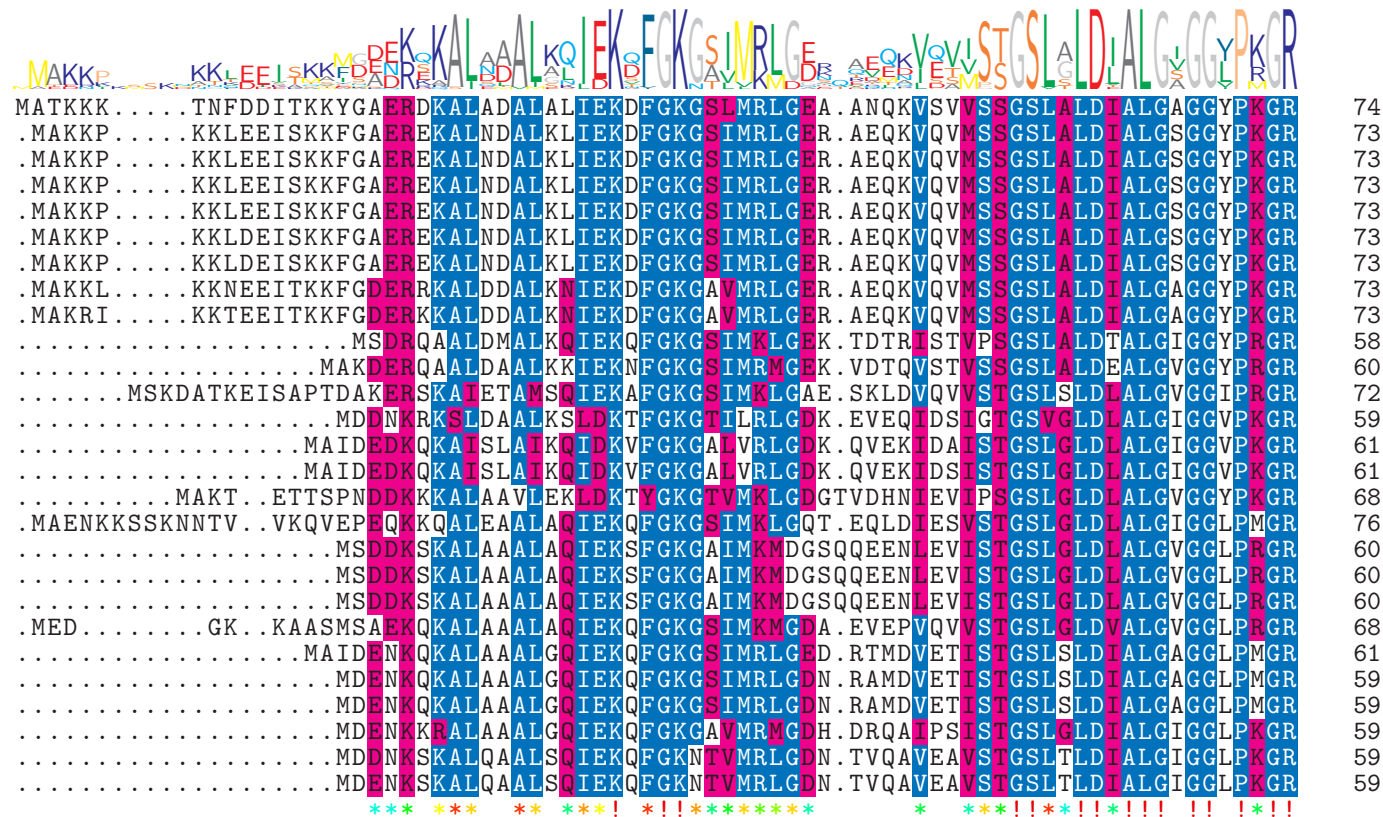


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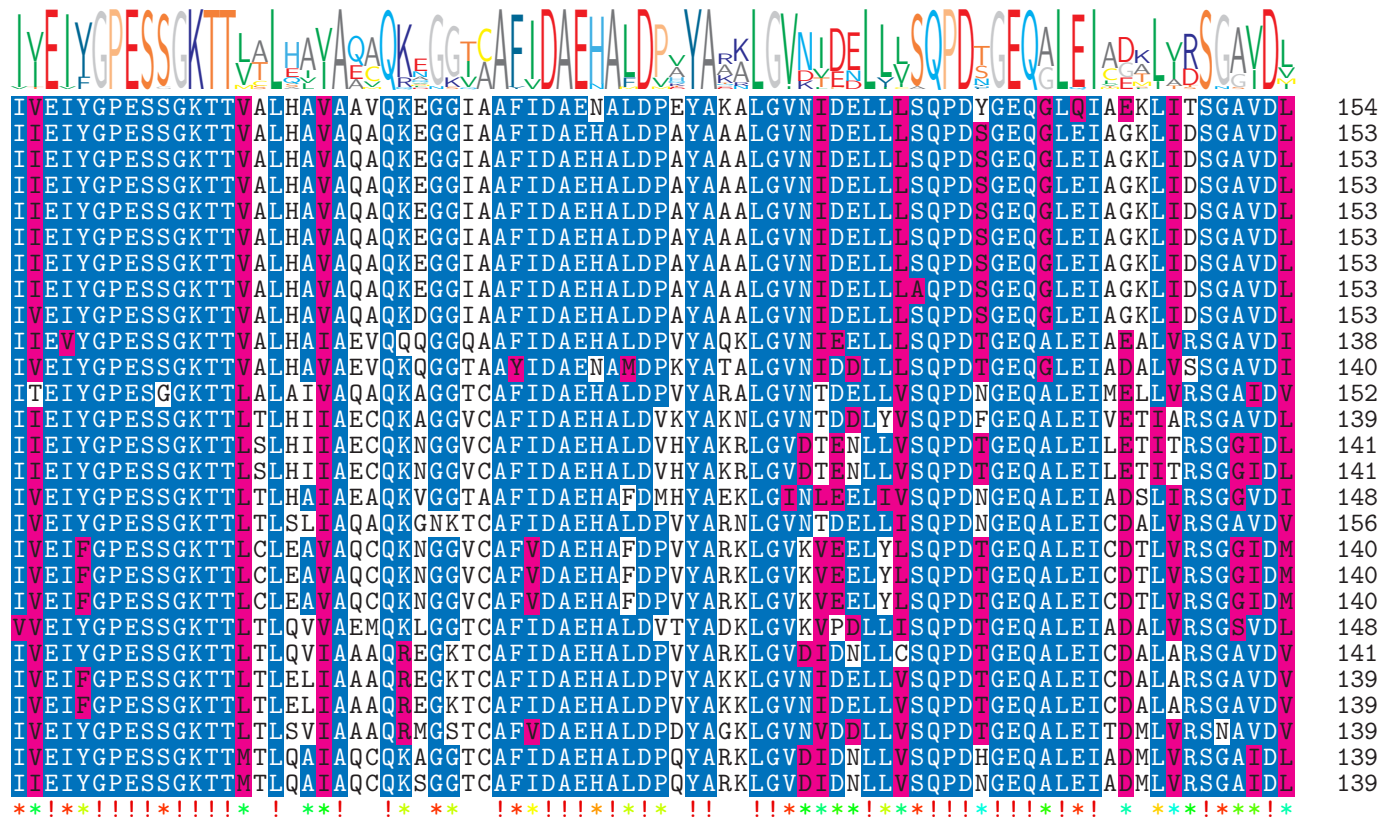
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus



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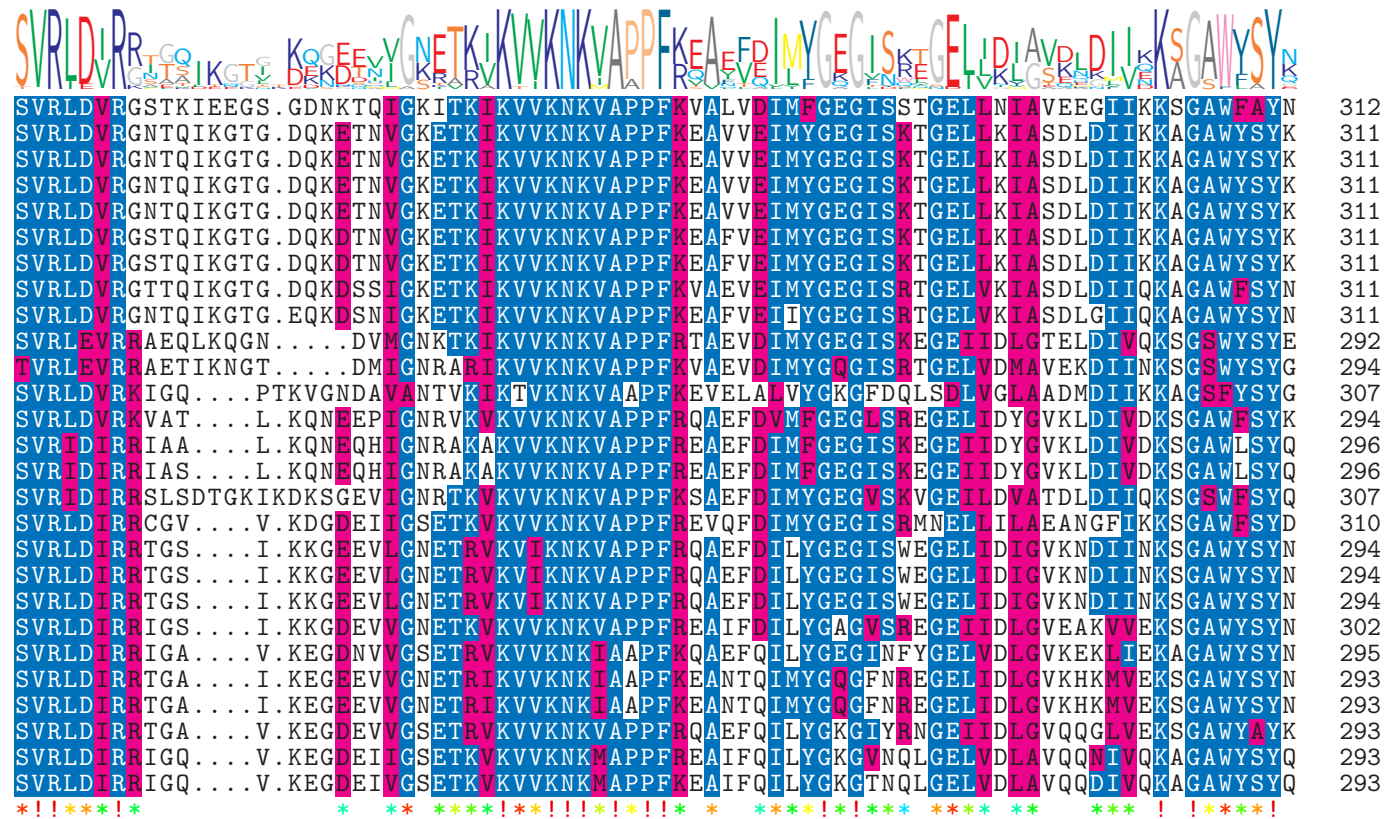
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NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKK	DE....PKKEEA..	VNEEVPLDLGDELEIE		385
NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKK	DE....PKKEEA..	VNEEVPLDLGDELEIE		385
NTDB id 159 SP RS09750 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKK	DE....PKKEEA..	VNEEVPLDLGDELEIE		385
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NTDB id 499 SM12261 RS08315 WP 078228268.1	DEKIGQGS	ENAKKYLAADNPEIF	DEIDHQVRVQFGL	IDGEEIAE....SKK	DE....AAQVDS..	VNEEVTLDLGDELEIE		381
NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGS	ENAKRYLADHPQLF	DEIDRKVRVKFGL	LEESEEEESAMAVASE	E.....	TDDLALDLNNGIEIE		377
NTDB id 408 SMU RS09530 WP 002262392.1	GEKIGQGS	ENAKKFLADNPEIF	DDIDHKVRVQYGL	IEEDNTEEEKQSSKEK	ETDEK..	ADKNLVLELDDTIELE		382
NTDB id 115 BSU 16940 NP 389576.2	EE	RLGQGREN	AKQFLKENKDIM	LMIQEQIREHYGL	DNNGVVQQAEETQEEL	EFEE.....		348
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	GT	KIGQGK	NNAVKWLKENPEI	ADKLEQEIRNTL	ASNPNSVLKATAS	.D...SNDES	DVEEFSEEE..L.....	372
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NTDB id 1427 RS RS02760 WP 011000510.1	GE	RIGQGRD	NCREFLRENAEL	AREIENKVRHL	GVTMPGAVTLAEE	.VE.ED.....		352
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	GD	KIGQGK	ANACKYLLKENPEI	AKTLDKKLREML	LNPNENMQLI..AE	TSSA	ADDVEFGAVPEEF.....	354
NTDB id 1205 PSJM300 05545 AFN77183.1	GSKIGQGK	ANAAKYLEENPEI	GQIEEQIRAKLL	VSANTKA..TP..	VA.EDLADADI			347
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	GN	KIGQGK	NNVIRYFEENN	QIAEEIERNIREQL	LTGTNGAV..Q...	I..EDEE	EPDLLES.....	349
consensus		***!!!*****	***	*****	*****	***	***	

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consensus

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