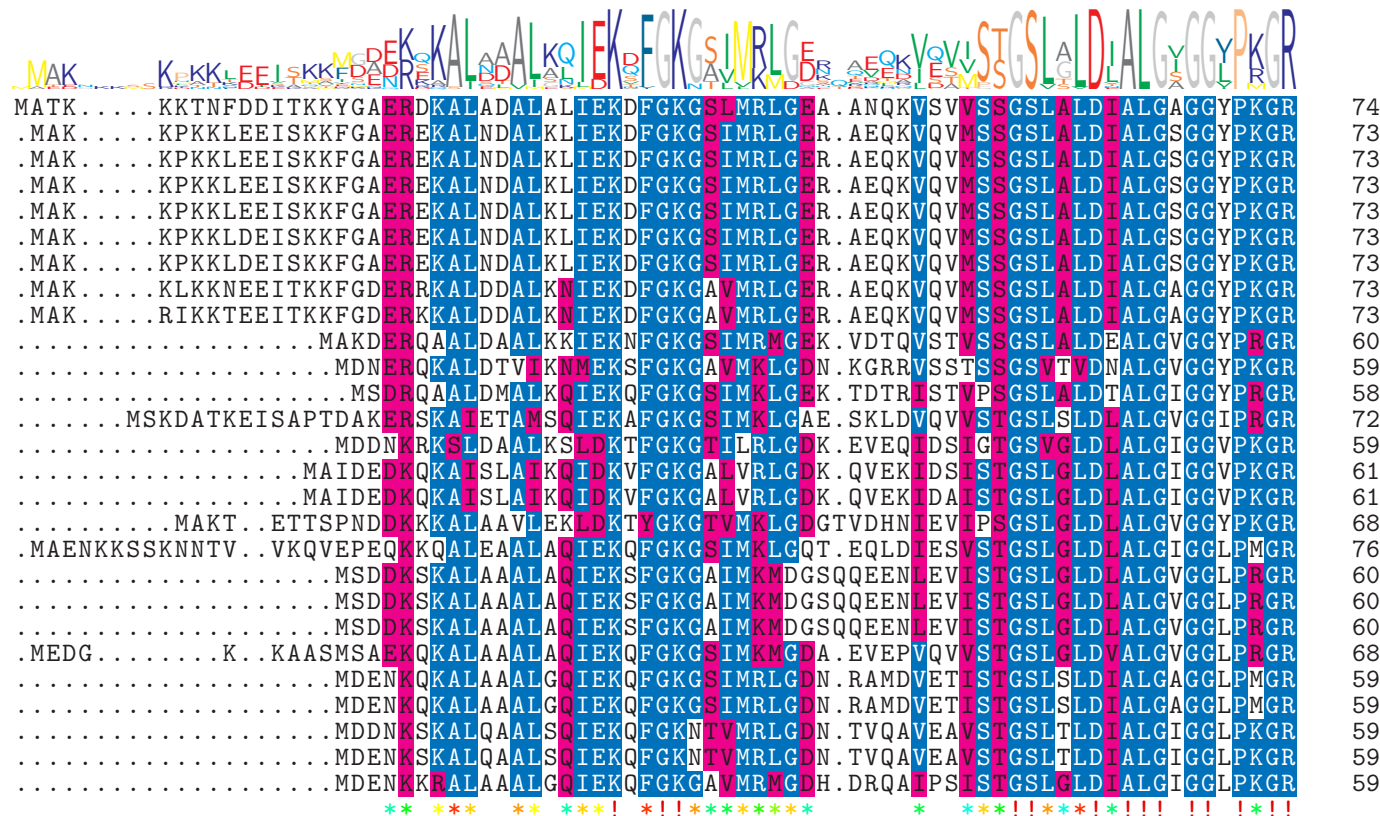


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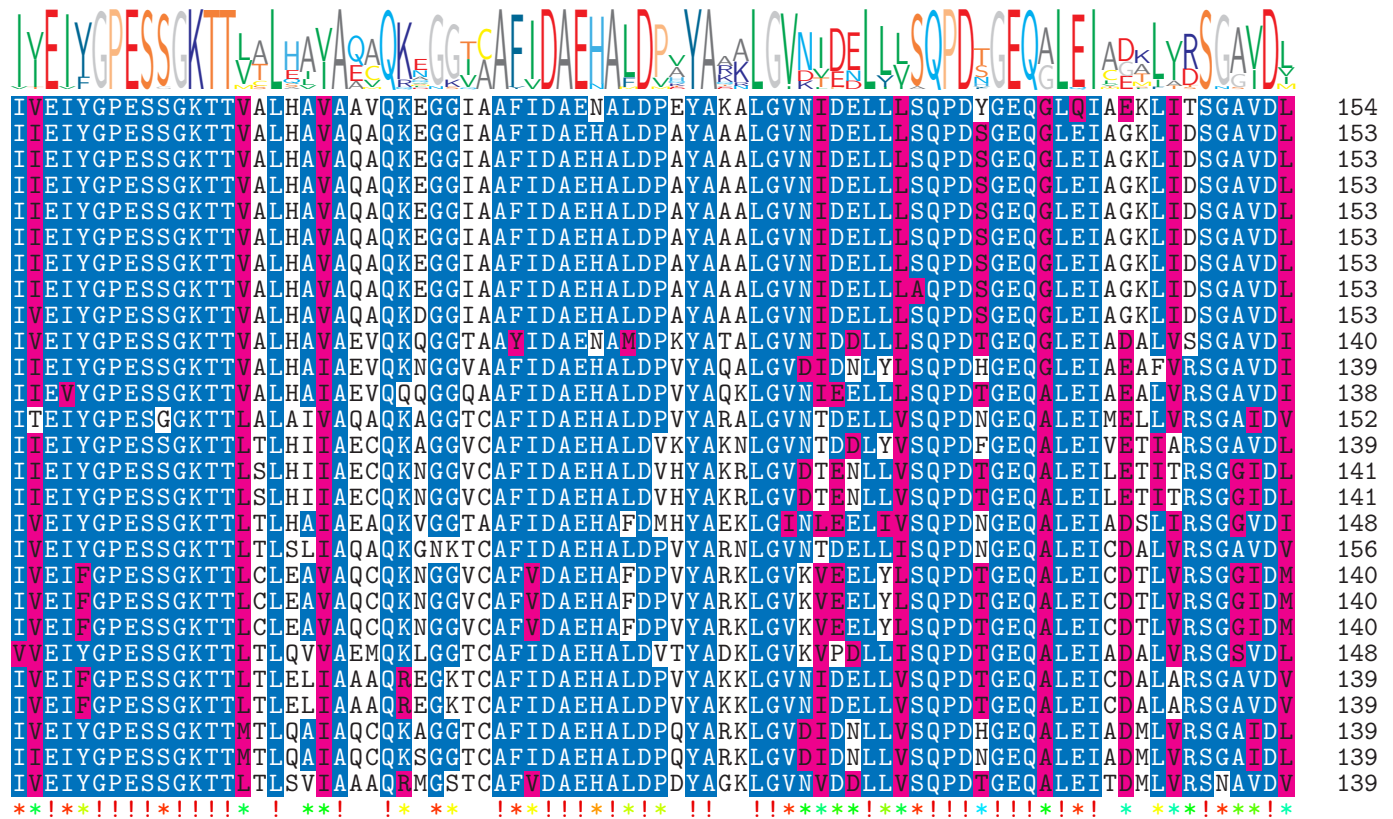
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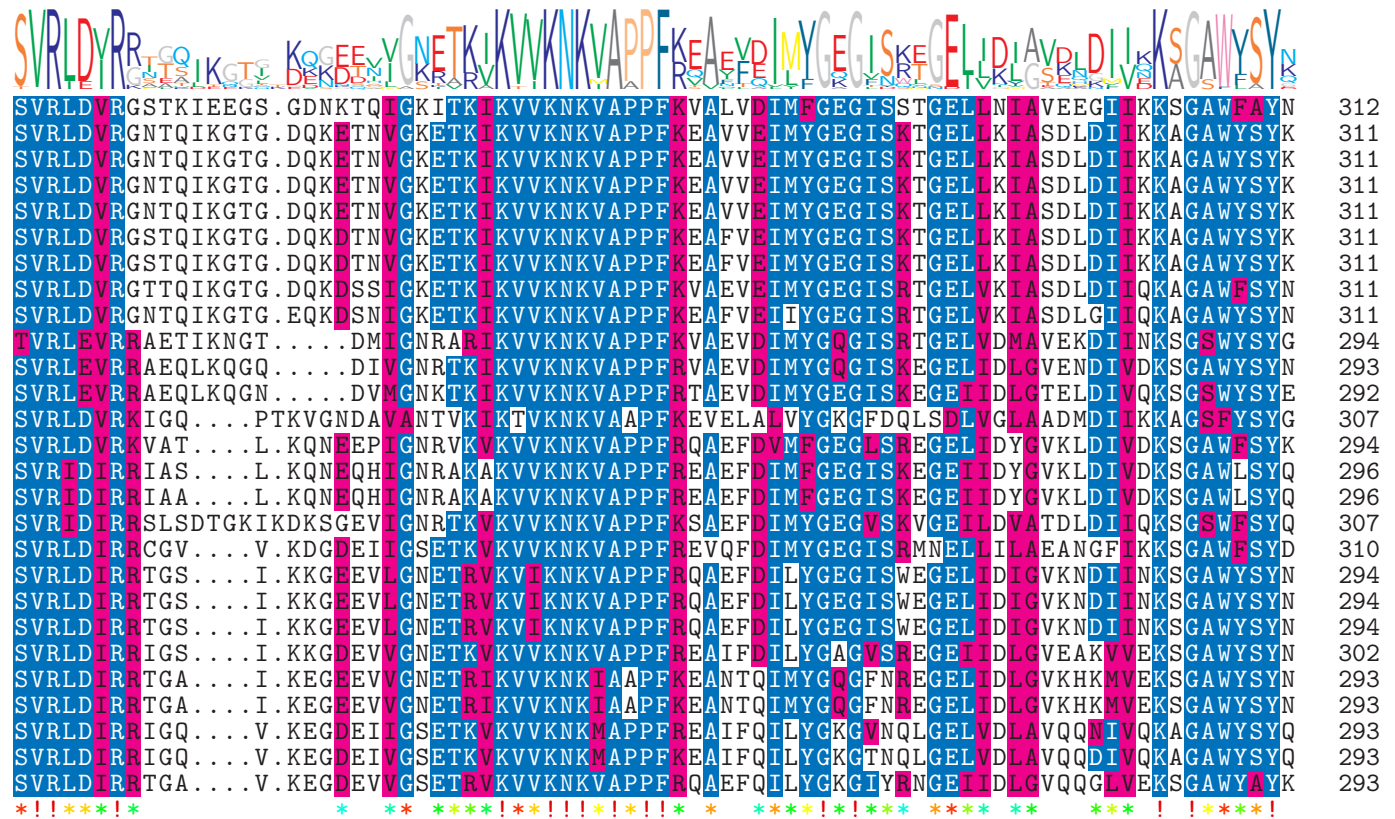
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consensus

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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
NTDB id 269 KZH43 RS08665 WP 001085462.1	DEKIGQGS	ENAKKYLA	EHPEIF	DEIDKQV	RSKF	GL	IDGEEV	SEQD	...	TEN	...	KKDE	...	PKKEEA	..VNEEVPLD	377
NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGS	ENAKKYLA	EHPEIF	DEIDKQV	RSKF	GL	IDGEEV	SEQD	...	TEN	...	KKDE	...	PKKEEA	..VNEEVPLD	377
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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	RVKF	GL	LEESE	EEESAM	...	AVA	...	SE	...	E	TDDLALD	369
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NTDB id 625 LCA RS02525 WP 011374196.1	SERIGQGS	RENAXNYLA	DHEDVE	DEVRLK	VRAAY	GIS	SDVPE	EDLPT	...	TED	...	EQINIL	PDDSTEE		355	
NTDB id 1055977 ACFDA8 RS07205 WP 002439552.1	GDRMGQGS	KENVKNYL	KENPQI	KEEIDR	KLREKL	GIF	DGDV	DENEN	...	EDD	...	SPKTLF	DE		349	
NTDB id 115 BSU 16940 NP 389576.2	EERLGGGS	RENAKQFL	KENKDI	MLMIQE	QIREHY	GL	NNGV	VQQA	...	EET	...	QEELEF	EE		348	
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NTDB id 1427 RS RS02760 WP 011000510.1	GERIGQGS	RDNCR	REFLRENA	EELARE	IENKVR	REHLGV	TPMG	...	AVTLA	..EEVEED		352				
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	GNKIGQGS	KNNVIRY	FEENNQIA	EIEIERN	IREQLL	TTGTN	...	GAVQ	..I	..EDEEE	PDLL..LES		349			
NTDB id 1205 PSJM300 05545 AFN77183.1	GSKIGQGS	KANAAKYLE	ENPEIGQ	EIEQQI	RAKLLV	TSAN	...	TKATP	VAED	LADADI		347				
consensus	**	!!	*****	**	*****	*****	**	**	**							

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved