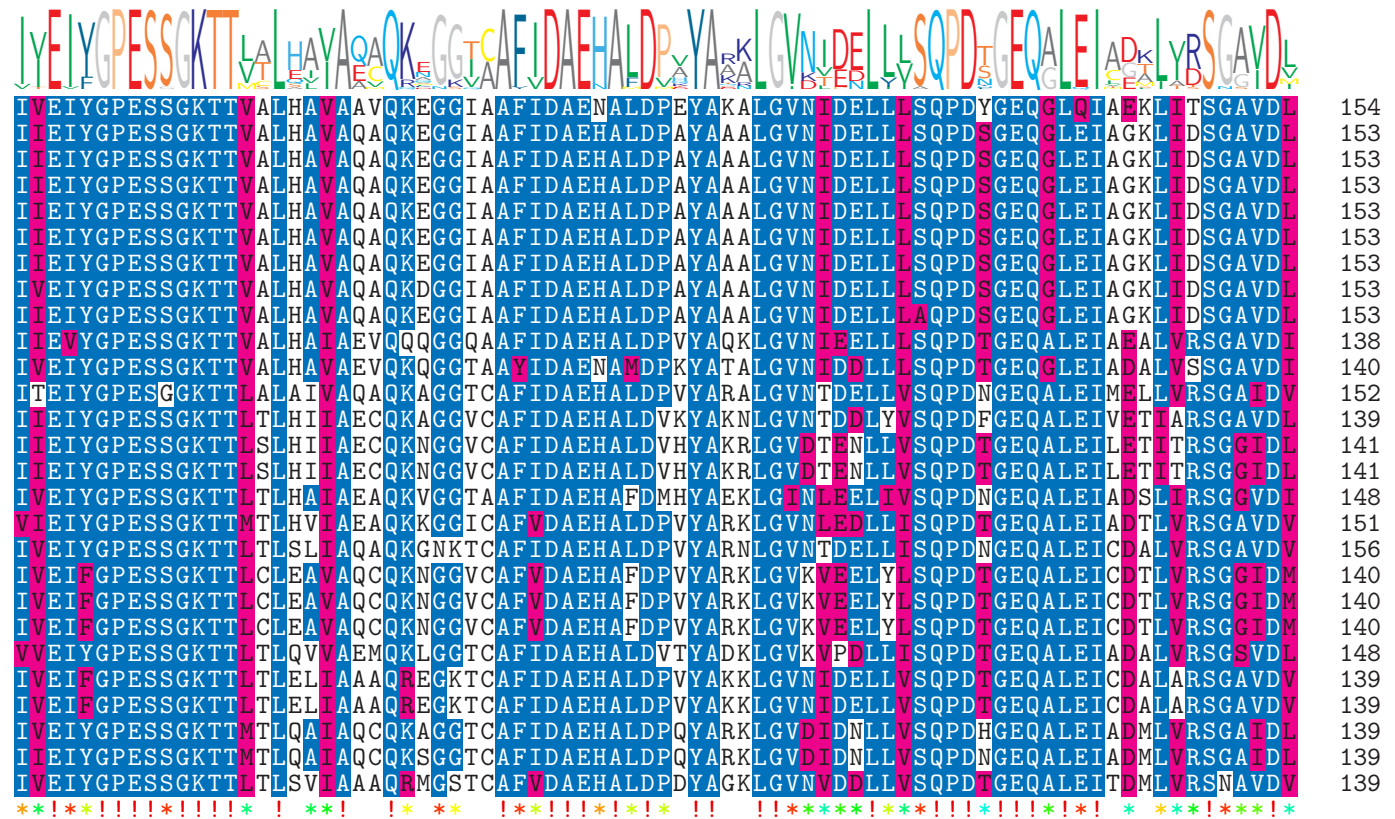


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NTDB	id	269	KZH43	RS08665	WP	001085462.1	MAKKPKKLEEISKKFGA	ERE	KAL	NDAL	KL	IEKD	FGKGS	SIMRLGER	.AEQK	VQVM	SSGSL	ALDI	I	ALGS	GGY	PKGR	73	
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NTDB	id	194	SPR	RS08825	WP	001085462.1	MAKKPKKLEEISKKFGA	ERE	KAL	NDAL	KL	IEKD	FGKGS	SIMRLGER	.AEQK	VQVM	SSGSL	ALDI	I	ALGS	GGY	PKGR	73	
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NTDB	id	453	Spy49	1753c	ACI	62003.1	MAKKLKKNEEITKKFGD	ERR	KAL	DDAL	KN	IEKD	FGKGA	VMRLGER	.AEQK	VQVM	SSGSL	ALDI	I	ALG	AGGY	PKGR	73	
NTDB	id	115	BSU	16940	NP	389576.2	MSDRQAALDMALKQ	IEK	Q	FGKGS	IMKLGEK	.TDTR	IST	VP	SGSL	ALD	T	ALGI	GGY	PRGR	58			
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NTDB	id	1334	RAOC	RS04870	WP	004918236.1	MA...K.TETT..SPND	DKKK	LA	AVLEK	LDK	T	FGKGT	VMKLGD	GTVDHN	IEV	IP	SGSL	GLDL	ALGV	GGY	PKGR	68	
NTDB	id	212850	BVL55	RS08920	WP	075996593.1	MAQMSL.RLVE..DNTV	DKSK	ALDA	ALSQ	IER	AFGKS	IMKLGKNEQ	VVEI	ET	VP	TGSL	GLDI	ALGV	GGL	PKGR	71		
NTDB	id	1389	A4U84	RS07560	WP	021112712.1	MAENKKS	SKNNT.VVKQ..VEPE	QKQ	AL	EAA	LAQ	IEKQ	FGKGS	IMKLGQT	.EQLD	IES	V	TGSL	GLDL	ALGI	GGL	PMGR	76
NTDB	id	1130	NGFG	RS03960	WP	003688695.1	MSDDKSKALAAAL	LAQ	IEKS	FGKGA	IMKMDGS	QQEEN	LEV	IST	GSL	GLDL	ALGV	GGL	PRGR	60				
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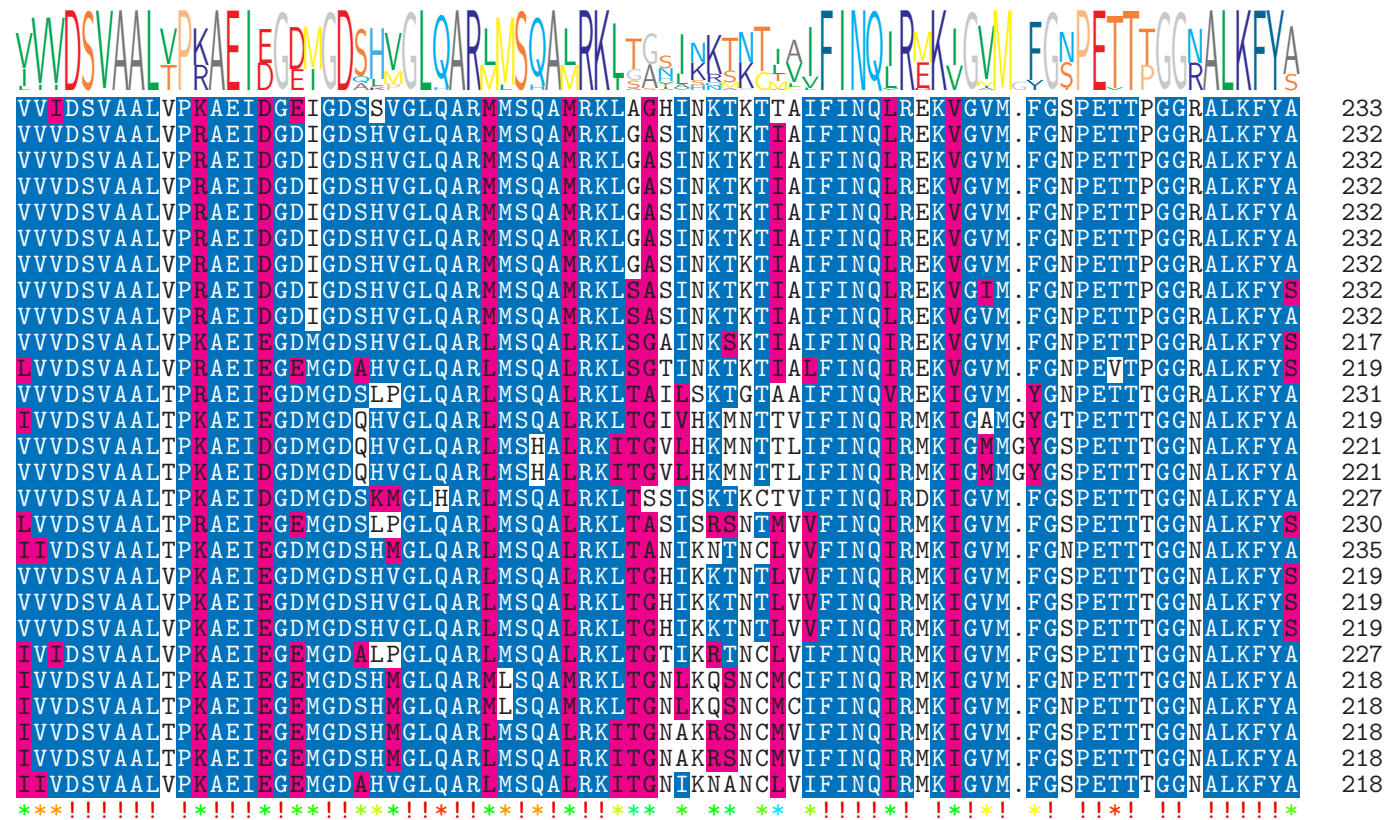
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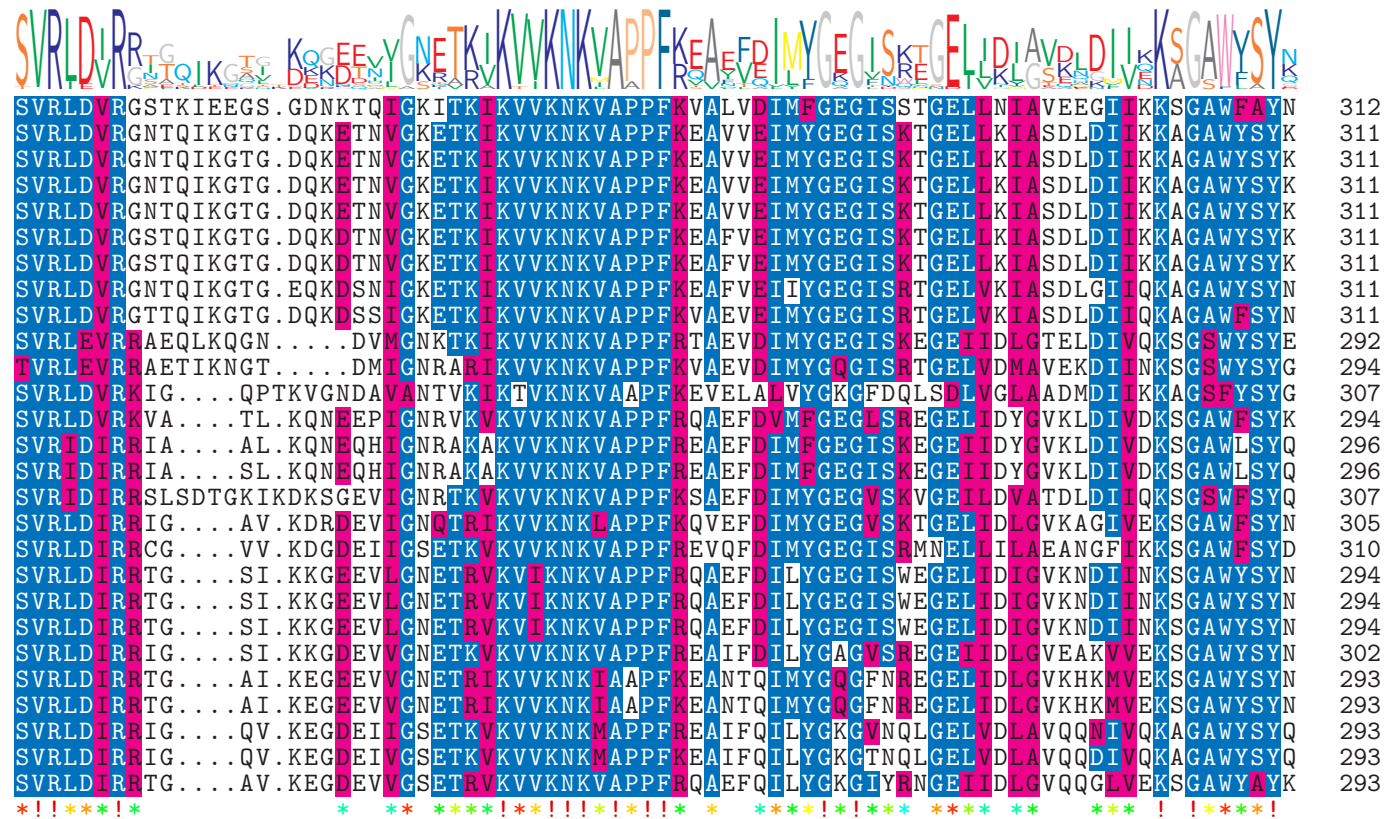
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consensus

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NTDB id 1123 OK783 RS03845 WP 003688695.1	GAKIGQG	KDNVRVWLKENPEIS	DEIDAKIRALNGVEM	HIT	EGTQDET	DGER.....PEE.....	348	
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consensus

 non conserved
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 ≥ 50% conserved