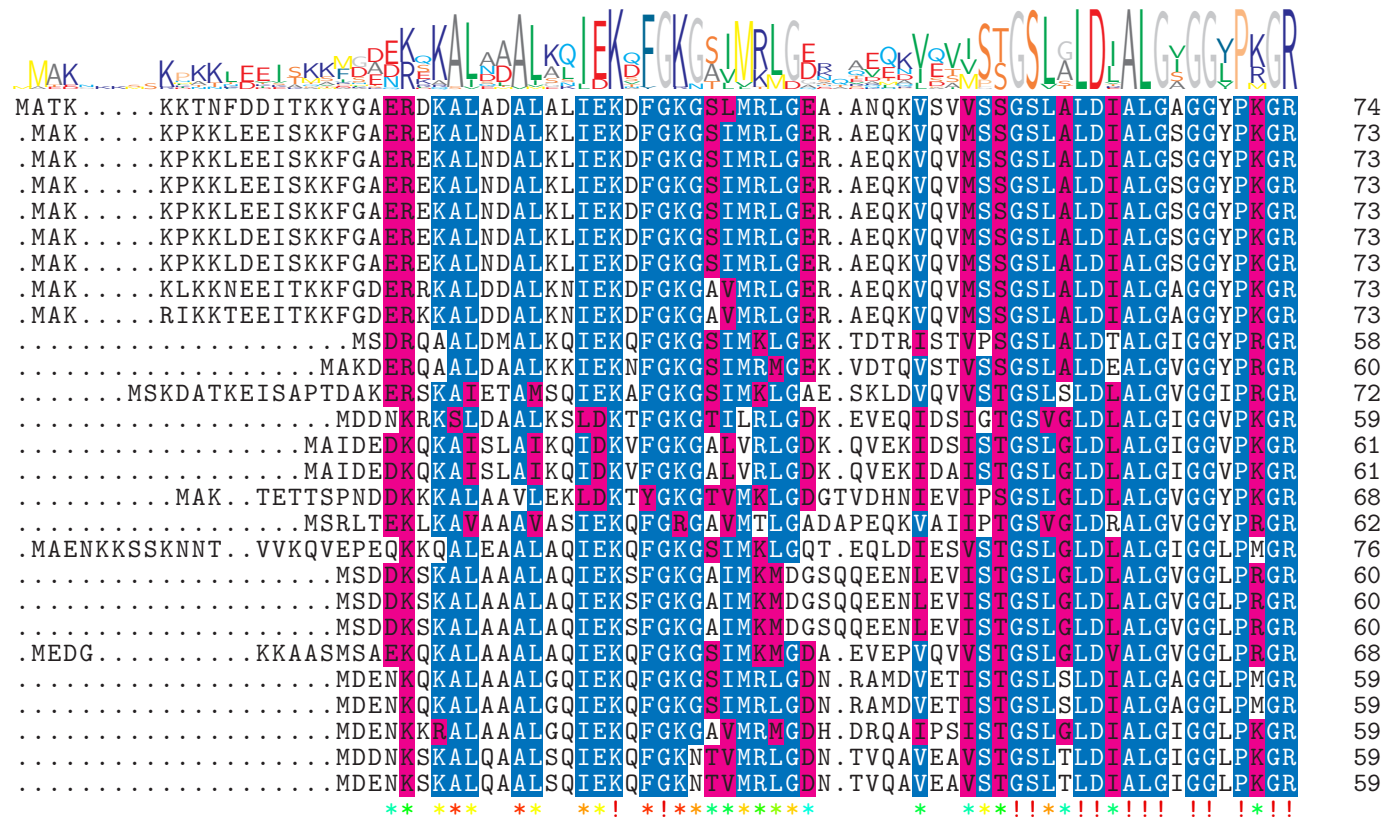


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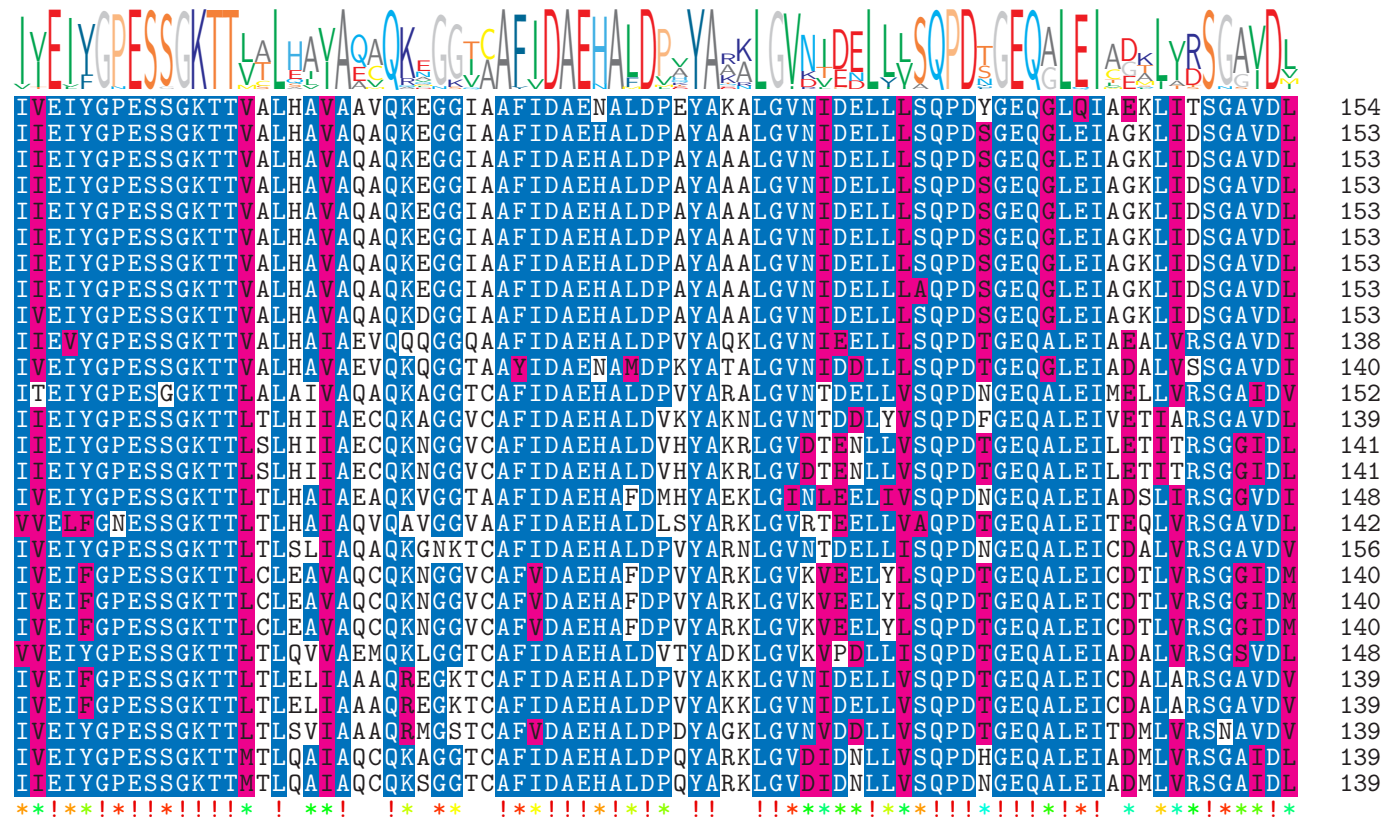
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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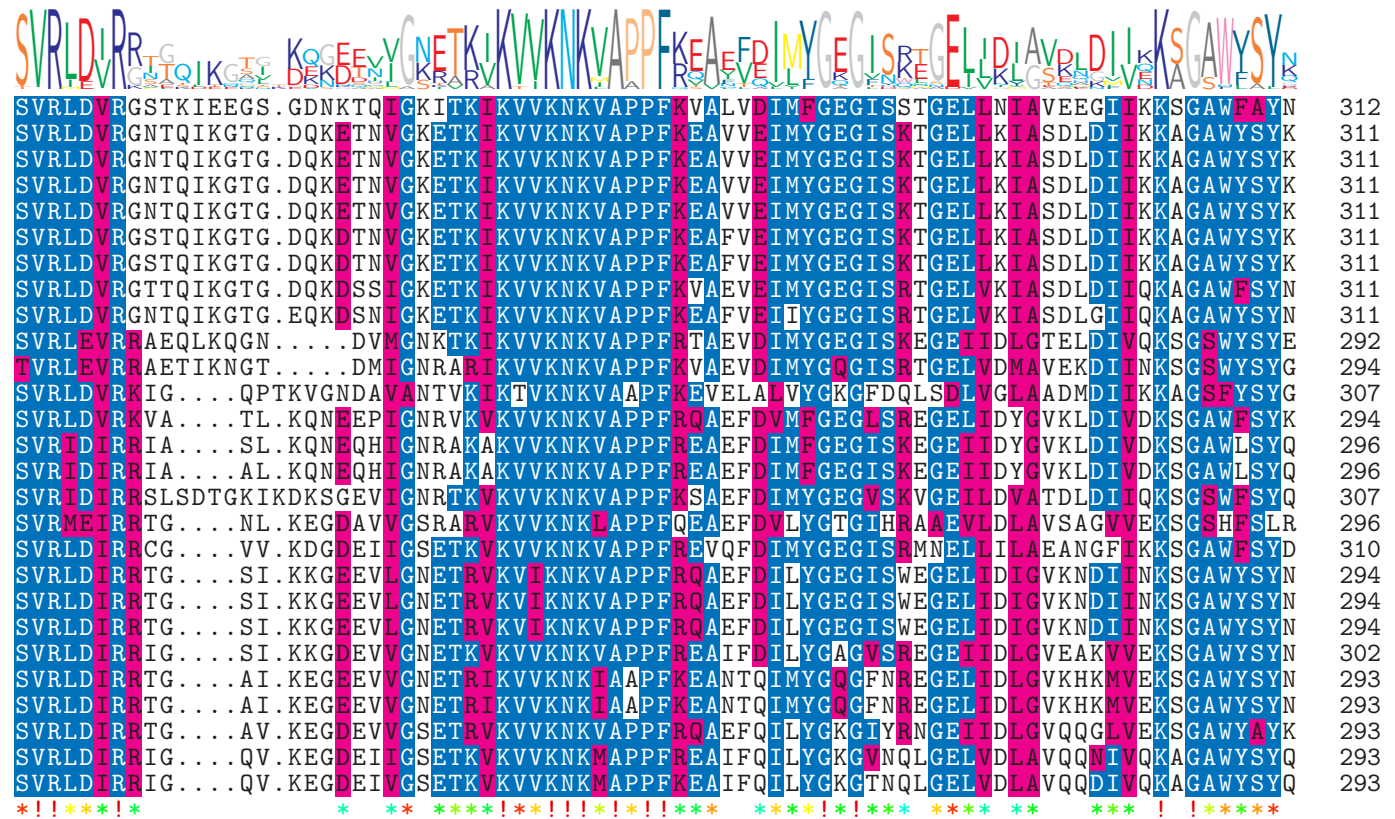


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NTDB	id	194	SPR	RS08825	WP	001085462.1	VVV	DSVAALVP	RA	EI	DGD	IGDSHVGLQAR	MSQAMRKL	GA	S	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GG	RALKFYA	232		
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NTDB	id	499	SM12261	RS08315	WP	078228268.1	VVV	DSVAALVP	RA	EI	DGD	IGDSHVGLQAR	MSQAMRKL	GA	S	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GG	RALKFYA	232		
NTDB	id	527	SMSK321	RS11065	WP	001085454.1	VVV	DSVAALVP	RA	EI	DGD	IGDSHVGLQAR	MSQAMRKL	GA	S	INKTKT	TA	IFINQLREK	VGVM	.FGNP	PETTP	GG	RALKFYA	232		
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NTDB	id	625	LCA	RS02525	WP	011374196.1	LV	VDSVAALVP	RA	EI	EGEM	GD	AHVGLQAR	LSQALRKL	SGT	INKTKT	TA	LFINQIREK	VGVM	.FGNPE	VTP	GG	RALKFYS	219		
NTDB	id	1313	DR	RS12030	WP	010888966.1	VVV	DSVAAL	TP	RA	EI	EGDM	GD	SLPGLQAR	LSQALRKL	TA	ILSKTGT	AA	IFINQVREK	IGVM	.YGN	PETTT	GG	RALKFYA	231	
NTDB	id	1245	Cj1673c	YP	002345041.1	IV	VDSVAAL	TP	KA	EI	EGDM	GD	QHVGLQAR	LSQALRKL	TG	IVHKMNTT	V	IFINQIR	MKIGAM	GYGT	PETTT	GG	NALKFYA	219		
NTDB	id	1211	C694	RS00795	WP	000963128.1	VVV	DSVAAL	TP	KA	EI	DGDM	GD	QHVGLQAR	LSH	ALRKIT	GV	LHKMNTT	L	IFINQIR	MKIGMM	GYG	SPETTT	GG	NALKFYA	221
NTDB	id	1214	NCTC11637	00359	SQJ03206.1	VVV	DSVAAL	TP	KA	EI	DGDM	GD	QHVGLQAR	LSH	ALRKIT	GV	LHKMNTT	L	IFINQIR	MKIGMM	GYG	SPETTT	GG	NALKFYA	221	
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NTDB	id	1389	A4U84	RS07560	WP	021112712.1	II	VDSVAAL	TP	KA	EI	EGDM	GD	SHMGLQAR	LSQALRKL	TAN	IKNTNCL	LV	VFINQIR	MKIGVM	.FGNP	PETTT	GG	NALKFYA	235	
NTDB	id	1130	NGFG	RS03960	WP	003688695.1	VVV	DSVAALVP	KA	EI	EGDM	GD	SHVGLQAR	LSQALRKL	TG	HIKKTNTL	V	VFINQIR	MKIGVM	.FGSP	PETTT	GG	NALKFYS	219		
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consensus

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[illegible]

logo

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consensus			

- non conserved
- similar
- ≥ 50% conserved