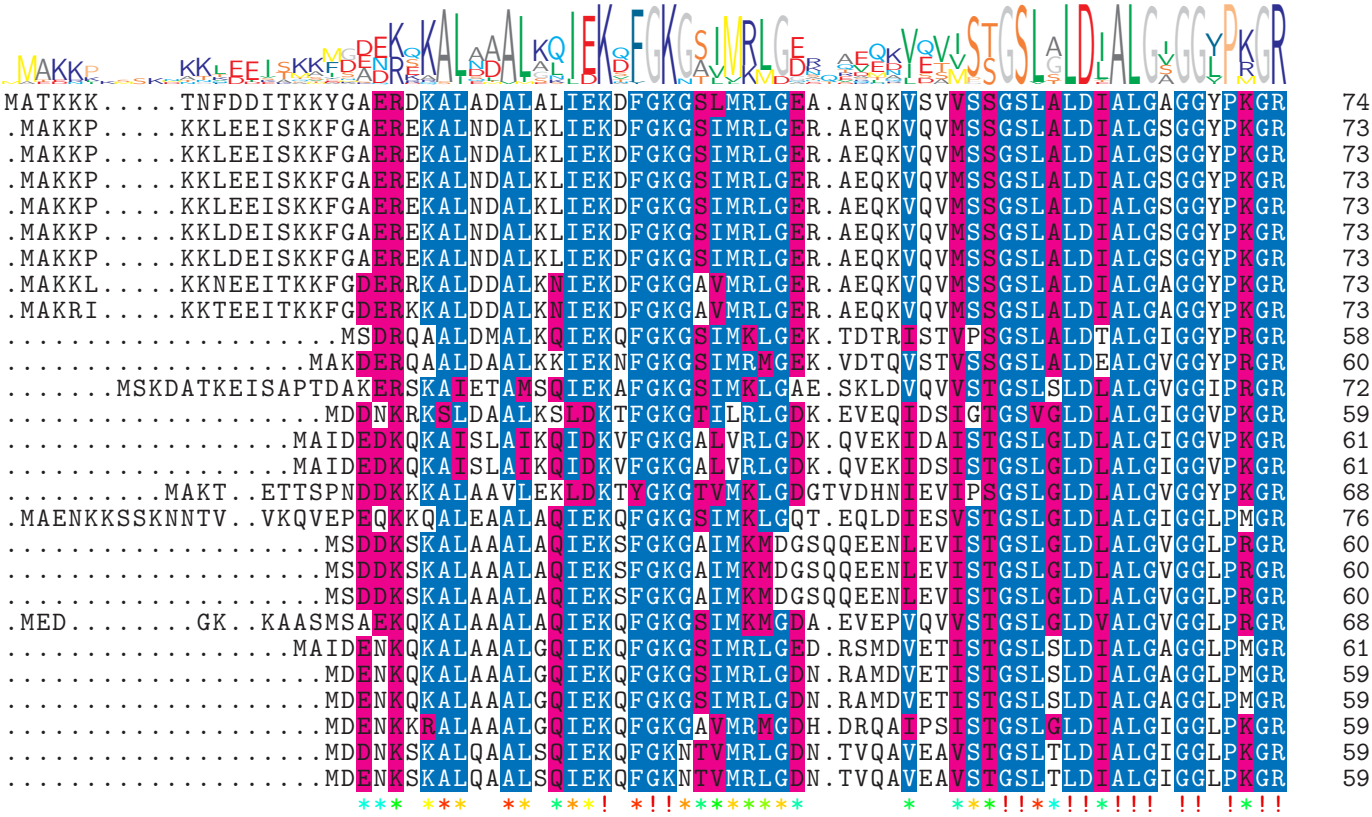
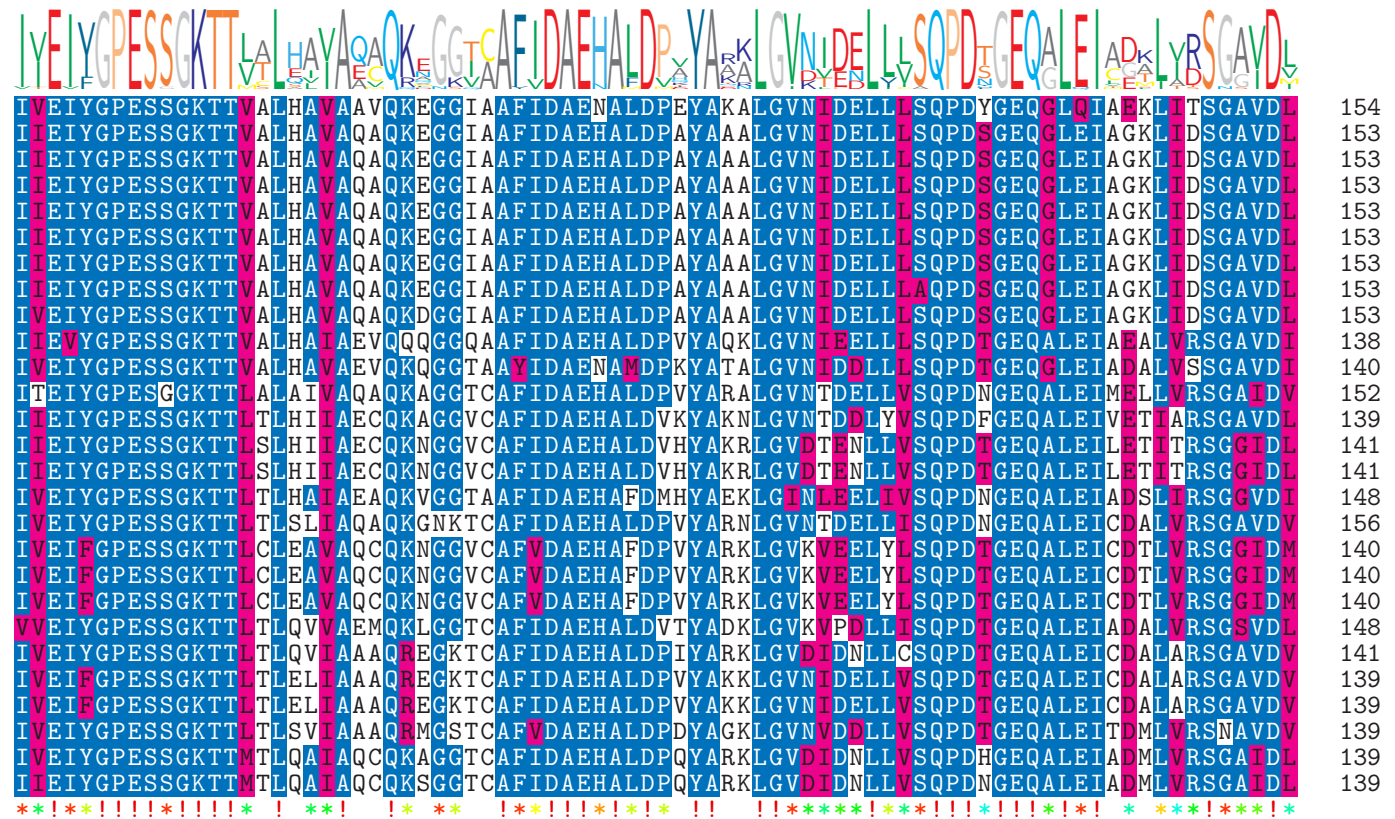


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus



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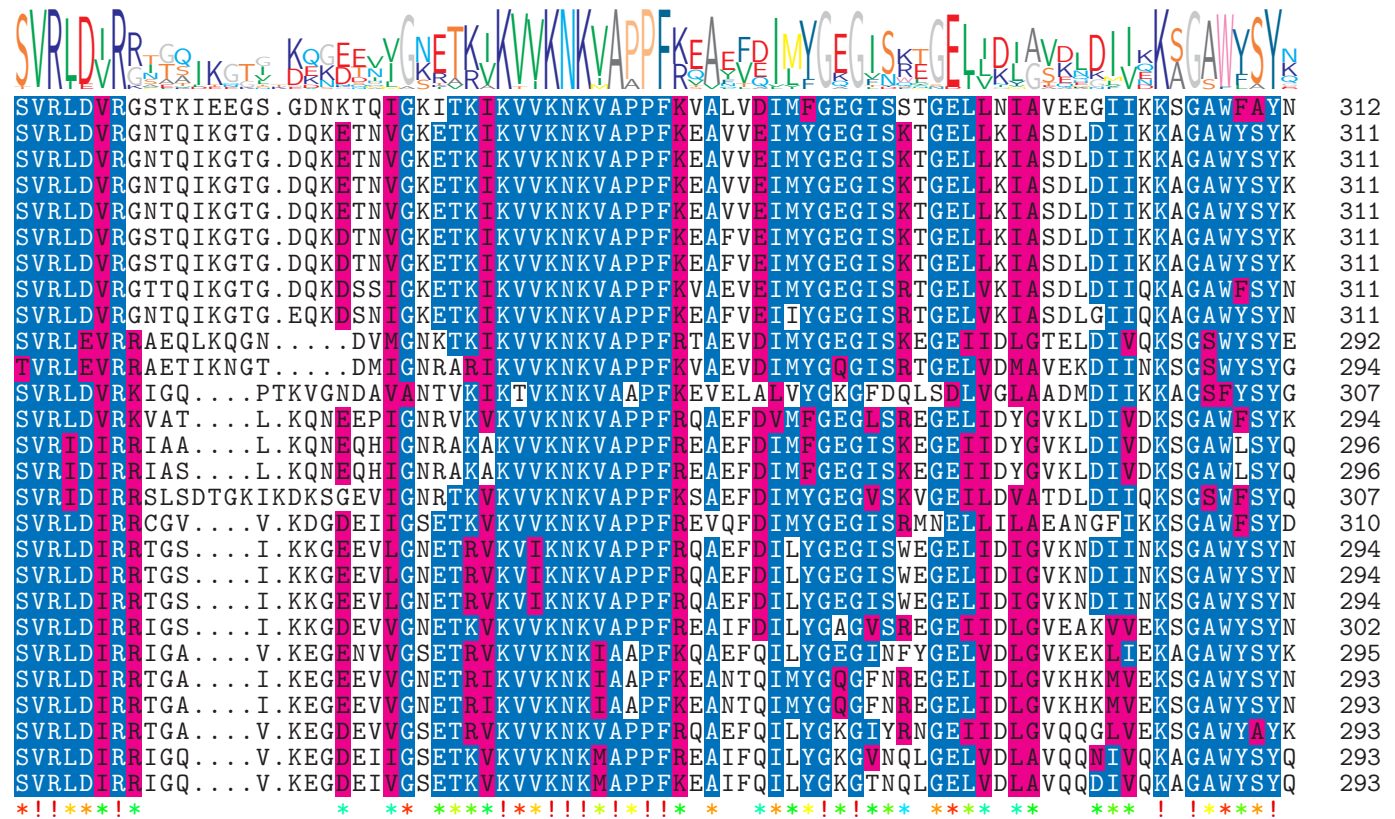


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NTDB	id	269	KZH43	RS08665	WP	001085462.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	GA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
NTDB	id	228	SPD	RS09265	WP	001085462.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	GA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
NTDB	id	194	SPR	RS08825	WP	001085462.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	GA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
NTDB	id	159	SP	RS09750	WP	001085462.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	GA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
NTDB	id	499	SM12261	RS08315	WP	078228268.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	GA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
NTDB	id	527	SMSK321	RS11065	WP	001085454.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	GA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
NTDB	id	453	Spy49	1753c	ACI	62003.1	VV	DSVAAL	VP	RA	EID	GD	IGDSH	VGLQAR	MS	QAMR	KL	SA	S	INKTKT	TA	IFIN	QLREK	VGV	M	FGN	PETTP	PGGR	ALKFYA	232			
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consensus

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NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKKDE	PKKEEA	.VNEEVPLDLGDELEIE		385
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NTDB id 1123 OK783 RS03845 WP 003688695.1	GAKIGQGS	KDNVRVWLKENPEIS	DEIDAKIRALNG	VEMH	ITEG	T...QDET	DGERPEE	348
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NTDB id 399490 GJE14 RS05785 WP 000963143.1	GEKIGQGS	KANATAWLKDN	PETAKEIEKKV	RELLLSNP	N....STPDFSV	...D...DSEG	VAETNEDF	353
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consensus		***!!!*****	***	*****	*****	***	***	

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NTDB id 600 KW2 RS01765 WP 011675413.1	...	387
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consensus

 non conserved
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 ≥ 50% conserved