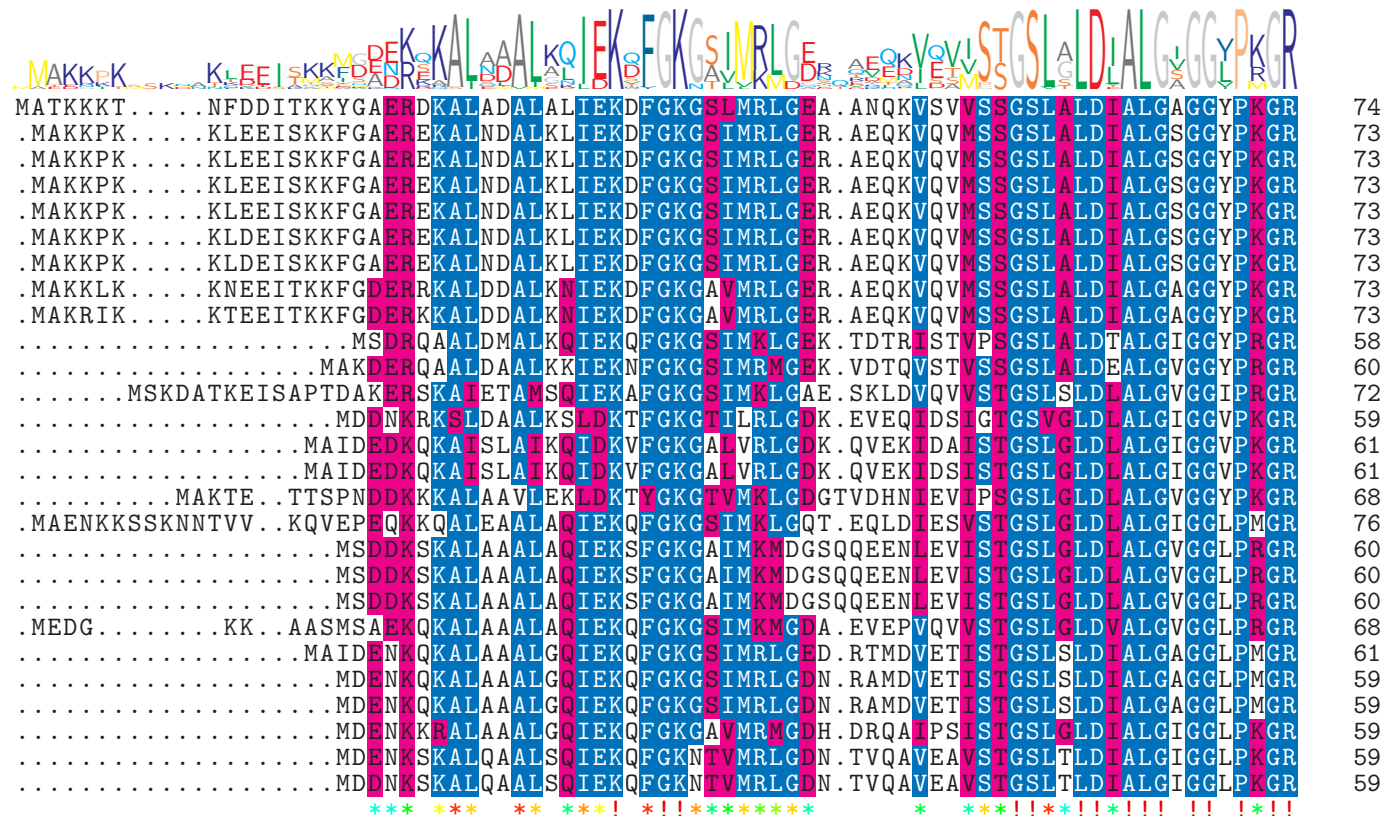


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consensus

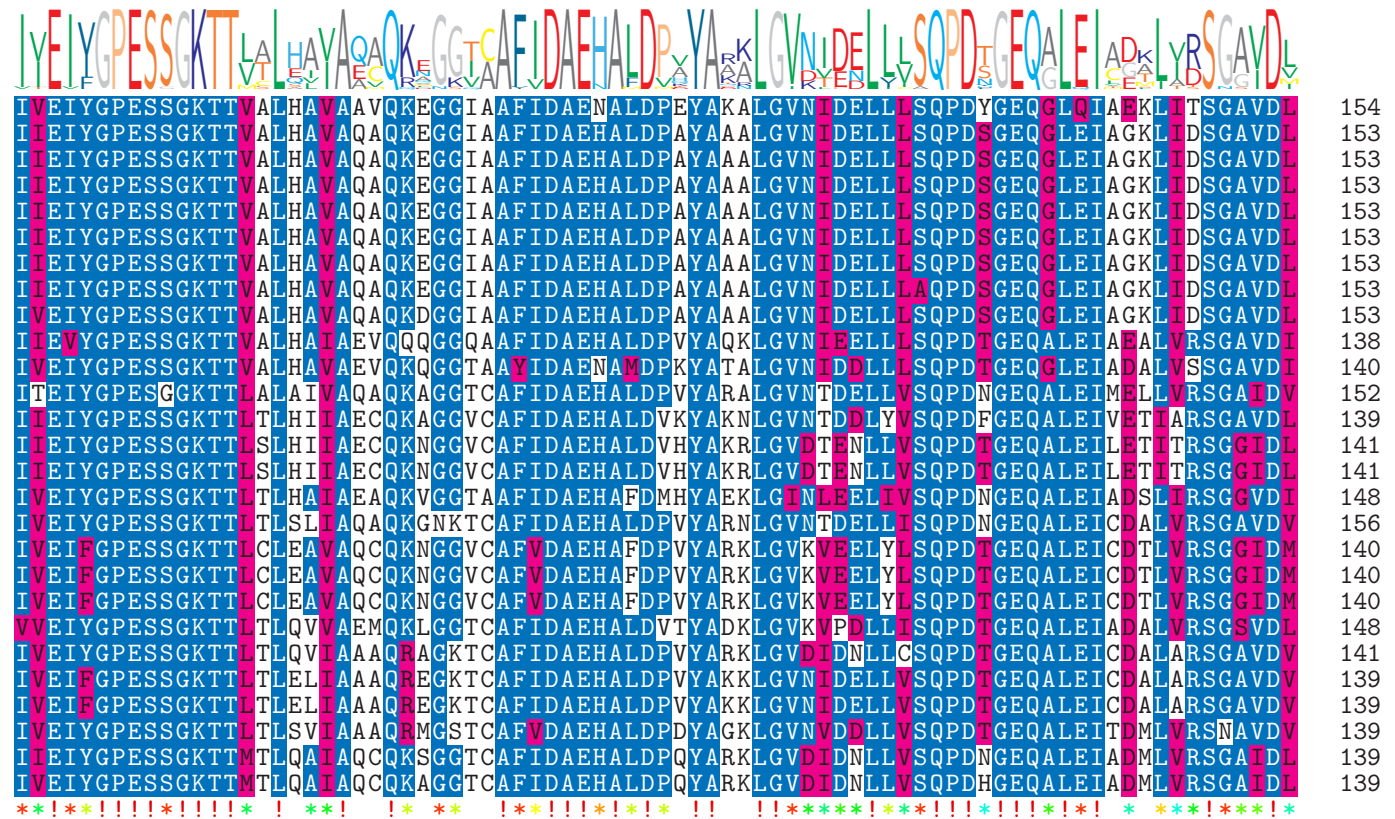
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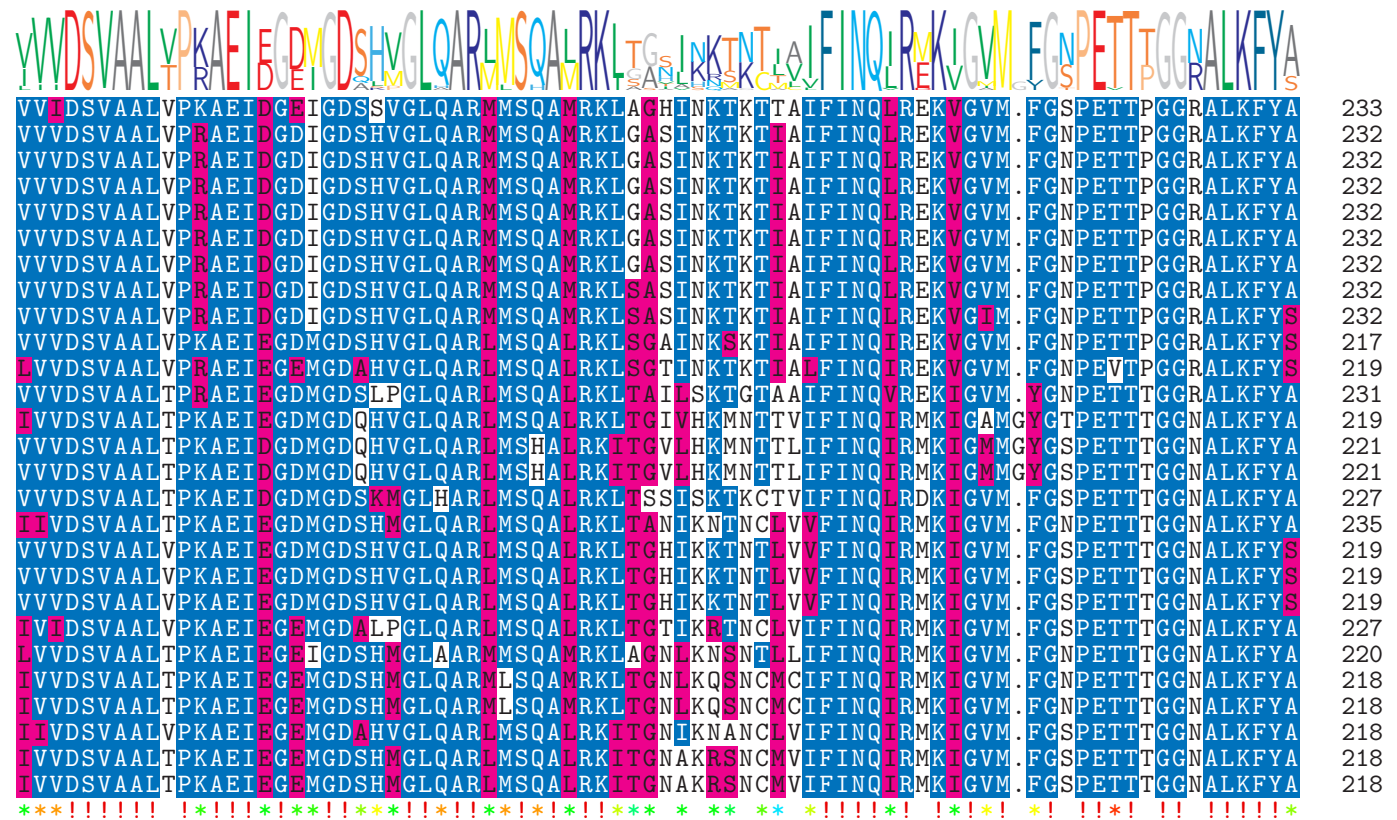
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consensus

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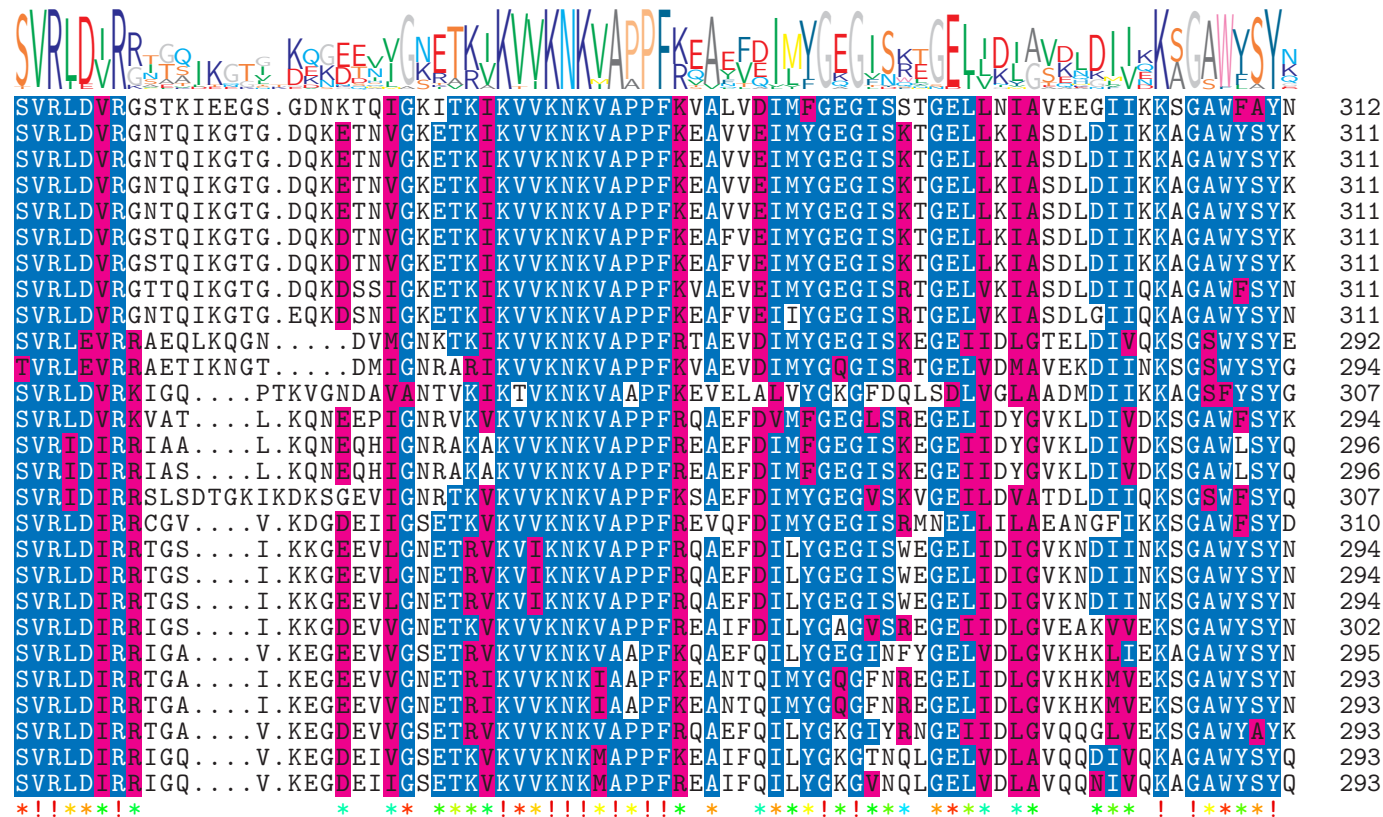
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
consensus



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NTDB id 1022 ACIAD RS06390 WP 004925639.1
consensus

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NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGQGA	EKAKNYLKEHQDV	FEIDHKVRAAHG	LDDSEVAETEEETT	ASKTKAKAKKEEK	AVETEEIELELED		387
NTDB id 269 KZH43 RS08665 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTEN	KKDE....PKKEEA	..VNEEVPLDLG	DELEIE	385
NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTEN	KKDE....PKKEEA	..VNEEVPLDLG	DELEIE	385
NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTEN	KKDE....PKKEEA	..VNEEVPLDLG	DELEIE	385
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NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGS	ENAKRYLADHPQLF	DEIDRKVRVKFGL	LEESEEEESAMAV	ASE.....E.....	TDDLALDLDN	GIEIE	377
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NTDB id 1313 DR RS12030 WP 010888966.1	DERIGQG	KEKTIAYIAERPE	MEQEIRDRVMAA	IRAGNAGEAPAL	APAPAAPEAAEA			363
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NTDB id 1334 RAOC RS04870 WP 004918236.1	DTKLGGQ	RDITVKELIKDN	PELAEELEAKIRE	KLQAK.....				343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	GTKIGQG	KNNAVKWLKENPE	IADKLEQEIRNTL	ASNPNS....	VLKATASDSN	DES.....DVEEF	..SEEEL.....	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	GAKIGQG	KDNVRVWLKENPE	ISDEIDAKIRALN	GVEMHI....	TEGTQ.....	DET	DG.ERPEE.....	348
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NTDB id 1165 A1552VC RS01565 WP 000344154.1	GDKIGQG	KANACKYLLKEN	PEIAKTLDKKL	REMLLNPNEM	QLIAETSS	..AADDVEFG	.AVPEEF.....	354
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	GNKIGQG	KNNVIRYLEENP	QISTEIEAVIRE	QLLTKASD....	QTAA.HDETE	EEEPDLLES.....		349
consensus	***!!!	*****	**	*****	*****	**	**	*

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		IEE	
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consensus			

- X non conserved
- X similar
- X ≥ 50% conserved