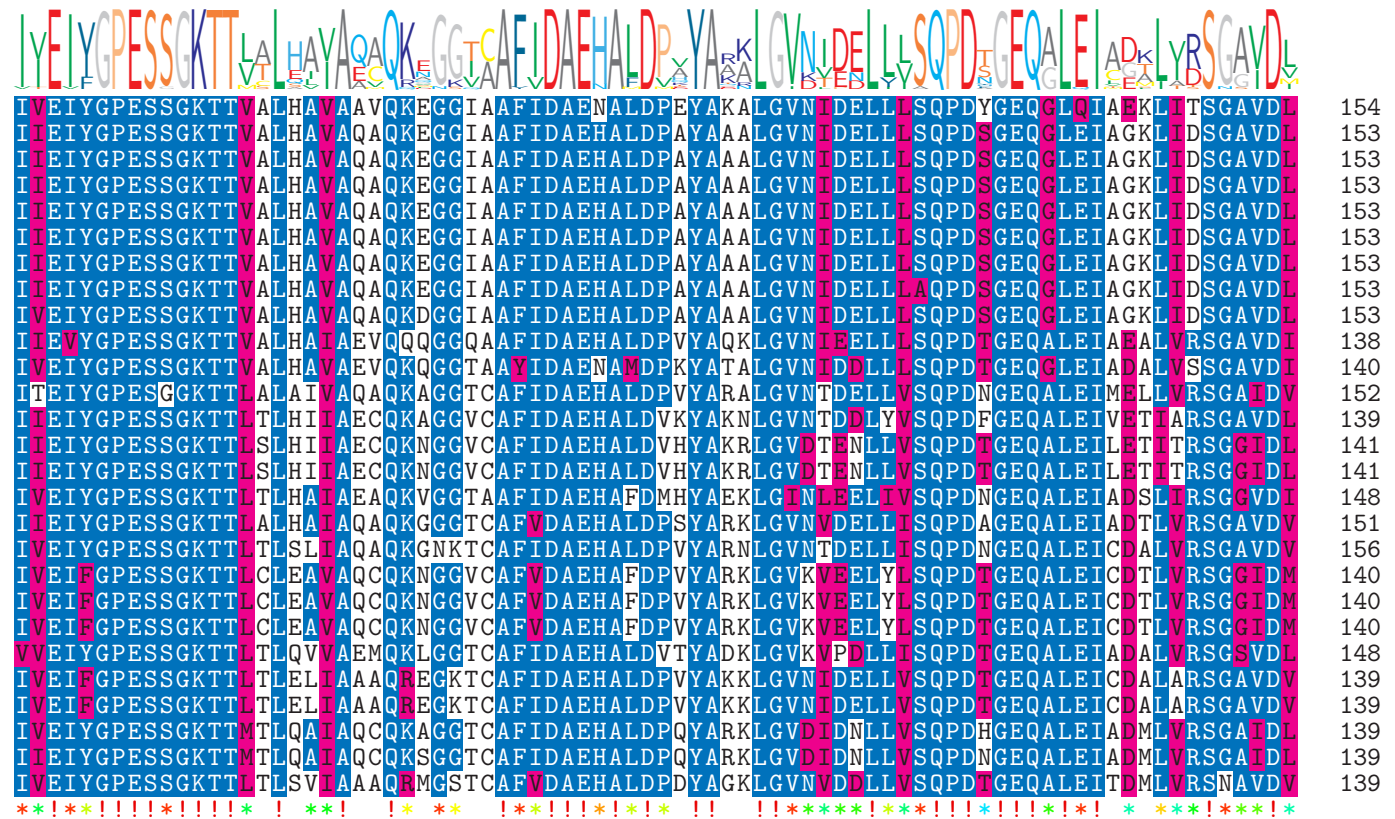


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NTDB id 408 SMU RS09530 WP 002262392.1	.MAKRI.....KKTEEITKKFGD	ERK	KAL	DDAL	KN	IEKD	FGKGA	VMR	LGER	.AEQK	VQVM	SSG	SLALD	I	ALG	AGGY	PKGR	73						
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NTDB id 1334 RAO	C RS04870 WP 004918236.1	MAKTET	.TSPND	DKKK	LA	AVLEK	LD	KT	YVGK	TVM	KLGD	GTVD	HN	.IEV	IPS	GS	GLDL	E	ALGV	GGY	PKGR	68		
NTDB id 158250 AMK58 RS10040 WP 014240962.1	MSSAQLRLV	.EKDSM	DKQ	KAL	DAAL	QA	IE	AFGKGS	IMKL	G	ARENT	VET	EV	.VST	GS	GLDL	E	ALGI	GG	LPGR	71			
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NTDB id 1130 NGFG RS03960 WP 003688695.1	MSDD	KS	KAL	AAAL	Q	IEKS	FGKGA	IMKM	D	GS	Q	Q	EE	N	.LEV	IST	GS	GLDL	E	ALGV	GG	LP	GR	60
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NTDB id 1123 OK783 RS03845 WP 003688695.1	MSD	KS	KAL	AAAL	Q	IEKS	FGKGA	IMKM	D	GS	Q	Q	EE	N	.LEV	IST	GS	GLDL	E	ALGV	GG	LP	GR	60
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....KKA	.ASMSA	EKK	KAL	AAAL	Q	IEKQ	FGKGS	IMKM	G	DA	.EVEP	VQV	.VST	GS	GLD	V	ALGV	GG	LP	GR	68		
NTDB id 1165 A1552VC RS01565 WP 000344154.1	MDEN	KQ	KAL	AAAL	G	IEKQ	FGKGS	SIMR	L	GD	N	.RAMD	VET	.IST	GS	LS	LD	I	ALG	AG	GL			

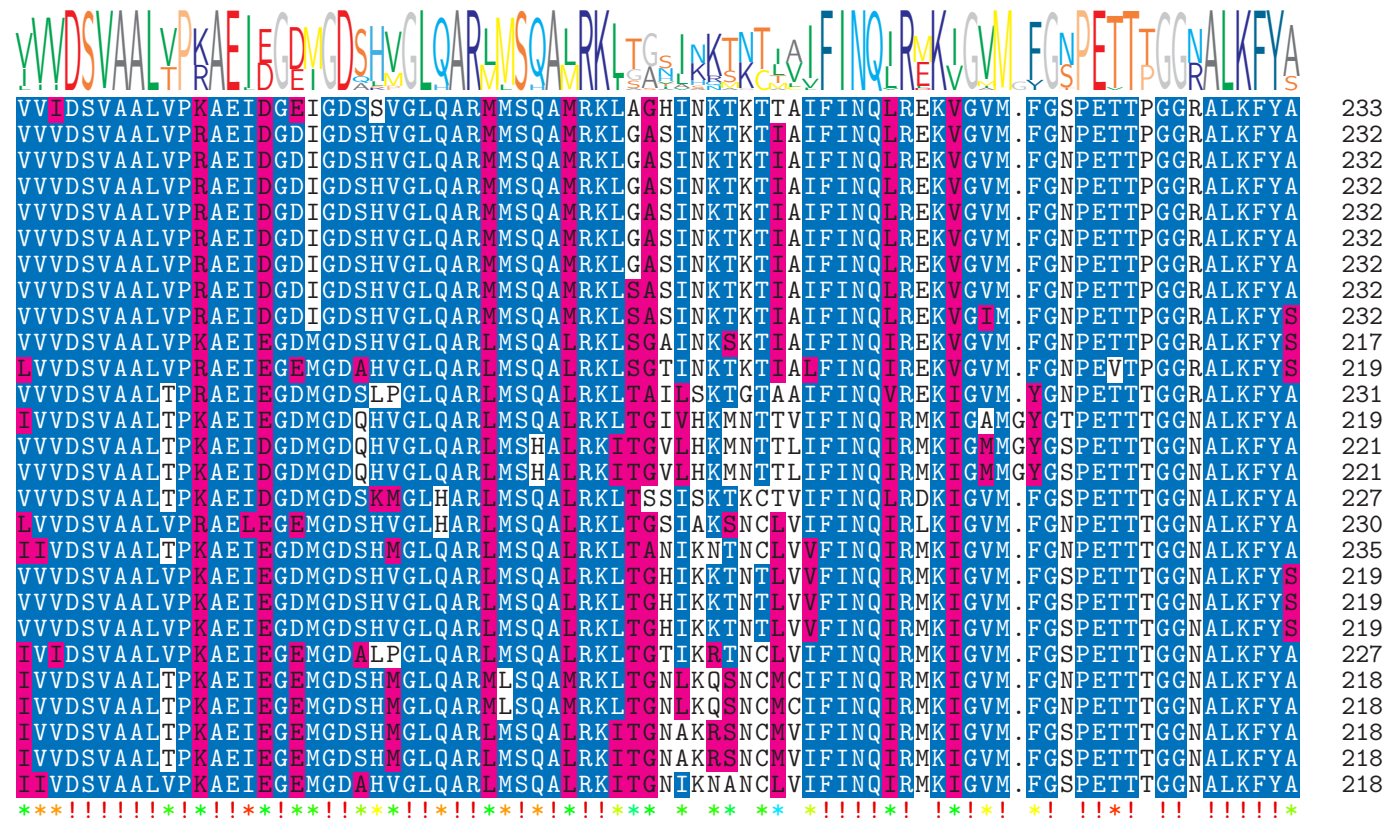
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consensus

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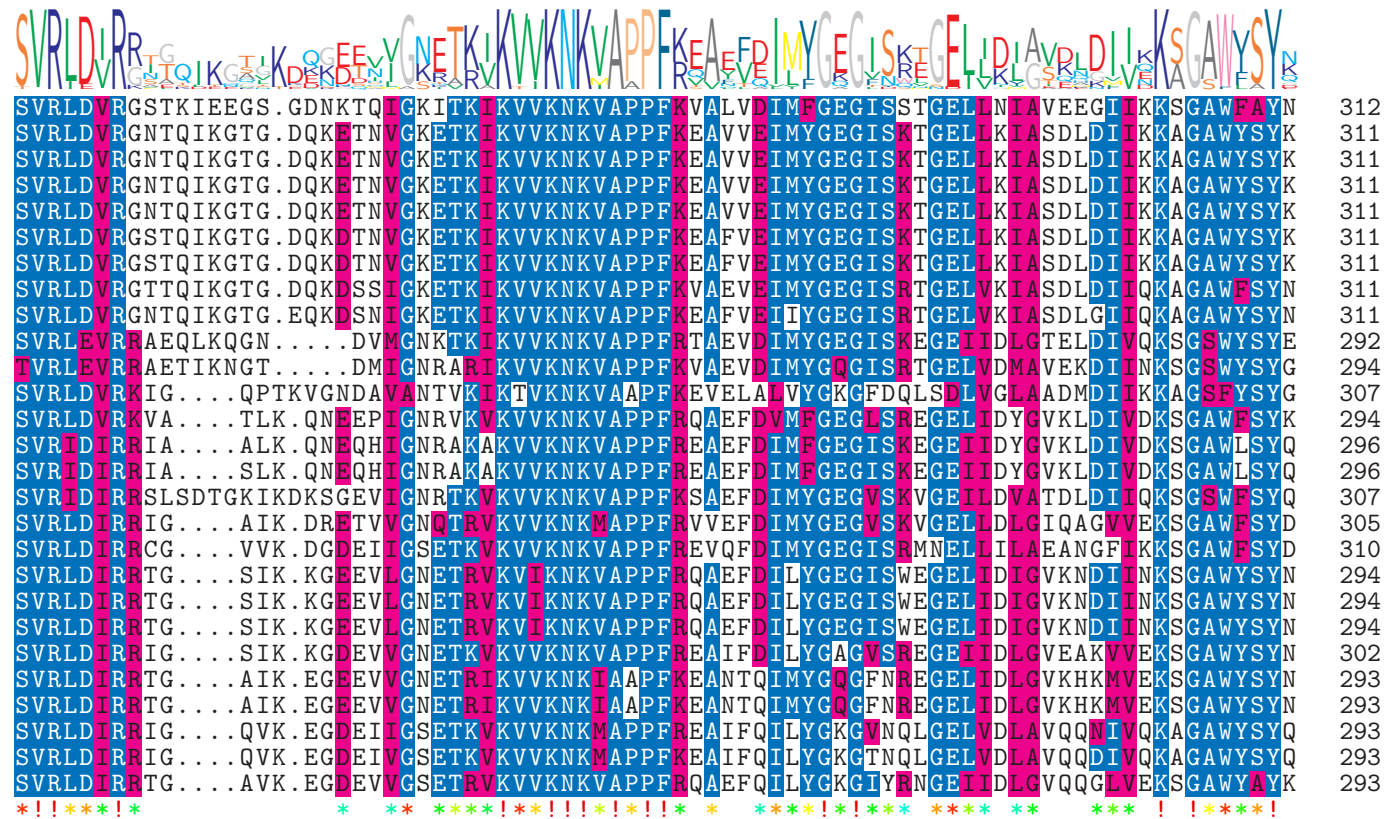
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consensus

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consensus	**!!!!***** ** ***** ***** ** *					*

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved