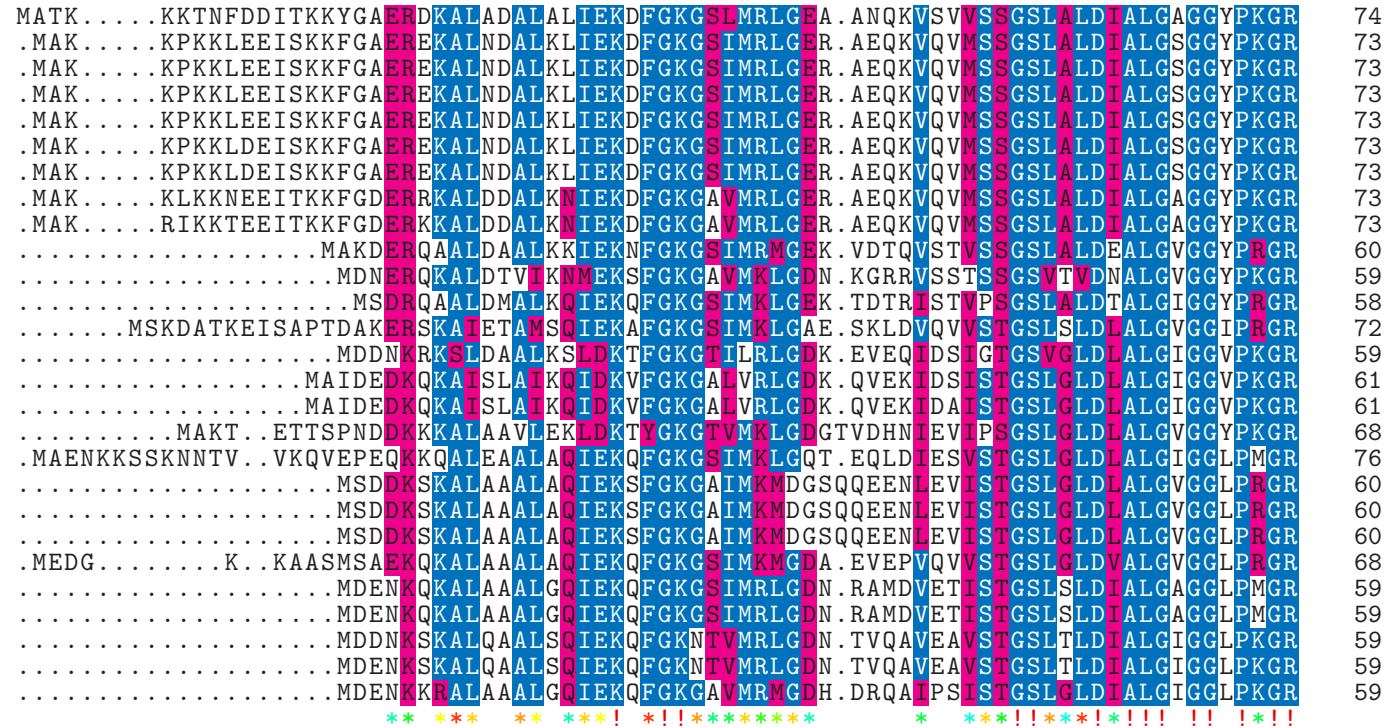


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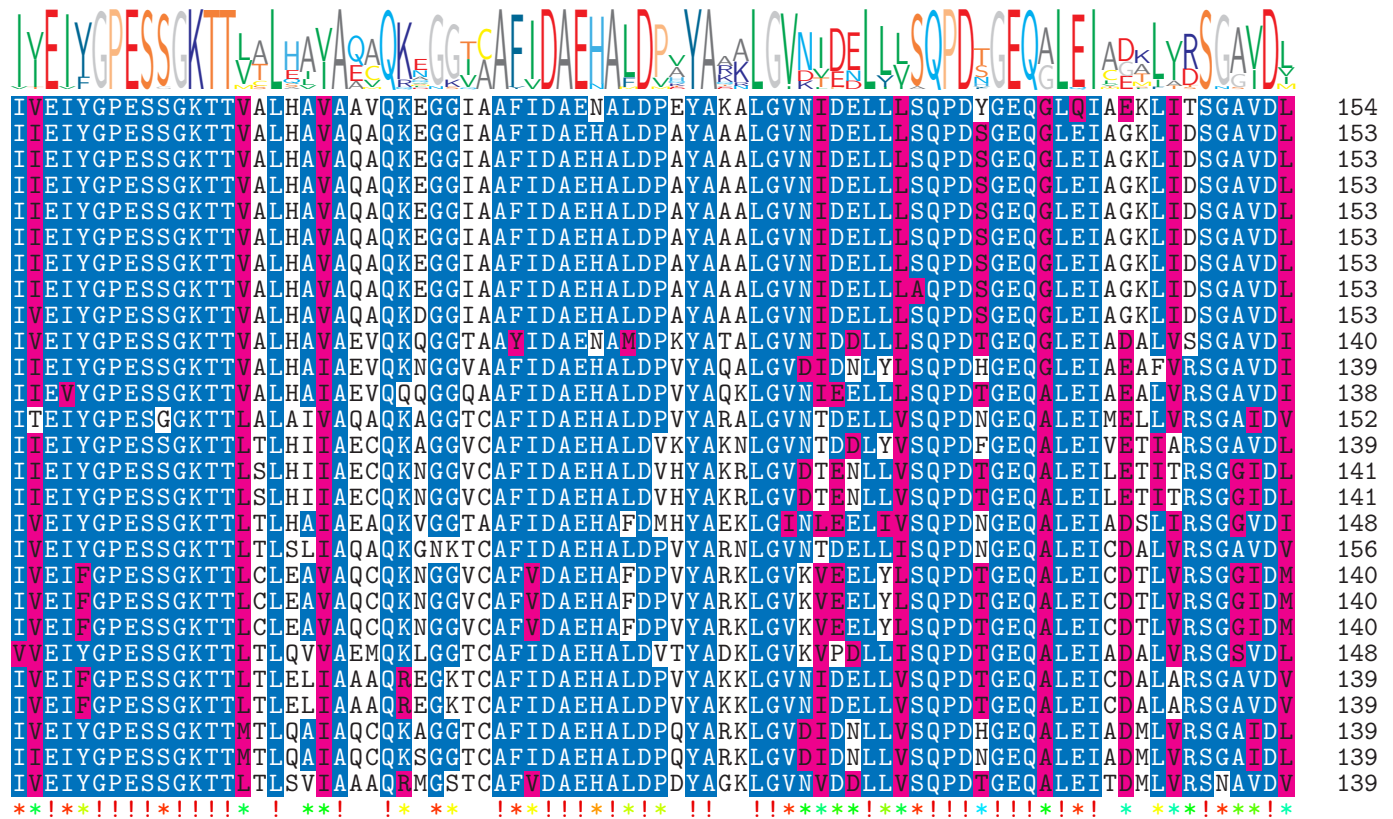
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consensus

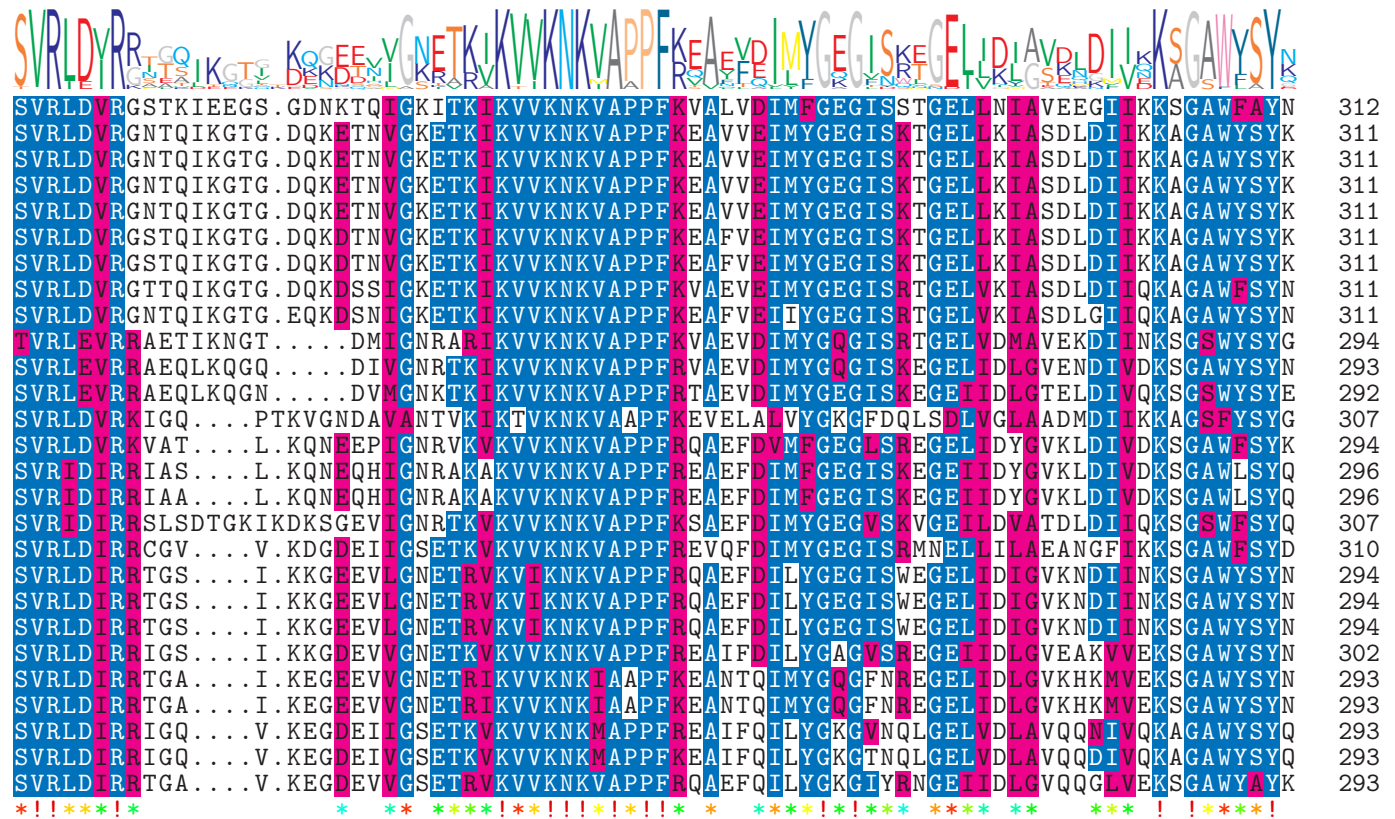
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGQGA	EKA	KNYL	KEH	QDV	FDE	IDHKV	RAAH	GL	DDSEVA	ETE	...	EET	...	TASKTKAKAKKEEKAVETEEIELE	384
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NTDB id 159 SP RS09750 WP 001085462.1	DEKIGQGS	ENAKKYLA	EHPEIF	DEIDKQV	RSKF	GL	IDGEEV	SEQD	...	TEN	...	KKDE	...	PKKEEA	..VNEEVPLD	377
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NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGS	ENAKRYLA	DHPQLF	DEIDRKV	RVVKF	GL	LEESEES	ESAM	...	AVA	...	SE	...	E	TDDLALD	369
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NTDB id 625 LCA RS02525 WP 011374196.1	SERIGQGS	RENAKNYLA	DHEDVE	DEVRLK	VRAAY	GIS	SDVPEED	LPT	...	TED	...	EQINIL	PDDSTEE		355	
NTDB id 1058553 ACFDBC RS07550 WP 002439552.1	GDRMGQGS	KENVK	NYLKEN	PQIKE	EIDRK	LREKL	GIFD	GDVDENEN	...	EDD	...	SPKTLF	DE		349	
NTDB id 115 BSU 16940 NP 389576.2	EERLGGGS	RENAKQFL	KENKD	IMLM	IQEQ	IREHY	GL	NNGVVQQA	...	EET	...	QEELEF	EE		348	
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NTDB id 1245 Cj1673c YP 002345041.1	DKKLGGGS	RENSKAFL	KENPEIA	DEITKAI	QNSMG	IEGMI	...	SGSEDD	..EGEE		343					
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NTDB id 1123 OK783 RS03845 WP 003688695.1	GAKIGQGS	KDNVRVWL	KENPEIS	DEIDAKI	RALNGVEMHI	...	TEGTQ	DETDGER	PEE		348					
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NTDB id 1205 PSJM300 05545 AFN77183.1	GSKIGQGS	KANAAKYLE	ENPEIGQ	EIEQQ	IRAKLL	VT	SAN	...	TKATP	VAEDLADADI		347				
consensus	**!	!!	*****	**	*****	*****	**	**	**							

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved