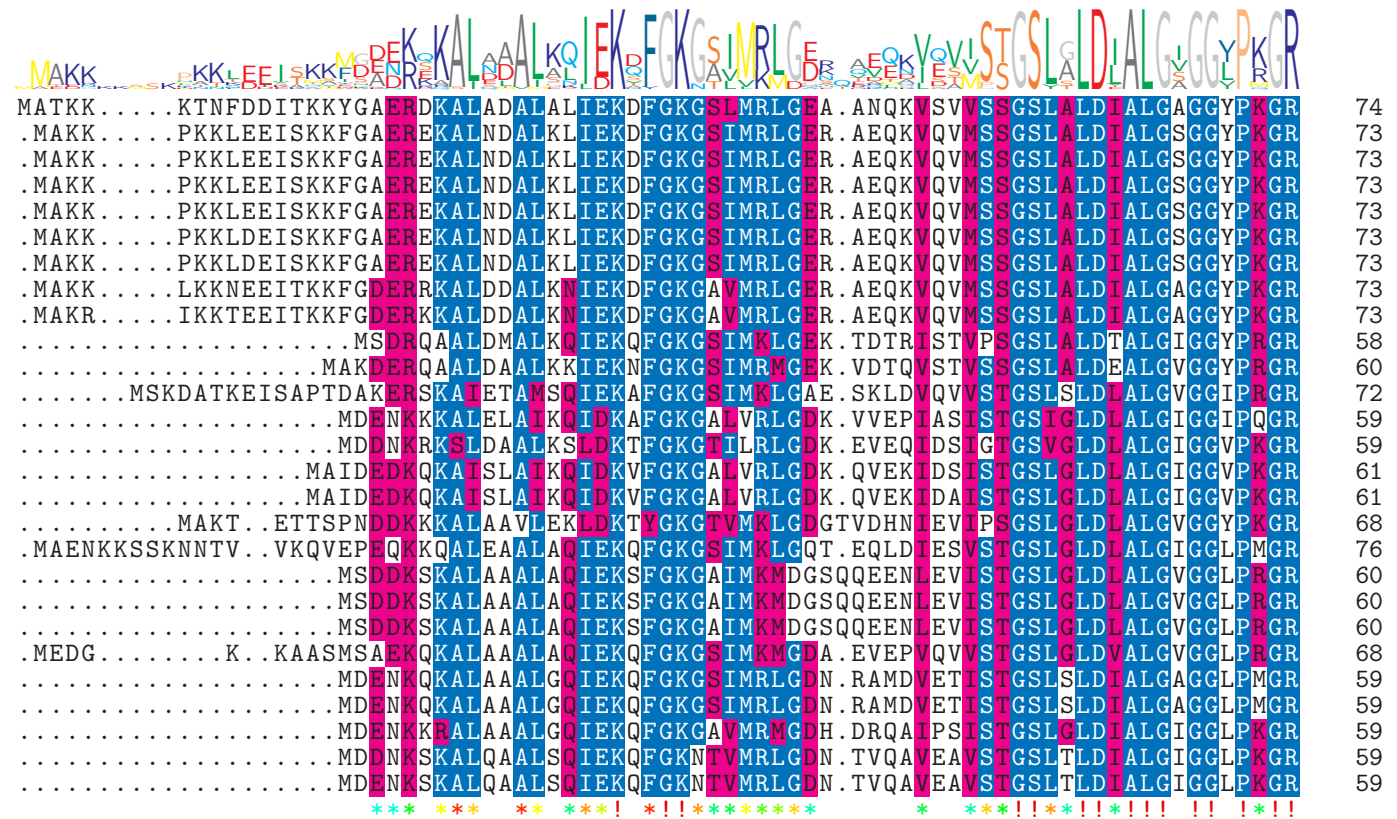


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consensus

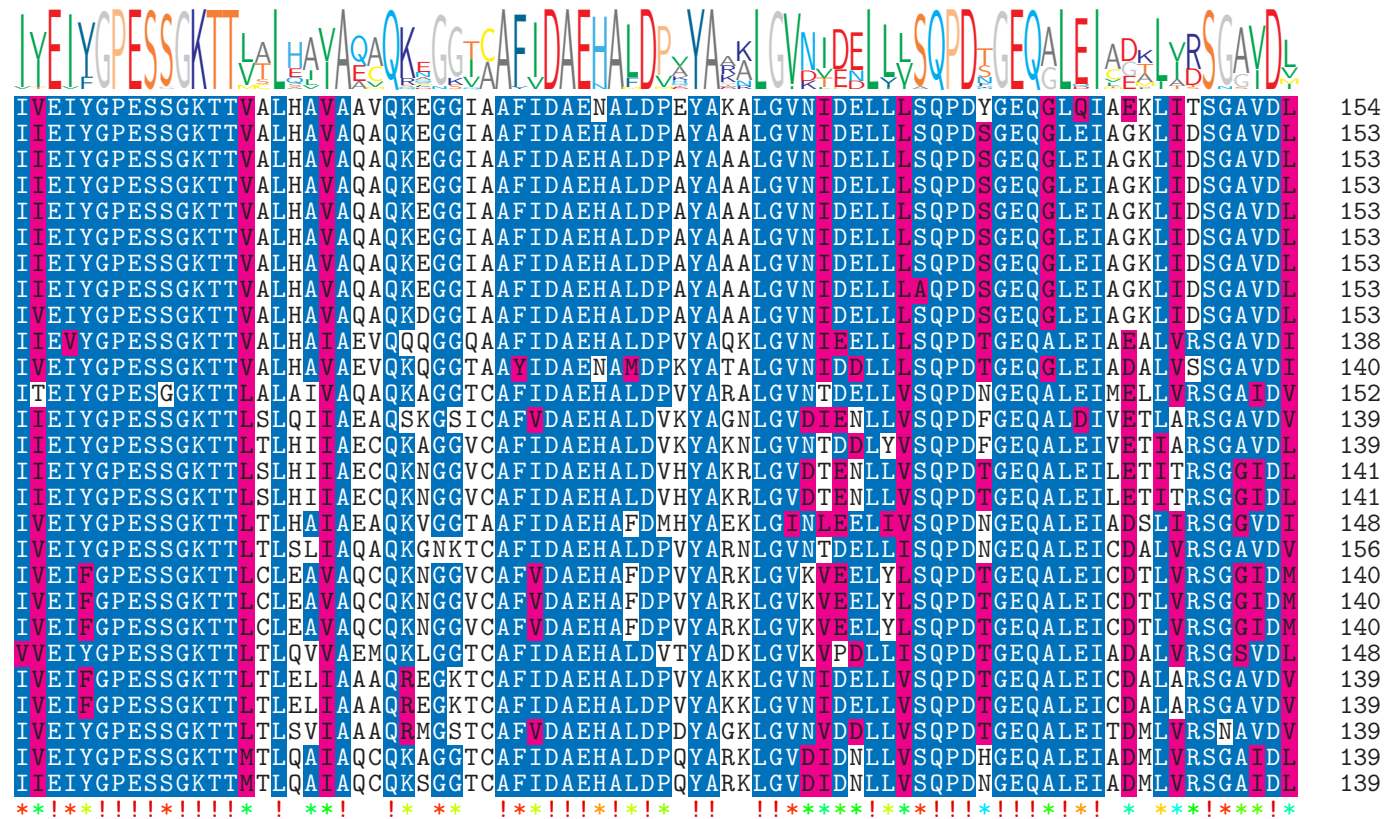
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consensus

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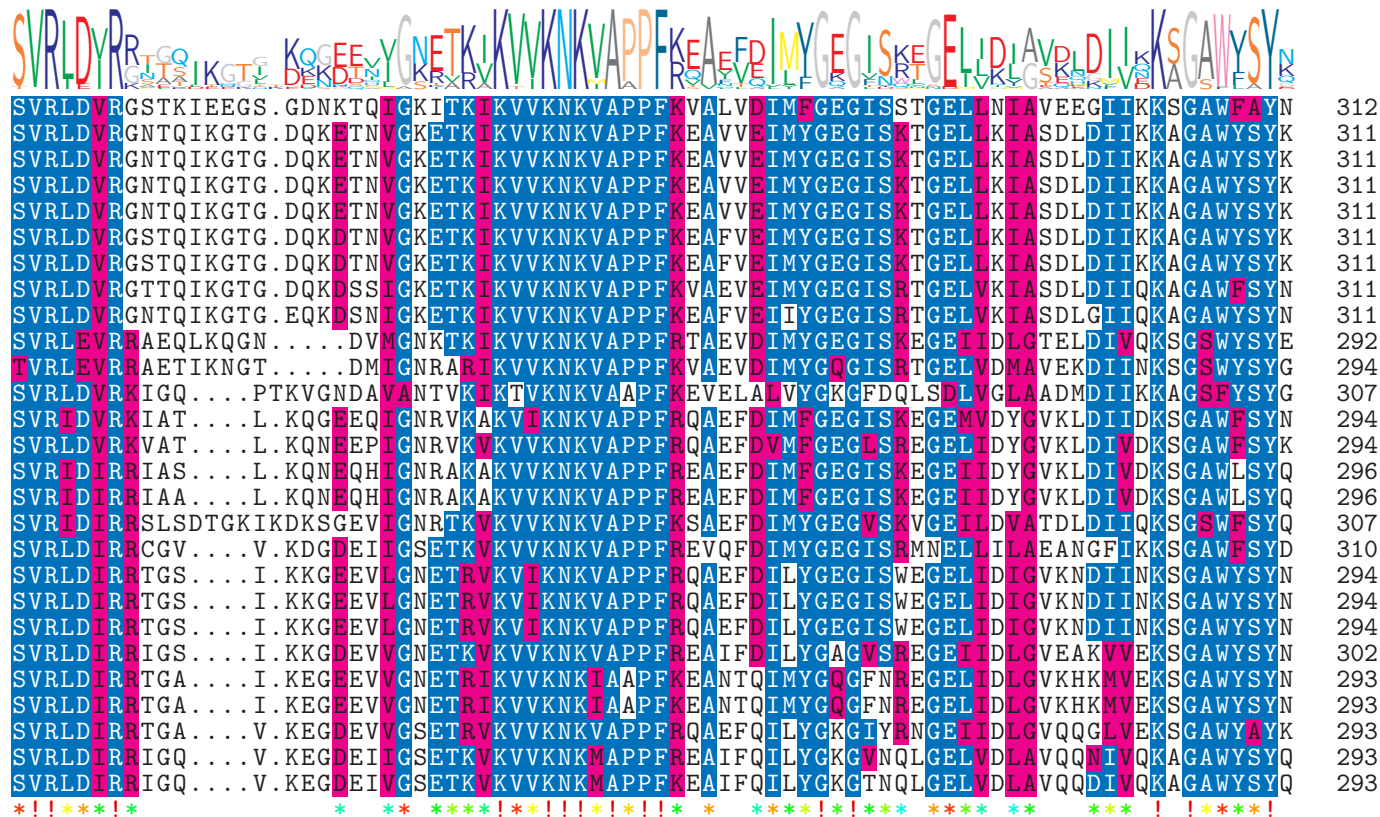


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NTDB	id	228	SPD	RS09265	WP	001085462.1	VVV	DSVAALV	PRAEI	DGDI	GDShVGLQAR	MMSQAMRKL	AGS	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GGRA	LKFYA	232				
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NTDB	id	527	SMSK321	RS11065	WP	001085454.1	VVV	DSVAALV	PRAEI	DGDI	GDShVGLQAR	MMSQAMRKL	AGS	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GGRA	LKFYA	232				
NTDB	id	499	SM12261	RS08315	WP	078228268.1	VVV	DSVAALV	PRAEI	DGDI	GDShVGLQAR	MMSQAMRKL	AGS	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GGRA	LKFYA	232				
NTDB	id	453	Spy49	1753c	ACI	62003.1	VVV	DSVAALV	PRAEI	DGDI	GDShVGLQAR	MMSQAMRKL	SAS	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GGRA	LKFYA	232				
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NTDB	id	625	LCA	RS02525	WP	011374196.1	LV	DSVAALV	PRAEI	EGEM	GDH	VGLQAR	LSQALRKL	SGT	INKTKT	TA	LFINQIREK	VGVM	.FGNPE	VTP	GGRA	LKFYS	219			
NTDB	id	1313	DR	RS12030	WP	010888966.1	VVV	DSVAAL	TPRAEI	EGDM	GD	SLPGLQAR	LSQALRKL	TA	ILSK	TGTAA	IFINQVREK	IGVM	.YGN	PETTT	GGRA	LKFYA	231			
NTDB	id	66888	UCH001	RS12645	WP	173636843.1	IV	VDSVAAL	TPKSEI	EGDM	GD	SHMGVQAR	LSQALRKL	TAV	VHKMNT	TV	IFINQIR	MKIG	TMG	YGS	PETTT	TGGN	NALKFYA	219		
NTDB	id	1245	Cj1673c	YP	002345041.1	IV	VDSVAAL	TPKAEI	EGDM	GD	QH	VGLQAR	LSQALRKL	TG	IVHKMNT	TV	IFINQIR	MKIG	AMG	YGT	PETTT	TGGN	NALKFYA	219		
NTDB	id	1211	C694	RS00795	WP	000963128.1	VVV	DSVAAL	TPKAEI	DGDM	GD	QH	VGLQAR	LSH	ALRKL	TG	VLHKMNT	TL	IFINQIR	MKIG	MMG	YGS	PETTT	TGGN	NALKFYA	221
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NTDB	id	1389	A4U84	RS07560	WP	021112712.1	II	VDSVAAL	TPKAEI	EGDM	GD	SHMGLQAR	LSQALRKL	TAN	IKNTNC	LV	VFINQIR	MKIG	VM	.FGNP	PETTT	TGGN	NALKFYA	235		
NTDB	id	1130	NGFG	RS03960	WP	003688695.1	VVV	DSVAALV	PKAEI	EGDM	GD	SHVGLQAR	LSQALRKL	TGH	IKKTNT	LV	VFINQIR	MKIG	VM	.FGSP	PETTT	TGGN	NALKFYS	219		
NTDB	id	1124	NGFG	RS03960	WP	003688695.1	VVV	DSVAALV	PKAEI	EGDM	GD	SHVGLQAR	LSQALRKL	TGH	IKKTNT	LV	VFINQIR	MKIG	VM	.FGSP	PETTT	TGGN	NALKFYS	219		
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GEKIGQGSENAKKFLADNPFIIDIDHKVRVQYGLIEEDNTEEKQSSKEK ETDEK . . ADKNLVLELDDTIELE	382
EERLGGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL . EFE . E	348
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logo

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved