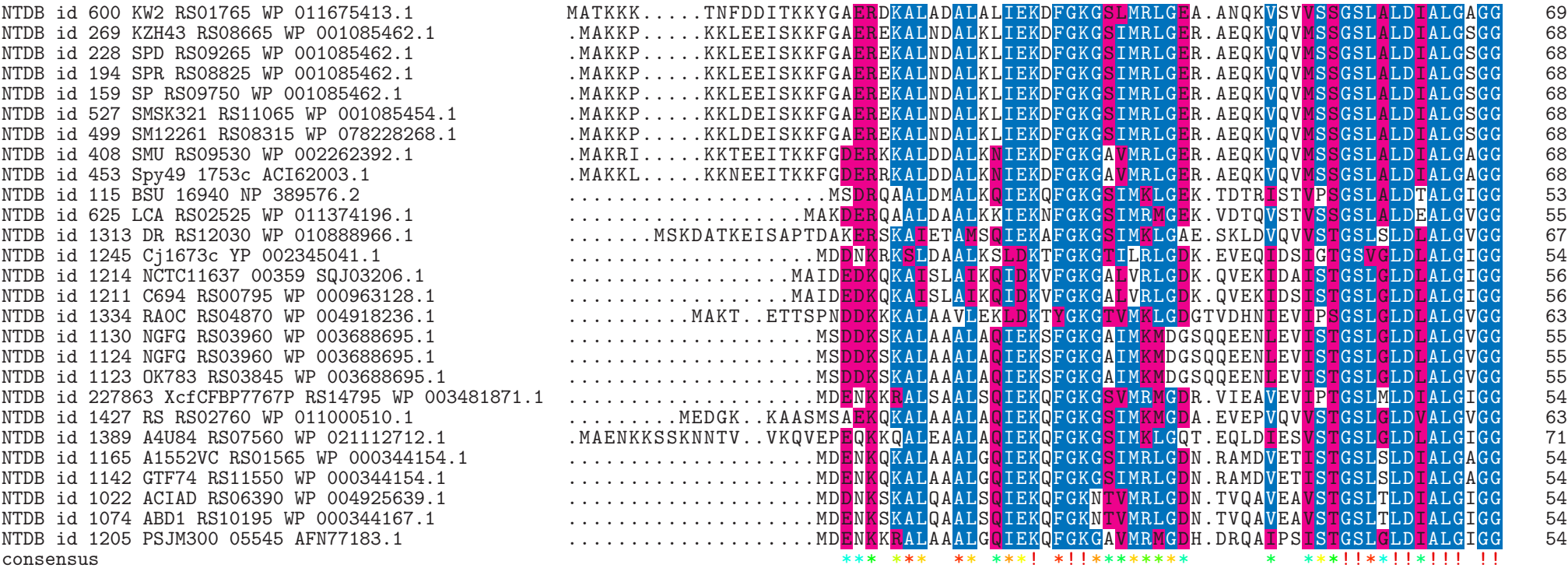


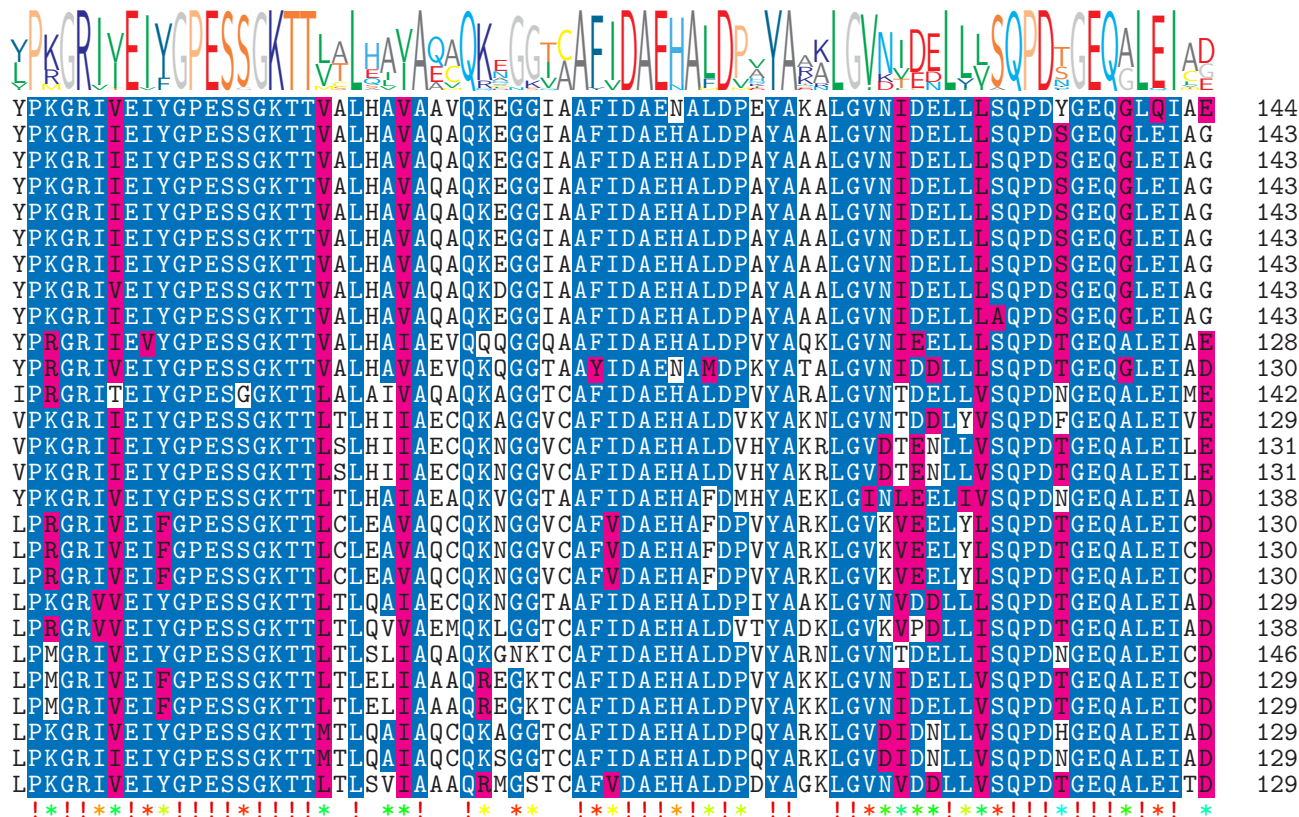
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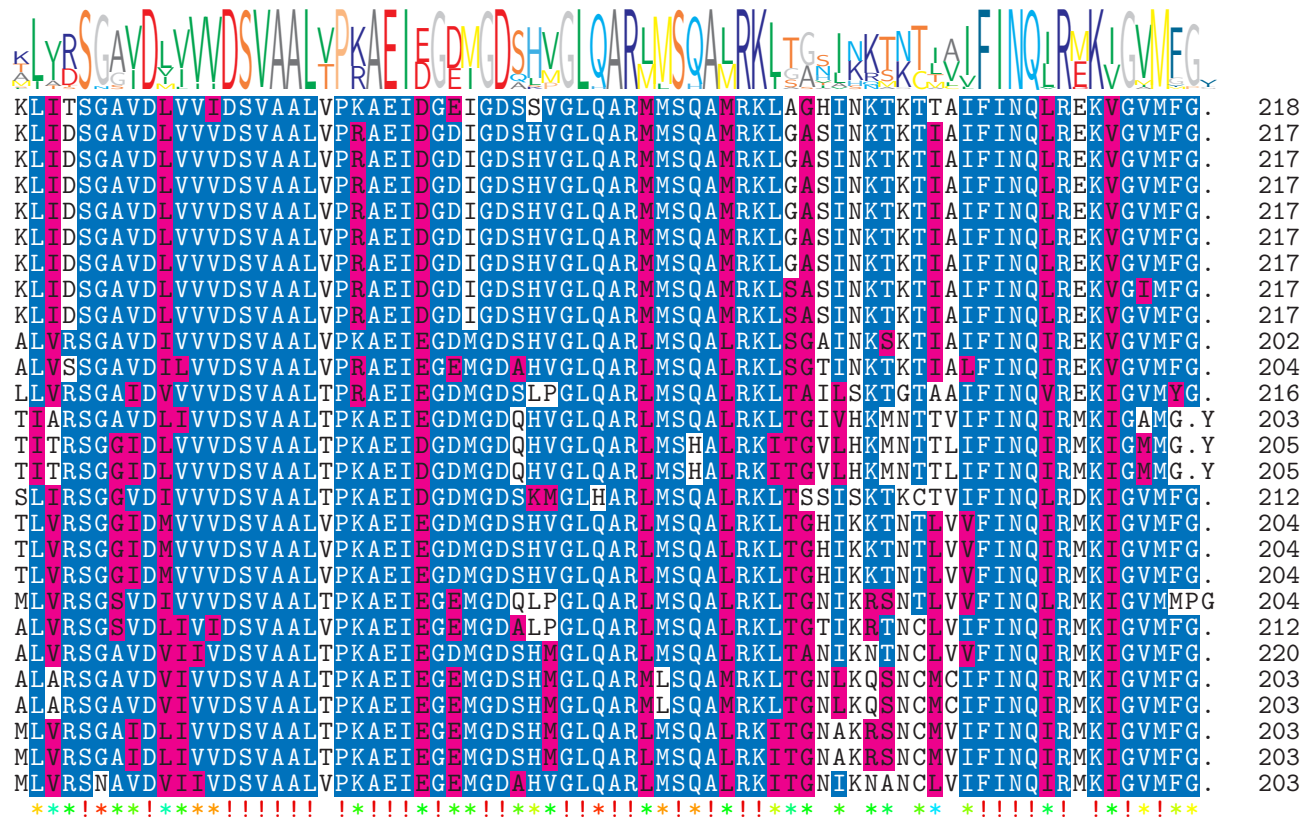


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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

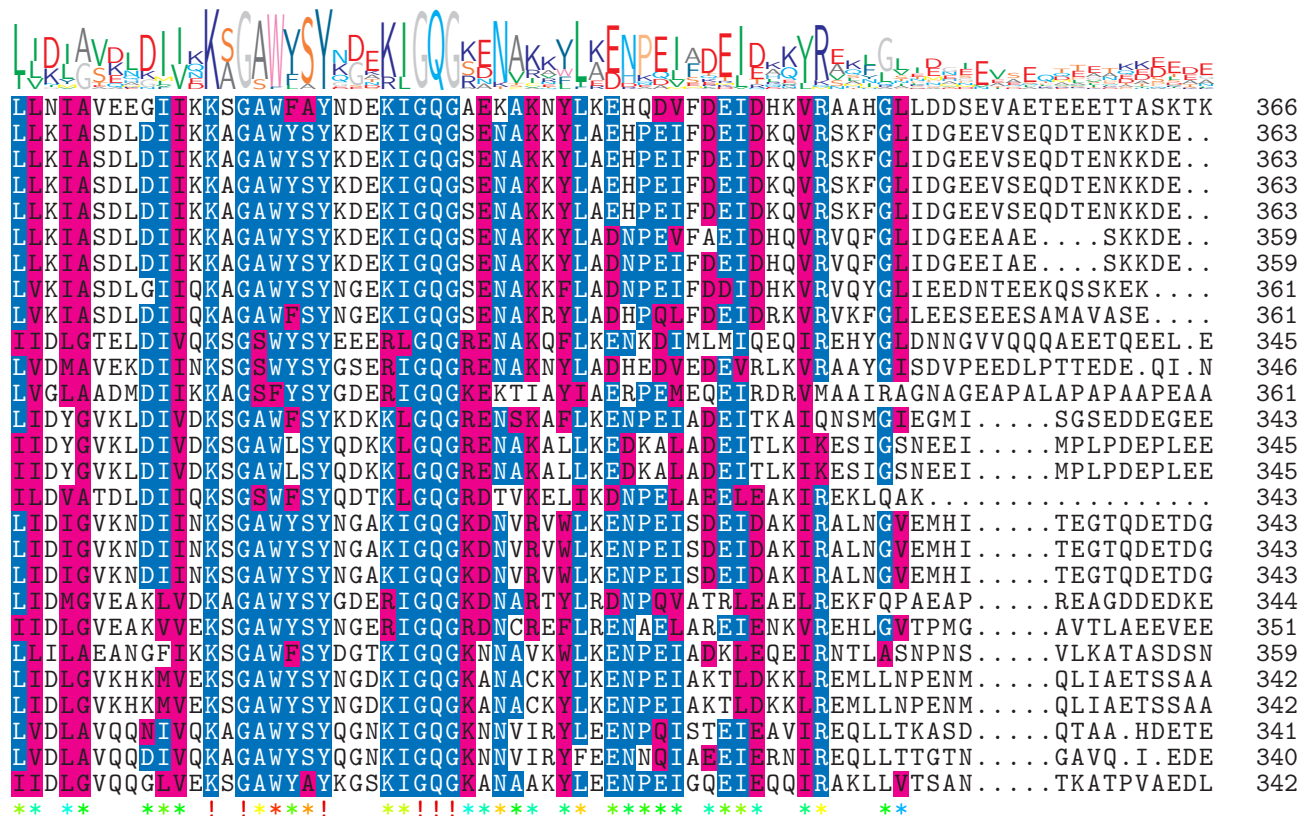
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logo

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NTDB id 625 LCA RS02525 WP 011374196.1	.NPEVTPGGRALKFYSTVRLDVR	RAETIKNGT.....DM	GNRARI	KVVKNKVAPPF	KVAEVDIMY	CGGIS	RTGE	273	
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NTDB id 1245 Cj1673c YP 002345041.1	GTPETTTGGRALKFYASVRLDVR	KVAT....L.KQNE	EP	GNRVK	KVVKNKVAPPF	RQAEFDVM	FGEGLS	REGE	273
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NTDB id 1334 RA0C RS04870 WP 004918236.1	.SPETTTGGRALKFYASVRLDIR	SLSDTGKIKDKSGE	IGNRTK	KVVKNKVAPPF	KSAEFDIMY	GEGVSK	VGE	286	
NTDB id 1130 NGFG RS03960 WP 003688695.1	.SPETTTGGRALKFYSSVRLDIR	TGS....I.KKG	EEVLGN	ETRVKV	IKNKVAPPF	RQAEFDILY	GEGISWE	GE	273
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NTDB id 1205 PSJM300 05545 AFN77183.1	.SPETTTGGRALKFYASVRLDIR	TGA....V.KEG	DEVVGS	ETRV	KVVKNKVAPPF	RQAEFDILY	GKGIY	RNGE	272
consensus	!!*!!	!!!!!!*	!!*	!!*	!!*	!!*	!!*	!!*	!!*

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consensus



logo

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NTDB id 527 SMSK321 RS11065 WP 001085454.1	..VAQVDS..VNEEVTLDLGDELEIEIEE	384
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved