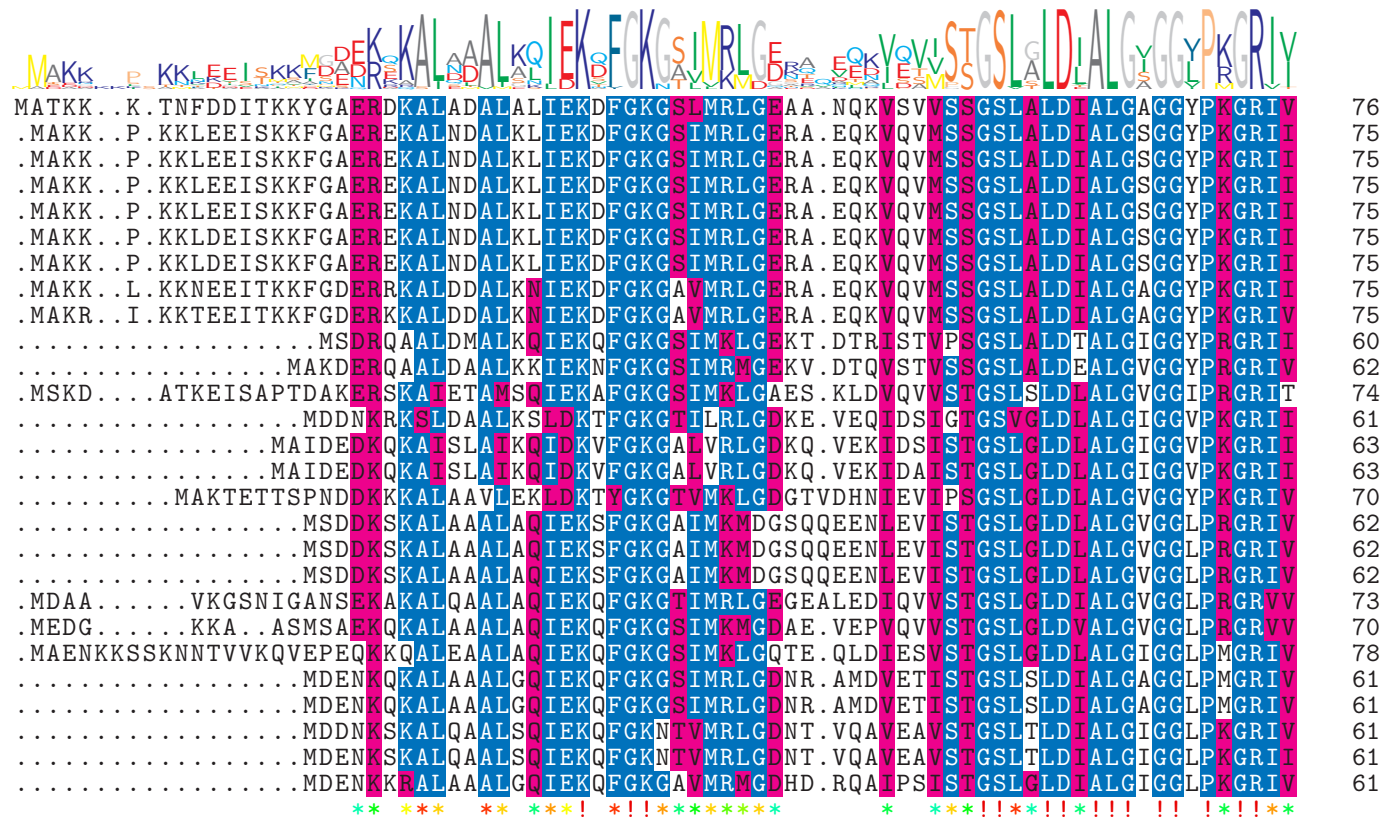


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

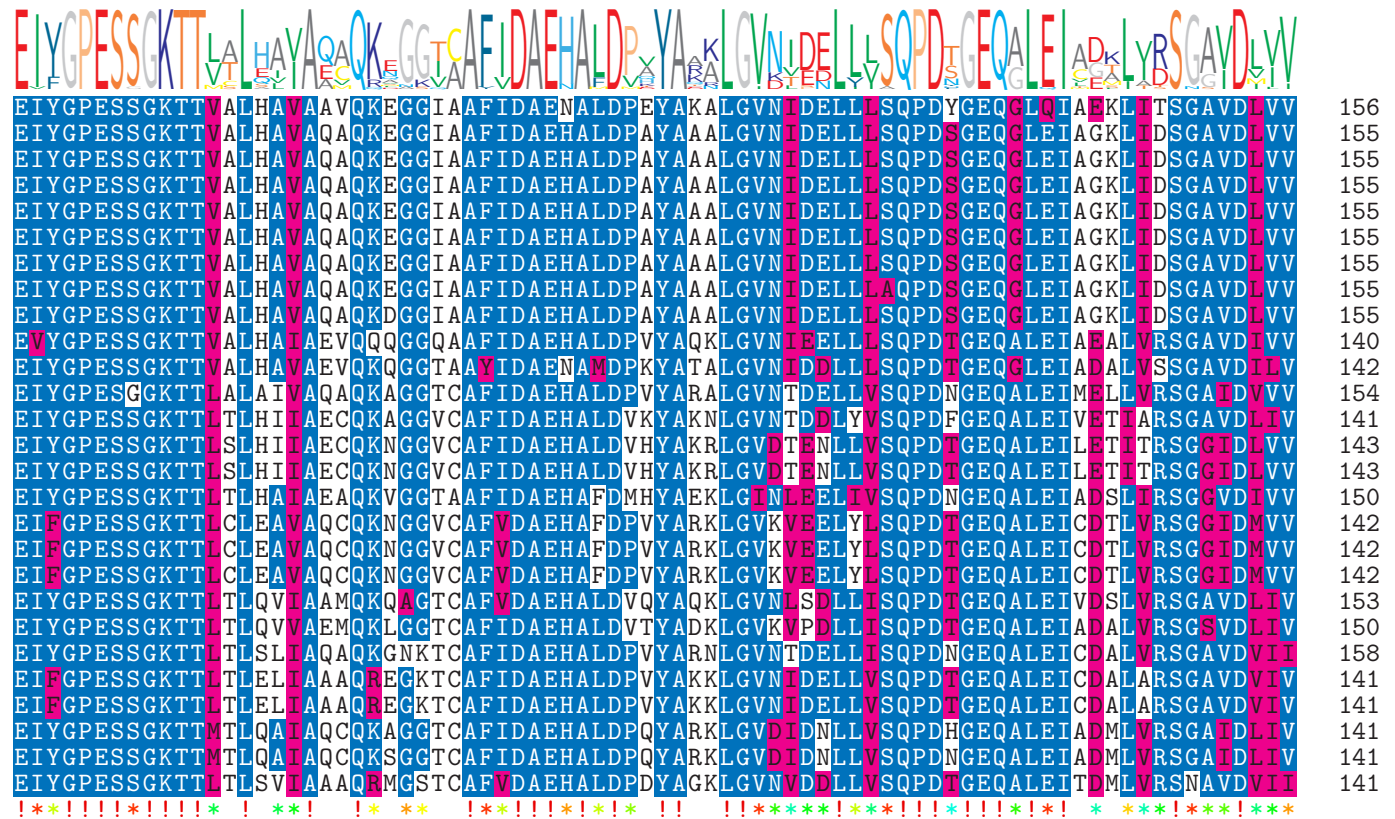
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
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consensus

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logo

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NTDB id 228 SPD RS09265 WP 001085462.1	RLDVRG	NTQIKGTG	.DQKETN	VGKETK	IKVV	KNKVAPPF	KEAVV	EIMYG	GEGISKT	GELLK	IASDL	DIIKK	KAGAW	YSYK	DE	313
NTDB id 194 SPR RS08825 WP 001085462.1	RLDVRG	NTQIKGTG	.DQKETN	VGKETK	IKVV	KNKVAPPF	KEAVV	EIMYG	GEGISKT	GELLK	IASDL	DIIKK	KAGAW	YSYK	DE	313
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NTDB id 453 Spy49 1753c ACI62003.1	RLDVRG	TTQIKGTG	.DQKDSS	IGKETK	IKVV	KNKVAPPF	KVAEV	EIMYG	GEGISRT	GELVK	IASDL	DIIQK	KAGAW	FSYNGE		313
NTDB id 408 SMU RS09530 WP 002262392.1	RLDVRG	NTQIKGTG	.EQKDSN	IGKETK	IKVV	KNKVAPPF	KEAFV	EIIYG	GEGISRT	GELVK	IASDL	GIIQK	KAGAW	SYNGE		313
NTDB id 115 BSU 16940 NP 389576.2	RLEVR	RAEQLKQGN	DVM	GNKTKI	KVV	KNKVAPPF	RTAEV	DIMYG	GEGISKEGE	IIDL	LGTEL	DIVQK	SGSW	SYSEEE		294
NTDB id 625 LCA RS02525 WP 011374196.1	RLEVR	RAETIKNGT	DM	IGNRARI	KVV	KNKVAPPF	KVAEV	DIMY	QGGSIR	TGELV	DMAVEK	DIINK	SGSW	SYSGSE		296
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NTDB id 1245 Cj1673c YP 002345041.1	RLDVR	KKVAT....	L.KQNEEP	IGNRV	KVKV	KNKVAPPF	RQAEF	DVMF	GEGLSR	EGELID	YGVKL	DIVDK	SGAW	FSYKDK		296
NTDB id 1211 C694 RS00795 WP 000963128.1	RIDIR	RIAS....	L.KQNEQH	IGNRA	KA	KVV	KNKVAPPF	REAEF	DIMF	GEGISKEGE	IIDYG	VKLD	DIVDK	SGAWLSYQDK		298
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NTDB id 1007810 ABL838 RS30810 WP 184639288.1	RLDIR	RIGHT....	I.KKGDEA	IGNETK	VKVV	KNKV	SPPFK	TAEF	DILF	GEGISREGE	IIDMG	VNAK	IIDK	SGAWYAYNGE		307
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NTDB id 1205 PSJM300 05545 AFN77183.1	RLDIR	RTGA....	V.KEGDEV	VGSETR	VKVV	KNKVAPPF	RQAEFQ	ILY	GKGIY	ENG	EIID	LG	VQQG	LVEK	SGAWYAYKGS	295
consensus	!***!*		*	**	*****!*****!*	*	*****!*****	*****!*****	*****	***	***	! !*****!				

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consensus

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved