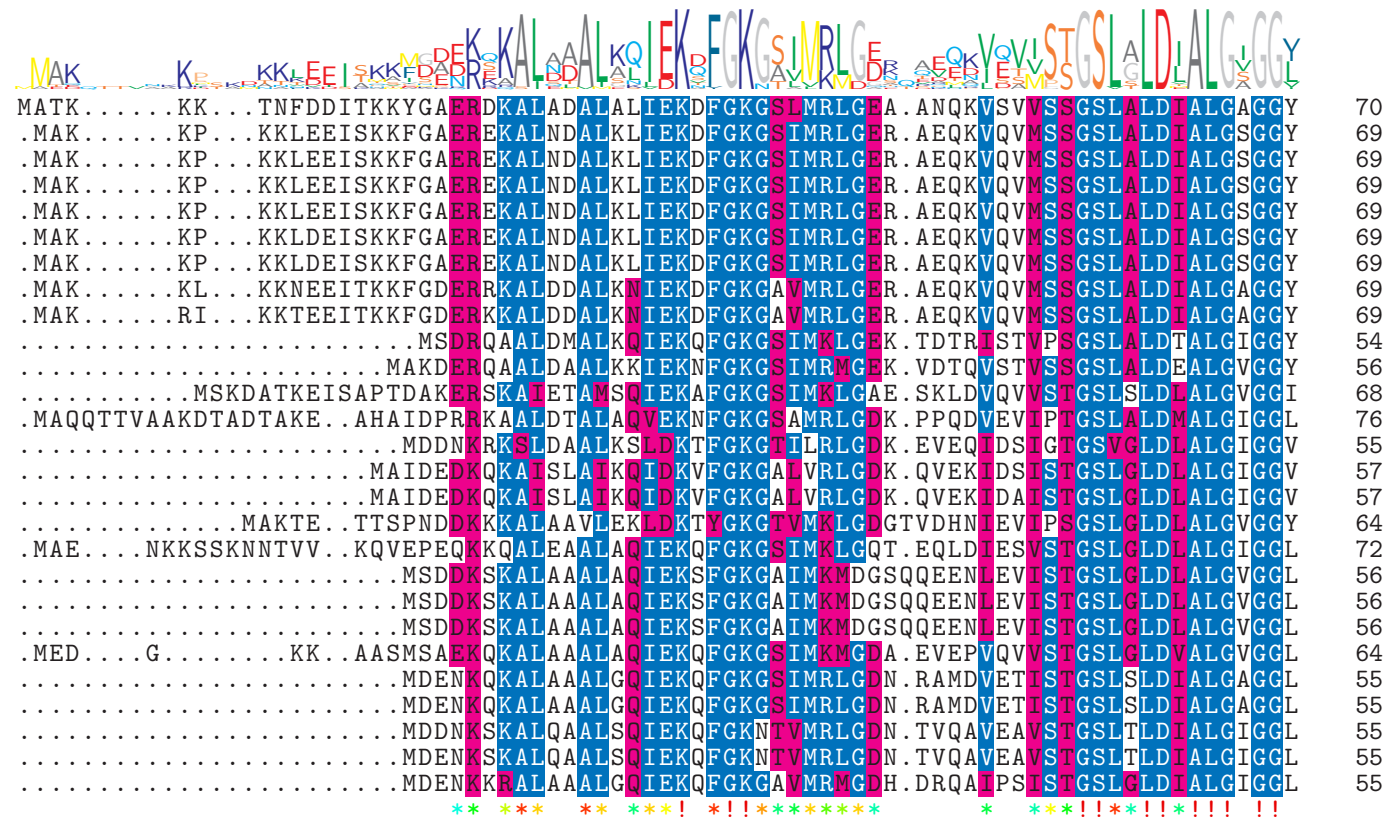


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consensus

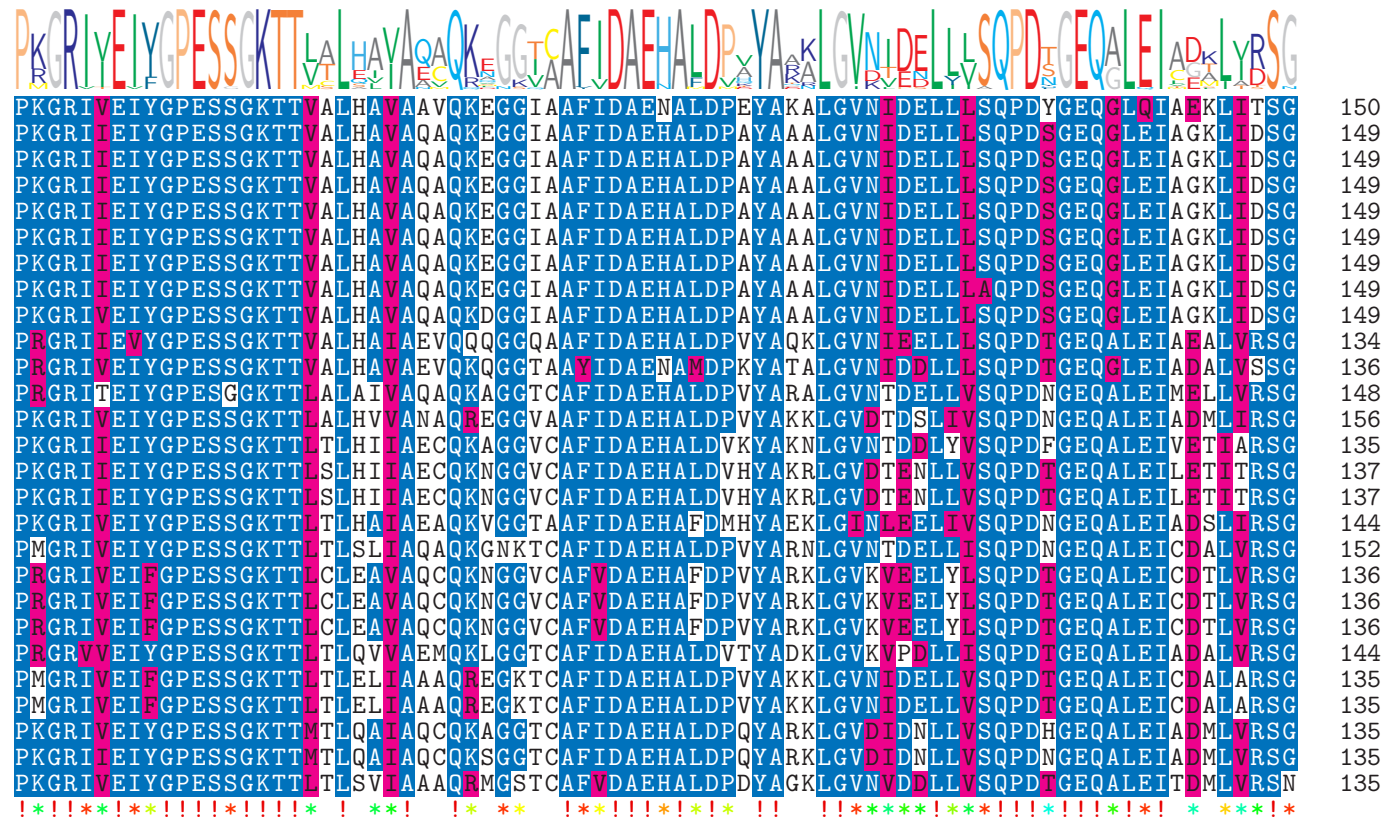
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AVD	LVV	DS	VAA	LP	KA	E	EG	MD	SH	VGL	QAR	MS	QAL	RKL	TG	S	INK	T	KT	IA	F	I	N	Q	L	R	E	K	V	G	V	M	.	F	G	S	P	E	T	T	P	G	G	R	A	L																			
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AVD	LVV	DS	VAA	LP	RA	E	D	G	I																																																								

logo

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NTDB id 342612 ESN35 RS03830 WP 033509171.1	KFYASVRLDIRRVQT....	L.KSGDEAVGNRTRVKVVKNKMAPPFK	SAEFDVLYGEGISKEGSVLDMALQC	NVVKKSGSW	310					
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consensus	!!!**!!**!!*	*	**	*****!*****!!*!!*!!*	*	*****!!*!!**	*****	**	***	!!**

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NTDB id 194 SPR RS08825 WP 001085462.1	YSYKDEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKK	DE.....PKKEEA..VNEEVPLDLGD	380	
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NTDB id 408 SMU RS09530 WP 002262392.1	FSYNGEKIGQGS	ENAKKFLADNPEIF	DDIDHKVRVQYGL	IEEDNTEEKQSSKEK	ETDEK..ADKNLVLELDD	377	
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NTDB id 1334 RAOC RS04870 WP 004918236.1	FSYQDT	KLGGGR	DTVKELIKDNPEL	AEELEAK	IREKLQAK.....	343	
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consensus	**!	**!!!*****	**	*****	*****	**	*

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NTDB id 499 SM12261 RS08315 WP 078228268.1	ELEIEIEE		384
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consensus			

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