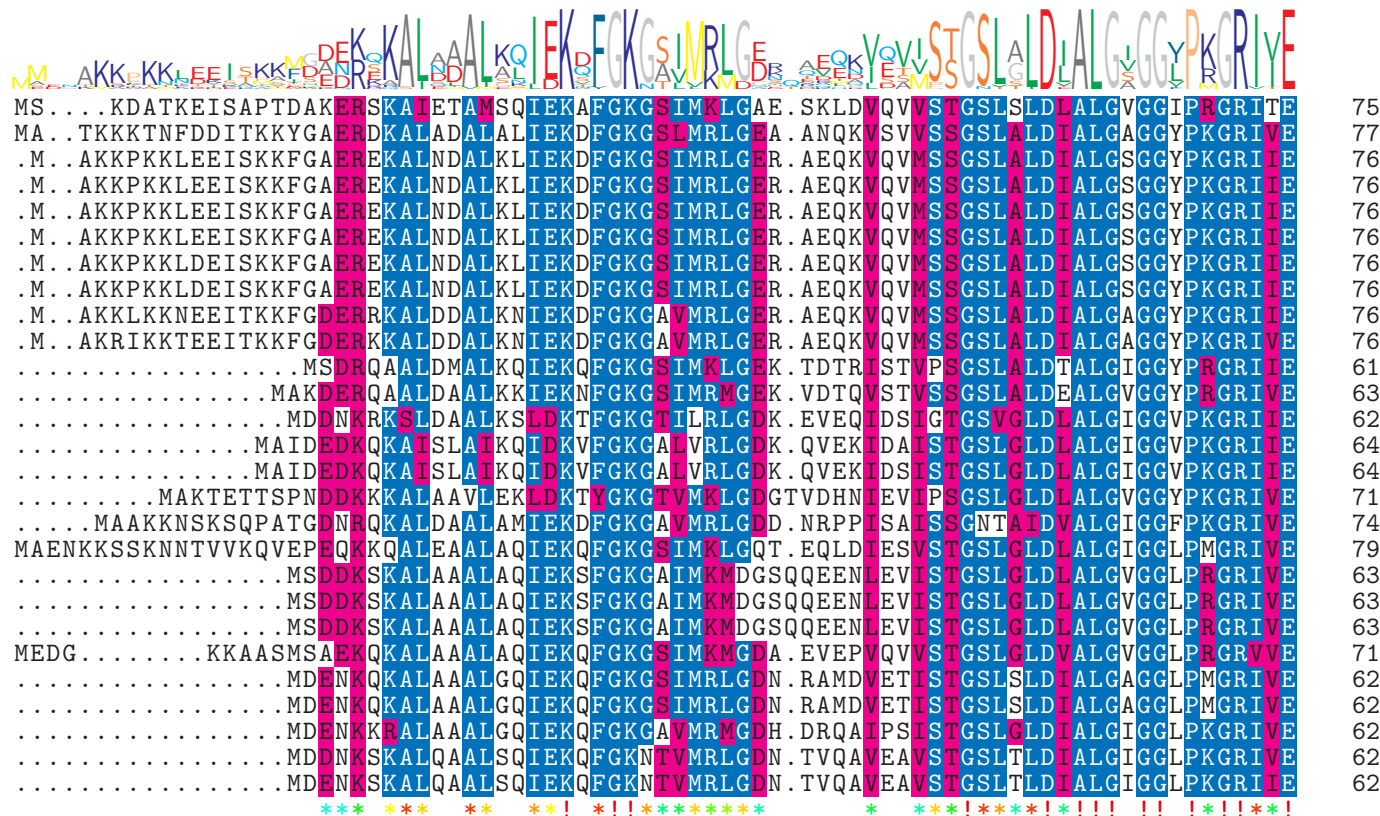


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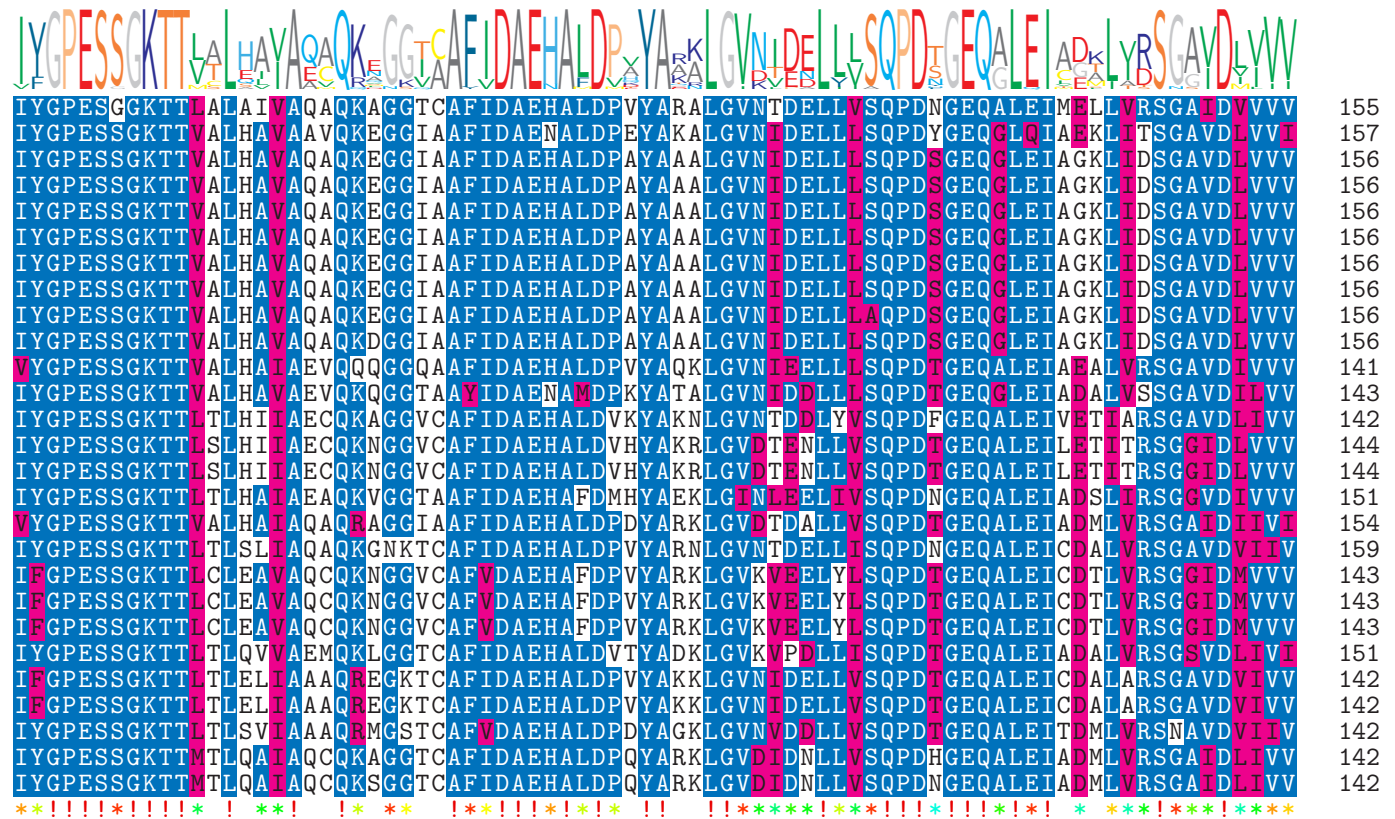
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consensus

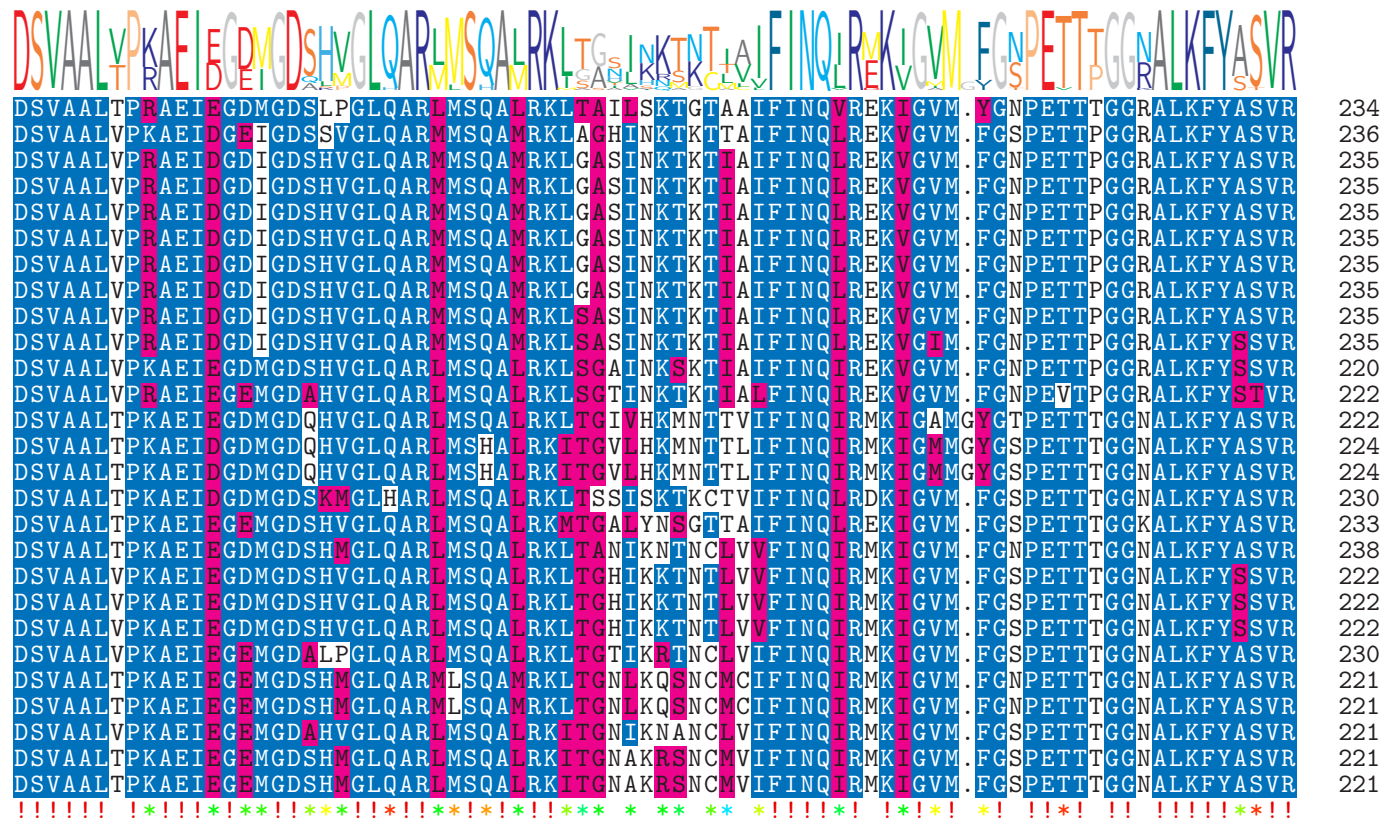
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consensus

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NTDB id 499 SM12261 RS08315 WP 078228268.1	LDVRGSTQIKGTG.DQKDTN	VGKETK	IKVVKNKV	APPFKE	AFVE	EIMY	GEGIS	KT	GELL	KIASDL	DI	IKKAGAWYSYKDEK	314			
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NTDB id 1334 RAOC RS04870 WP 004918236.1	IDIRRLSDTGKIKDKSGEV	IGNR	TKV	IKVVKNKV	APPFKS	AEFD	DIMY	GEG	VSKV	GE	ILD	VATDL	DI	IQKSGSWSYQDTK	310	
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	LDIRRCGV....V.KDGDEI	IGSET	KV	IKVVKNKV	APPFREV	QFD	DIMY	GEGIS	RM	NELL	I	LAENG	F	IKKSGAWFSYDGT	313	
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	LDIRRIGQ....V.KEGDEI	IGSET	KV	IKVVKNK	M	APPFKE	IFQ	ILY	GKGV	N	QL	GELVD	LAVQQ	D	IVQKAGAWYSYQGNK	296
consensus	***!*	*	**	*****!*****!*****!*	*	*****!*****!*****!*	*	*****!*****!*****!*	*	*****!*****!*****!*	*	*****!*****!*****!*	*			

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NTDB id 115 BSU 16940 NP 389576.2	LGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
NTDB id 625 LCA RS02525 WP 011374196.1	IGQGRENAKKNYLADHEDVEDEVRLKVRAAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
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NTDB id 1211 C694 RS00795 WP 000963128.1	LGQGRENAKALLKEDKALADEITLKIKEISIGSNEEIMPLP.....DEPLEE.....ME.....	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	LGQGRDITVKELIKDNPELAEELEAKIREKLQAK.....	343
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consensus	*!!!!***** ** * *** ***** ** ** *	

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