

logo

NTDB	id	135591	mycmcyITA	RS02245	WP	011166605.1	.MSTE....IQKIEDNNLKESQMWNS	KE	KE	AI	KE	IE	KM	FG	KG	SI	VL	GG	QS	.DNLN	VET	FS	SG	SL	LD	NAL	GI	GG	YP	KG	73				
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NTDB	id	194	SPR	RS08825	WP	001085462.1	.MAKK....PKKLEE.ISKKFGA	ERE	K	AL	ND	AL	KL	IE	KD	FG	KG	SI	MR	LG	ER	.AEQKV	QV	M	SS	GS	SL	AL	D	IAL	GS	GG	YP	KG	72
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NTDB	id	453	Spy49	1753c	ACI62003.1		.MAKK....LKKNEE.ITKKFGD	ERR	K	AL	DD	AL	KN	IE	KD	FG	KG	AV	MR	LG	ER	.AEQKV	QV	M	SS	GS	SL	AL	D	IAL	GAG	GY	PK	72	
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NTDB	id	1334	RAOC	RS04870	WP	004918236.1	MAKT...ETTSPND	DK	KA	LA	AV	LE	K	LD	KT	YG	KG	TV	ML	LG	DT	V</													

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NTDB id 135591 mycmcITA RS02245 WP 011166605.1
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

RI	EI	YG	PE	SS	GK	TT	SL	HA	ICEI	QK	GG	TA	CF	DA	EH	AL	DP	YA	AA	LG	VN	DE	LL	SQ	PD	GE	QA	LEI	AD	K	LY	R	SG	AV	D		
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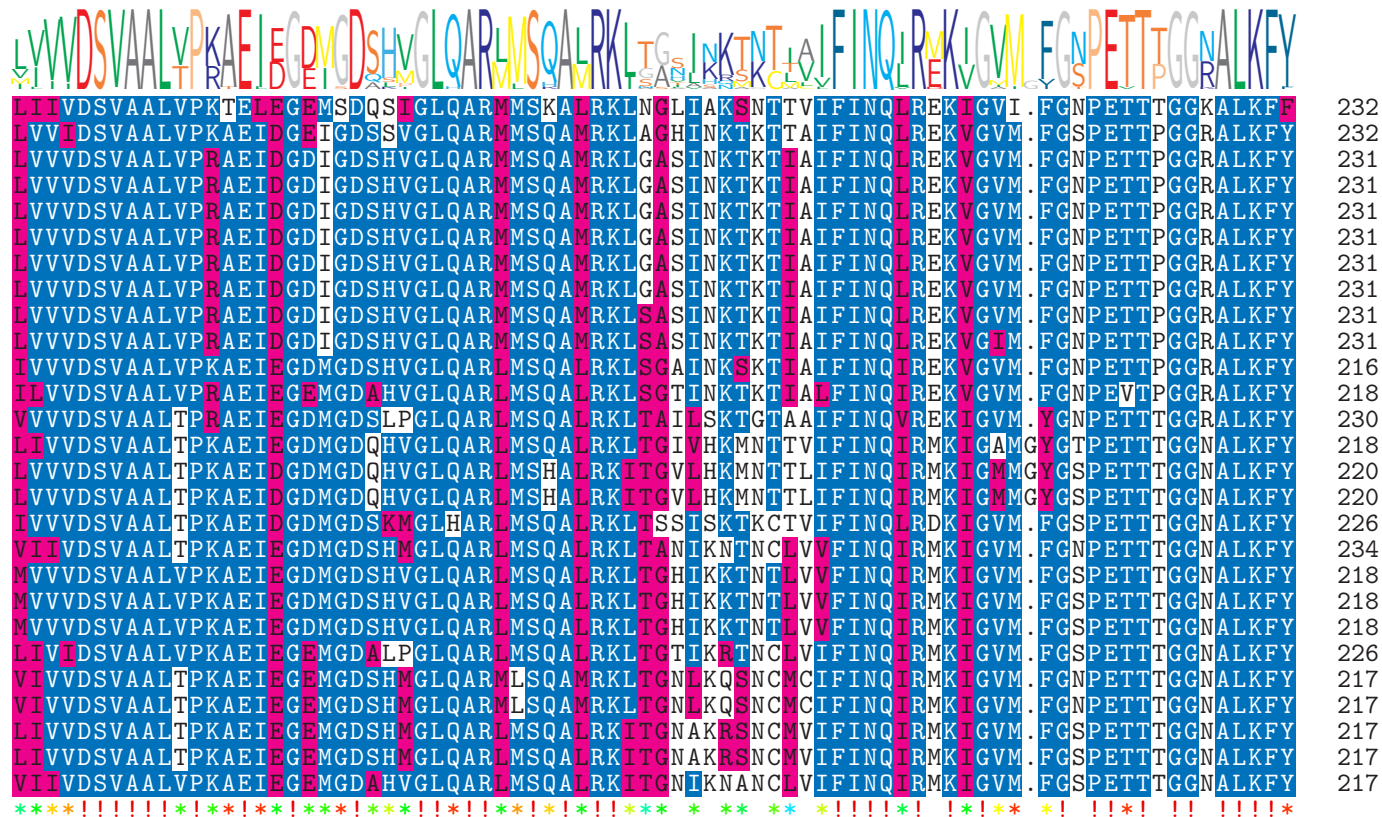
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NTDB id 1205 PSJ3M300 05545 AFN77183.1

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consensus

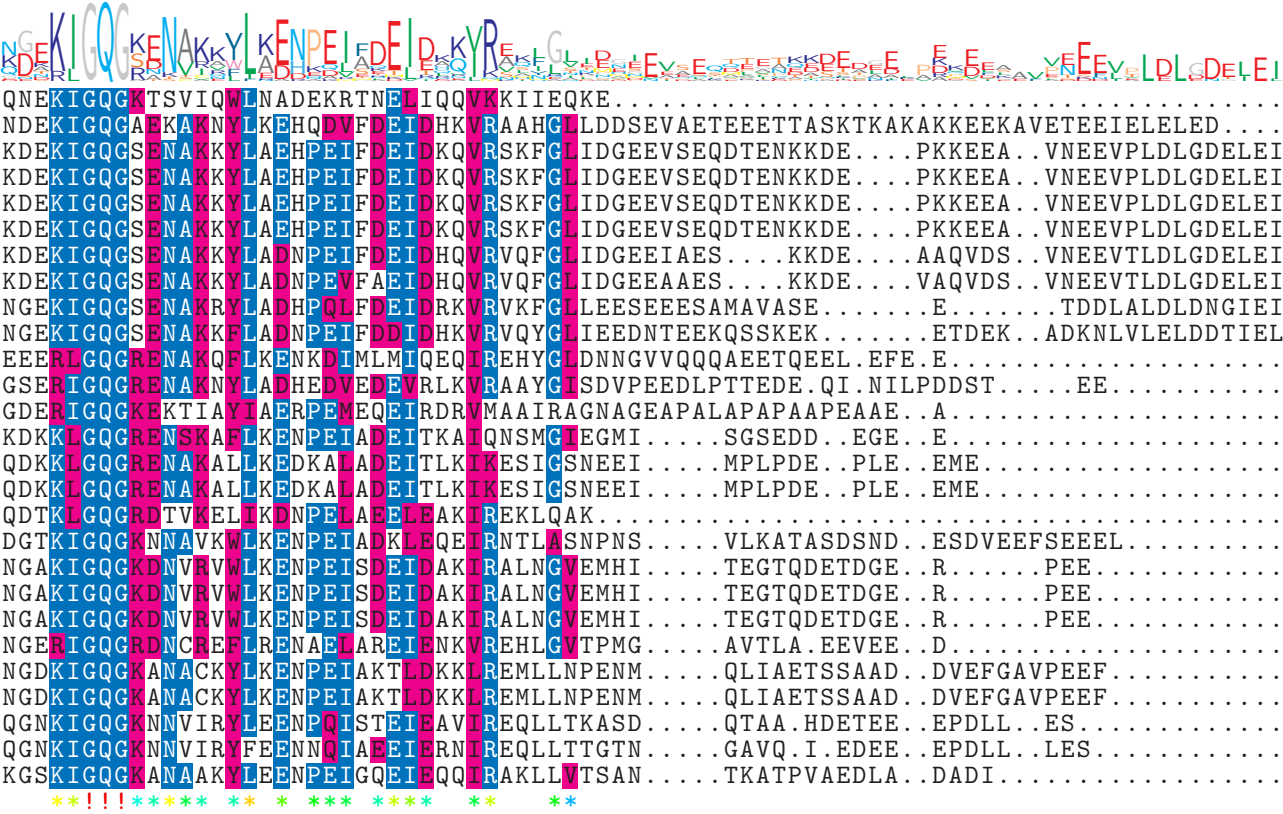


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NTDB id	269	KZH43	RS08665	WP	001085462.1	ASVRLDVRGNTQIKGTG.DQKETN	VGKETKIKVVKNKVAPPFKAEAVV	EIMYGEGISKTGELKLIASDLDI	IKKAGAWYSY	310
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consensus



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NTDB id 135591 mycm ycITA RS02245 WP 011166605.1	E.I.EE	345
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved