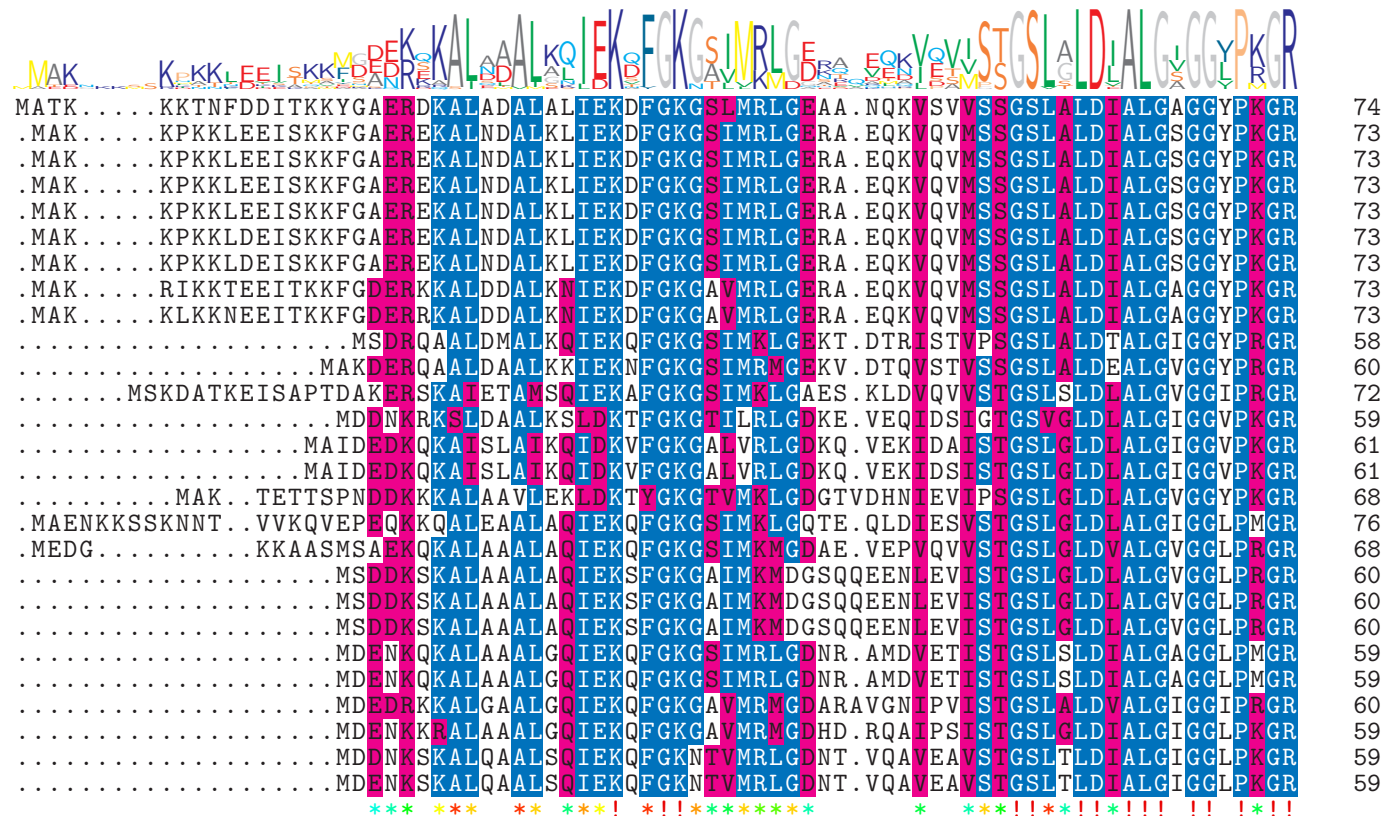


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consensus

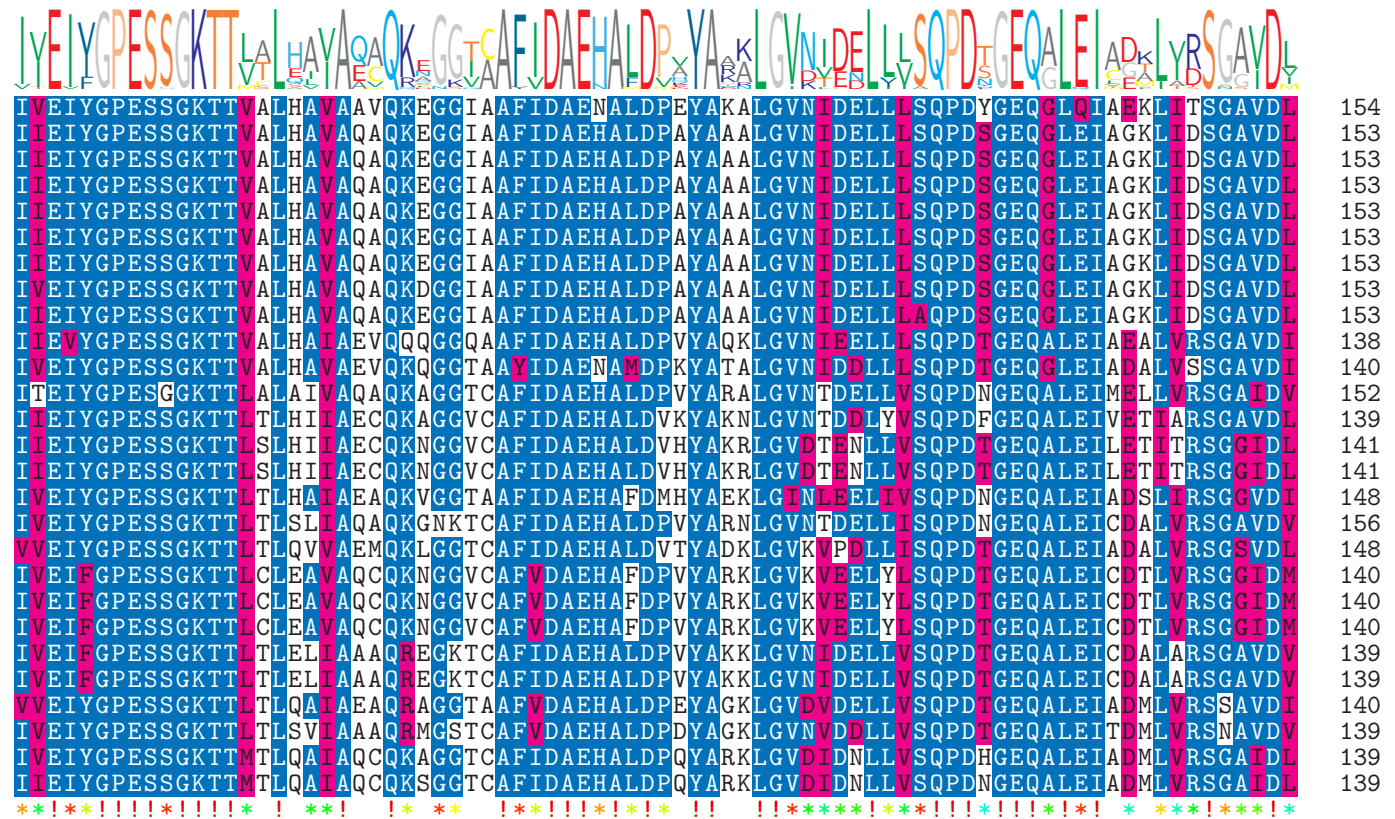
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consensus

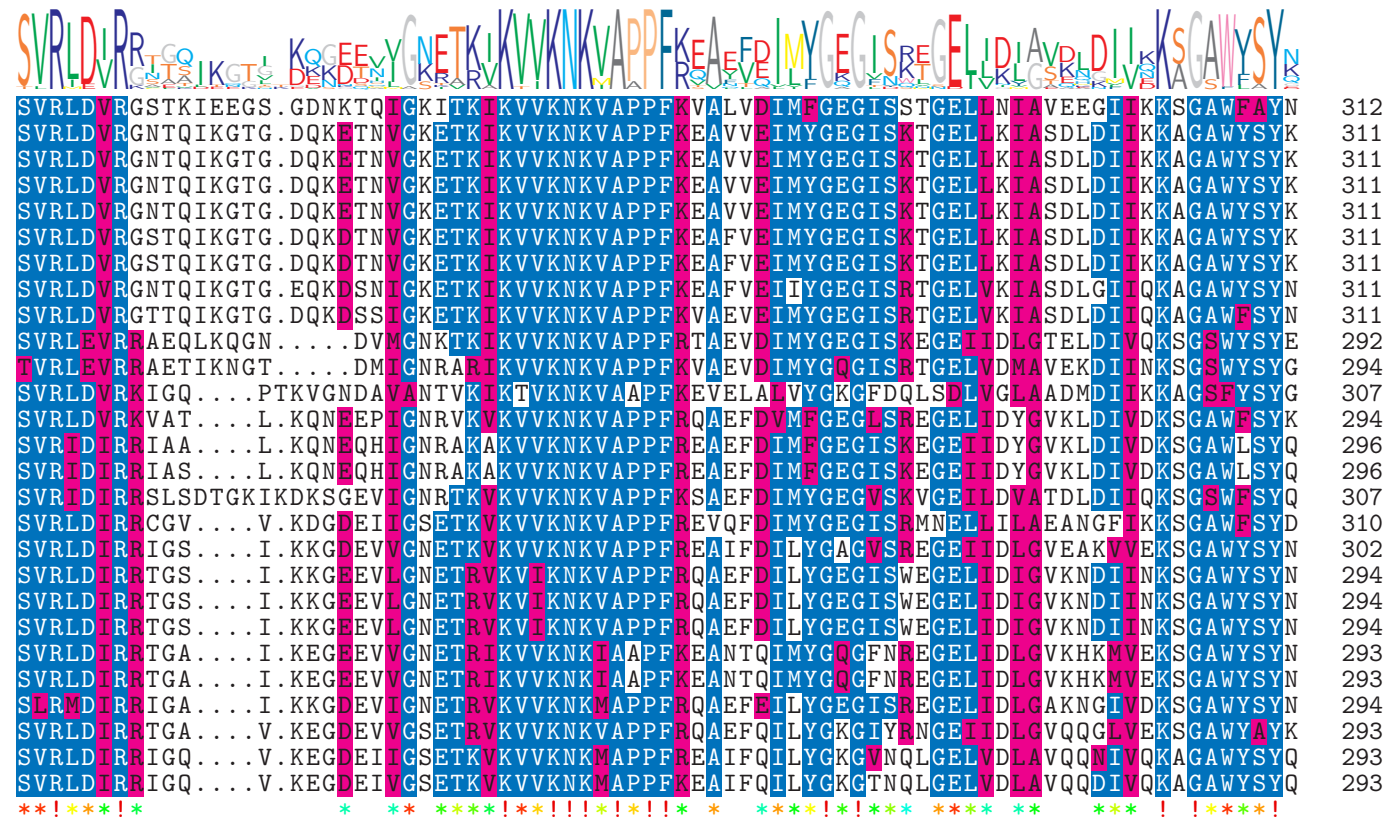
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGQGA	EKAKNYLKEHQDV	FDEIDHKVRAAHG	LDDSEVAETEEETT	ASKTKAKAKKKEEK	AVETEEIELELED		387
NTDB id 269 KZH43 RS08665 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTEN	KKDE....PKKEEA	.VNEEVPLDLGDE	LEIE	385
NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTEN	KKDE....PKKEEA	.VNEEVPLDLGDE	LEIE	385
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NTDB id 408 SMU RS09530 WP 002262392.1	GEKIGQGS	ENAKKFLADNPEIF	DDIDHKVVRVQYGL	IEEDNTEEKQSS	KEK.....ETDEK	.ADKNLVLELDDT	IELE	382
NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGS	ENAKRYLADHPQL	FDEIDRKVRVKFGL	LEESEESAMAV	ASE.....E.....	TDDLALDLDN	GIEIE	377
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NTDB id 625 LCA RS02525 WP 011374196.1	SERIGQGR	ENAKNYLADHED	VEDEVRLKVR	AAYGISDVPE	EDLPTTEDE.QI.NIL	PDDST.....EE		355
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	GTKIGQGG	KNNAVKWLKEN	PEIADKLEQEIR	NTLASNPNS....	VLKATASDSN	DESVDVEE...F	SEEL.....	372
NTDB id 1427 RS RS02760 WP 011000510.1	GERIGQGR	DNCREFLRENA	ELAREIENKVR	EHLGVTMPG....	AVTLAAEEVE	ED.....		352
NTDB id 1130 NGFG RS03960 WP 003688695.1	GAKIGQGG	KDNVRVWLKEN	PEISDEIDAKI	RALNGVEMHI....	TEGTQDETD.....	GERPEE.....		348
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	GDKIGQGG	KANACKYLKEN	PEIAKTLDKKL	REMLLNPNEM....	QLIA...ETSSA	ADDVE...FG	AVPEEF.....	354
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	GNKIGQGG	KNNVIRYFEEN	NQIAEEIERNI	REQLLT	TGTN....GAVQ	.I.EDEEEP	DLLE....S.....	349
consensus	***!!!	*****	***	*	***	*****	***	*

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consensus			

- non conserved
- similar
- ≥ 50% conserved