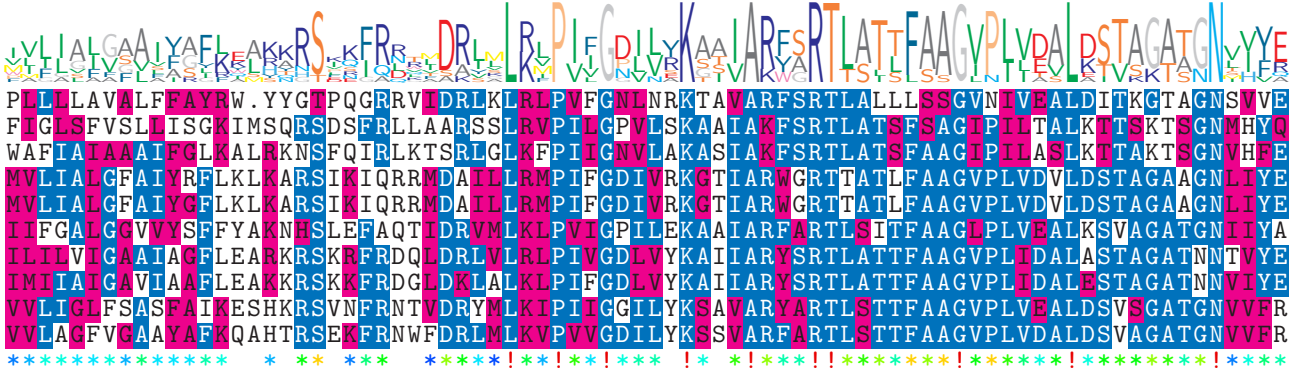


NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.L	VEATIEAEDLRTAARLRDRGLFVAETKEPGKG	LQAEVRIPALER	GPGLK	65
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NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLPLKNYRWKGINSGKK	VSGQMLAISEIEVRDKLKDQHIQIKL	KKGSVS	LLAR.....LTHRVK	68
NTDB id 1113	AAA85695.1 219..1451()	.MAKNGGFSLFAKKEKRFIFEGRHSADKL	VNGVEVSAFTEEEARKKLAKRGIRPLQIT	RVK....T...SSKR	KITQE	70
NTDB id 1112	NGFG RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRFIFEGRHSADKL	VNGVEVSAFTEEEARKKLAKRGIRPLQIT	RVK....T...SSKR	KITQE	70
NTDB id 1252	GCD85 RS07730 WP 011213805.1	.MDK.....NSPPLTLTFHYQGINKAGQK	MEGDIQARSLATAKADLRKQGIVTNK	VVKRKP	LFDR....KNKKITQA	67
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAK.....KTQVMPVFAYEGVDRKGIK	LKGELPARNMALAKVTLRKQGITIRTIRE	KRKNIFEG...LLK	KKVKPL	69
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVK.....KAQMMPTFAYEGVDRKGVK	IKGELPAKNMALAKVTLRKQGVTVRNIRE	KRKNILEG...LFK	KKVTTL	69
NTDB id 37195	G655 RS23370 WP 010792214.1	.MAD.....KALKTCVFWEGTDKKGAK	VKGELAGQNTMLVKAQLRKQGINPLKVR	KKGITLLG.....KG	KRVKPM	66
NTDB id 1198	PSJM300 03950 AFN76868.1	.MAQ.....KAIKNSVFTWGLDRQGA	IKGELSCVSPALVKAQLRKQGINPLKVR	KKGISLFS.....AG	KKIKPM	66
consensus		***	* * * * *	* * * * *	* * * * *	66

[illegible][illegible]

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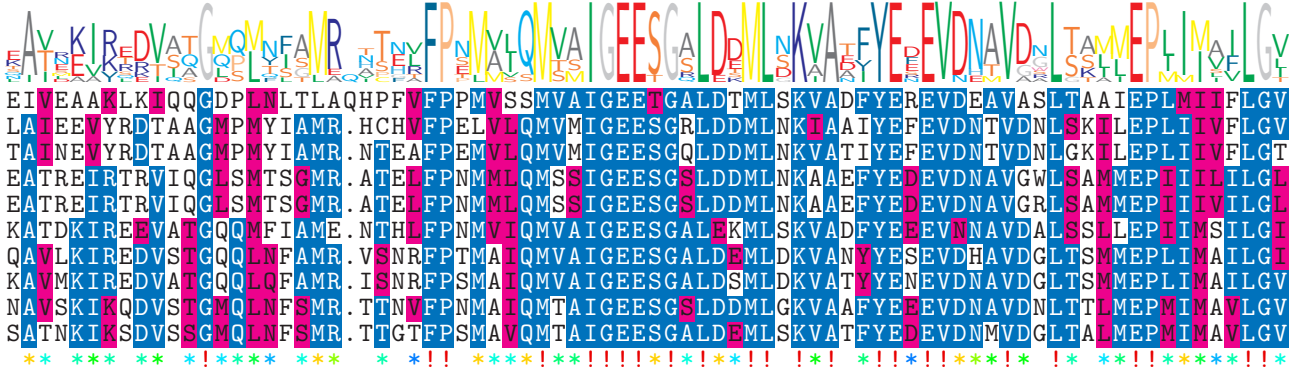
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NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 37195 G655 RS23370 WP 010792214.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus



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NTDB id 1029 TT RS02230 WP 011228203.1
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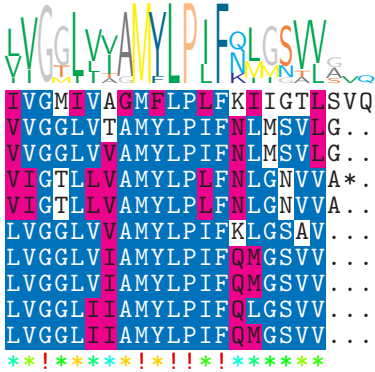


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NTDB id 37195 G655 RS23370 WP 010792214.1
NTDB id 1198 PSJM300 03950 AFN76868.1
consensus



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☒ non conserved
☒ similar
☒ ≥ 50% conserved