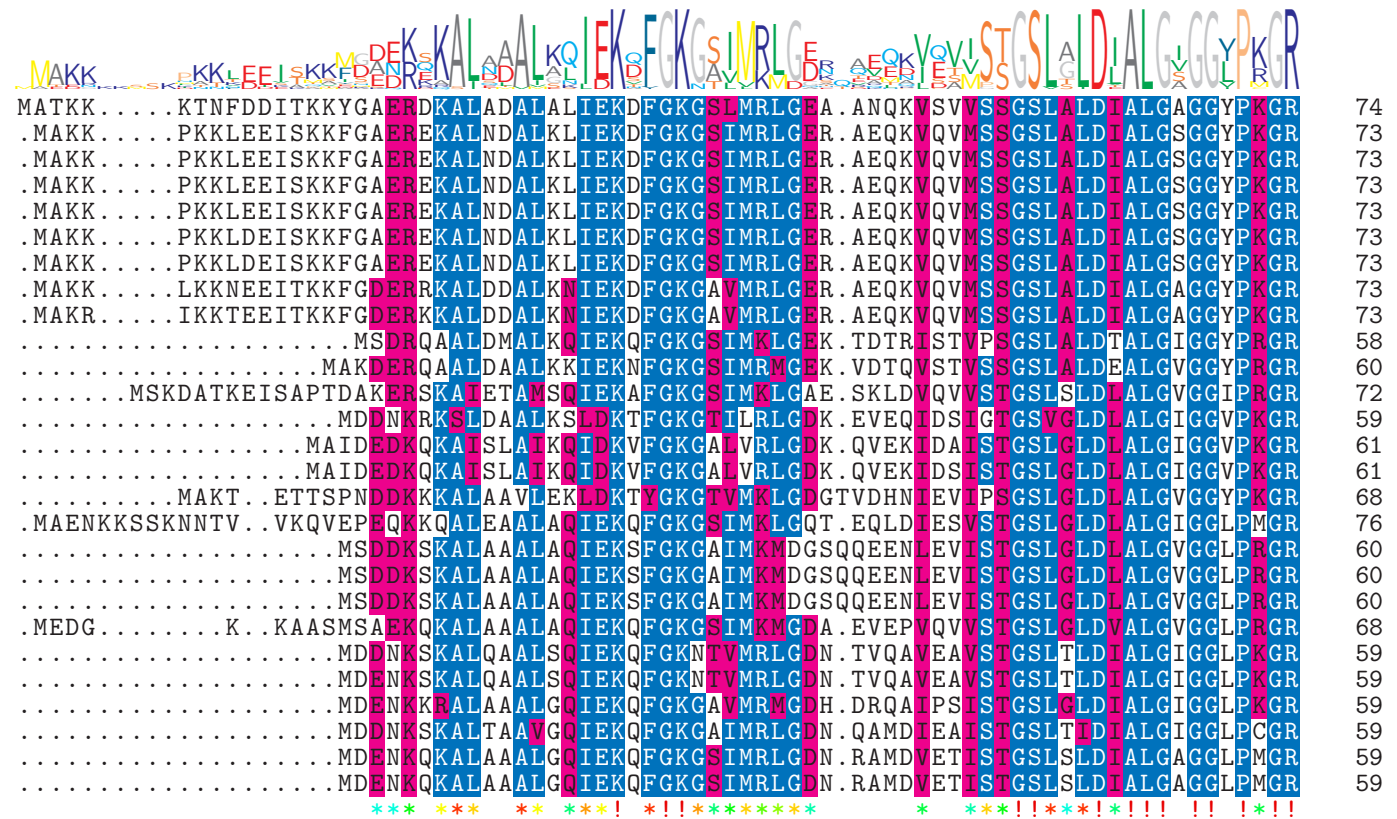


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consensus

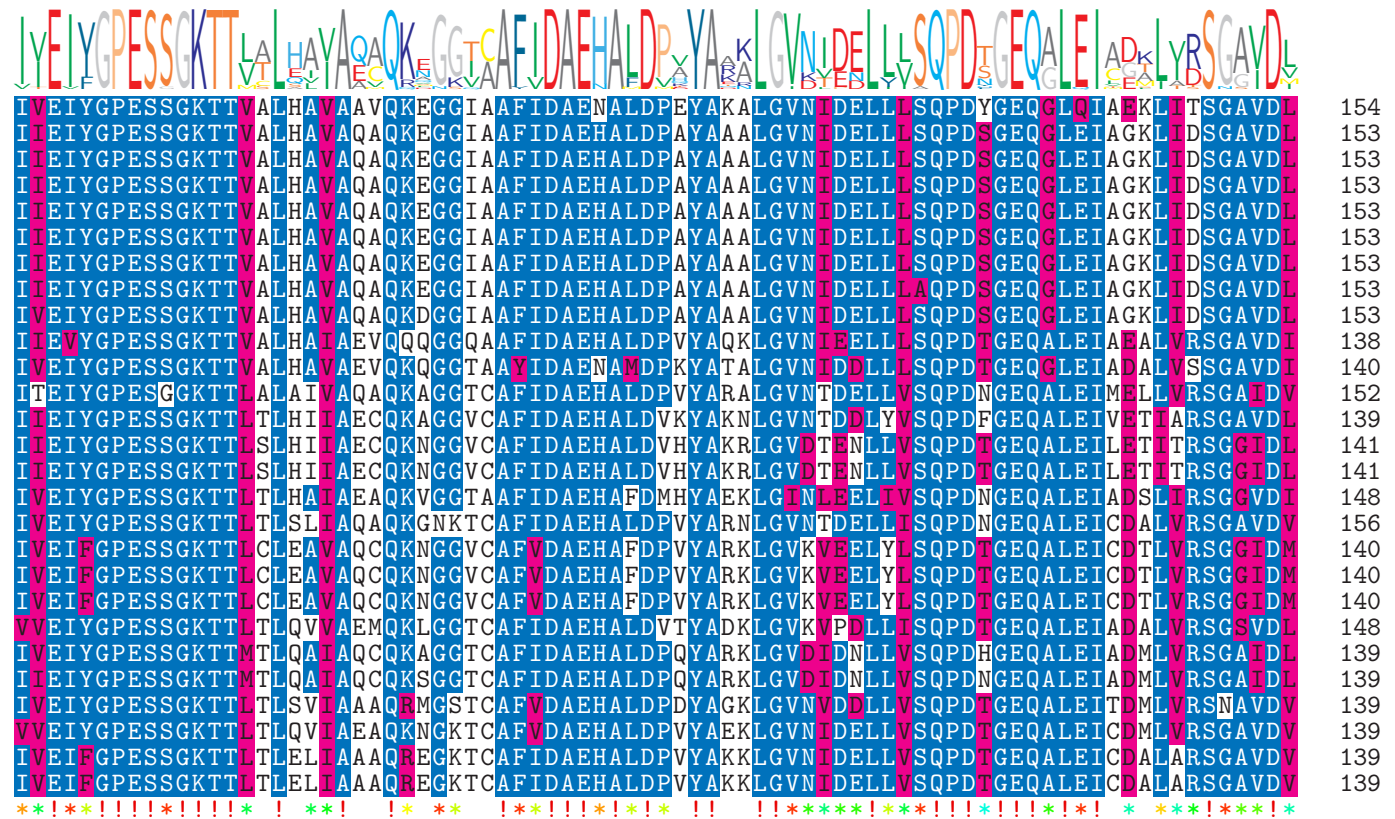
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consensus

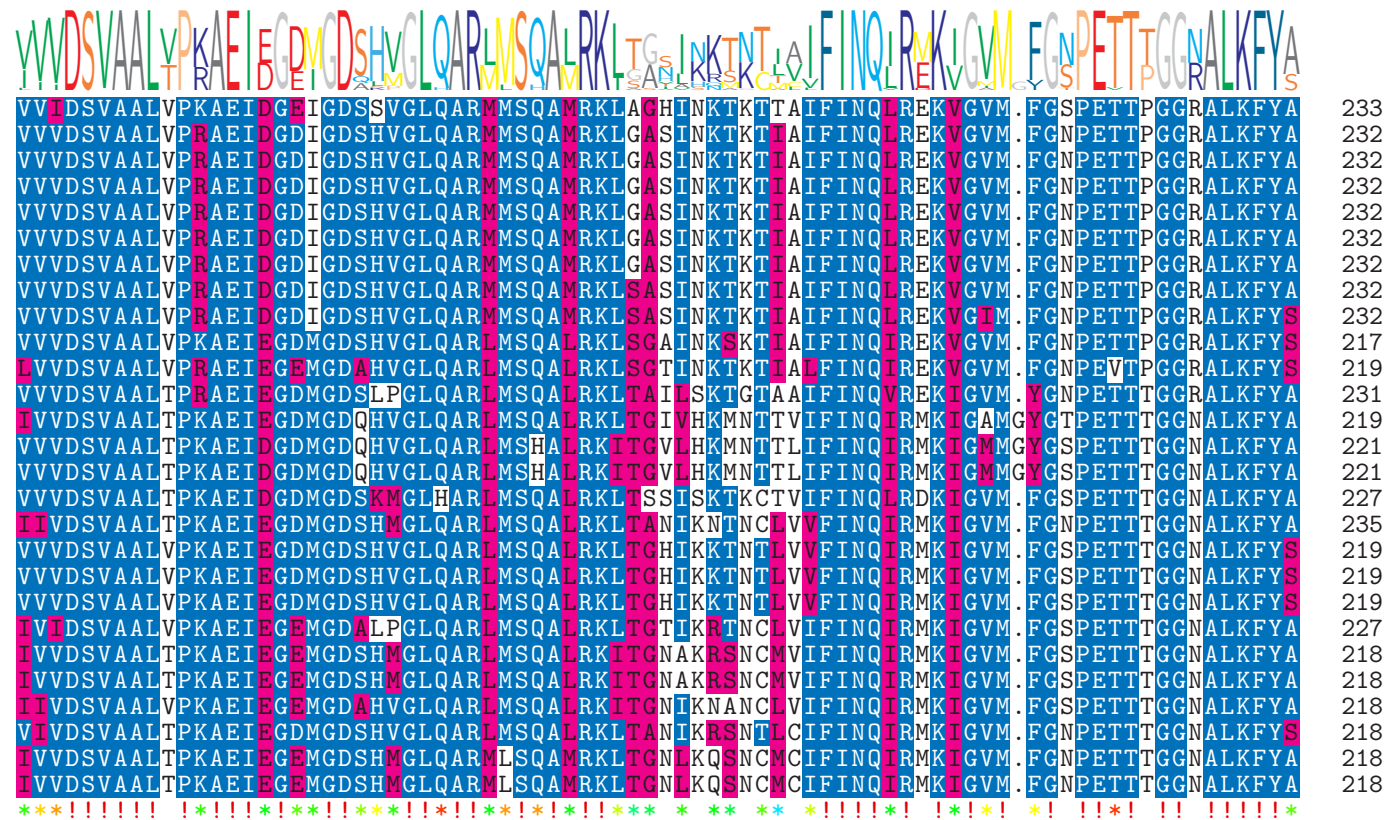
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NTDB id 1142 GTF74 RS11550 WP 000344154.1
consensus

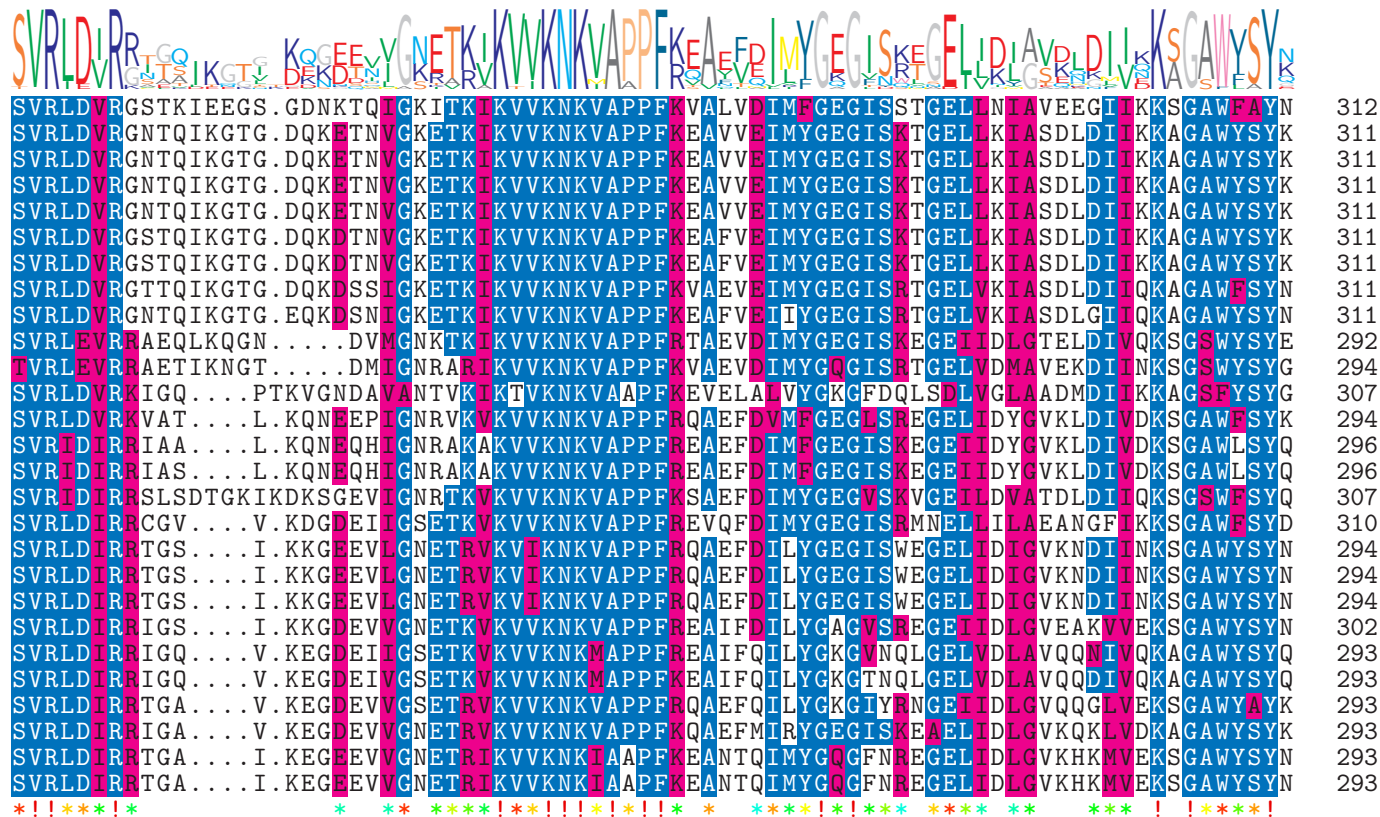
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NTDB id 1205 PSJM300 05545 AFN77183.1
NTDB id 370076 FJN13 RS03760 WP 012517447.1
NTDB id 1165 A1552VC RS01565 WP 000344154.1
NTDB id 1142 GTF74 RS11550 WP 000344154.1
consensus

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logo

NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGQGA	EKAKNYLKEHQDV	DEIDHKVRAAHG	LDDSEVAETEEEE	...TTASKTKAKAKKEEKAVETEEIELELED	387	
NTDB id 269 KZH43 RS08665 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	LIDGEEVSEQDTE	...NKKDE...PKKEEA..VNEEVPLDLGDE	381	
NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	LIDGEEVSEQDTE	...NKKDE...PKKEEA..VNEEVPLDLGDE	381	
NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	LIDGEEVSEQDTE	...NKKDE...PKKEEA..VNEEVPLDLGDE	381	
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NTDB id 499 SM12261 RS08315 WP 078228268.1	DEKIGQGS	ENAKKYLADNPEIF	DEIDHQVRVQFGL	LIDGEEIAE	...SKKDE...AAQVDS..VNEEVTLDLGDE	377	
NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGS	ENAKRYLADHPQLF	DEIDRKVRVKFGL	LEESEEEESAMAV	...ASE...E...TDDLALDLNG	373	
NTDB id 408 SMU RS09530 WP 002262392.1	GEKIGQGS	ENAKKFLADNPEIF	DDIDHKVRVQYGL	IEEDNTEEEKQSS	...KEK...ETDEK..ADKNLVLELDDT	378	
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NTDB id 625 LCA RS02525 WP 011374196.1	SERIGQG	RENAKNYLADHEDVE	DEVRLKVRVQYGL	ISDVPEEDLPTTE	...DE.QI.NILPDDST...EE...	355	
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NTDB id 1334 RAOC RS04870 WP 004918236.1	DTKLGGG	RDITVKELIKDNPE	LAELEAKIREKLQAK			343	
NTDB id 1389 A4U84 RS07560 WP 021112712.1	GTKIGQG	KNNVAVKWLKENPE	IADEIKLEQEI	RNTLASNPNSV	...LKATASDSNDES	VEEF...SEEEL...	372
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NTDB id 370076 FJN13 RS03760 WP 012517447.1	GD	RIGQGANVMKYLKEH	PETANEIETKIRQELL	LSKTVK	...AEEALPQGEDDV...L...PE...	348	
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	GD	KIGQGANACKYLKEN	PEIAKTLDKKL	REMLLNPENMQ	...LIAETSSAADDVEFGAV...PEEF...	354	
consensus	**!!!!***** ** * *** ***** ** **						

logo

NTDB id 600 KW2 RS01765 WP 011675413.1		387
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved