

logo

NTDB	id	600	KW2	RS01765	WP	011675413.1	MATKK....KTNFDDITKKYGA	ERD	KALAD	ALAL	IEKD	FGKGS	LMRLGEA	.ANQKVSV	VSSGSL	ALD	I	ALGAGGY	PKGR	74														
NTDB	id	269	KZH43	RS08665	WP	001085462.1	.MAKK....PKKLEEISKKFGA	ERE	KALND	ALKL	IEKD	FGKGS	SIMRLGER	.AEQKVQVM	SSSGSL	ALD	I	ALGSGGY	PKGR	73														
NTDB	id	228	SPD	RS09265	WP	001085462.1	.MAKK....PKKLEEISKKFGA	ERE	KALND	ALKL	IEKD	FGKGS	SIMRLGER	.AEQKVQVM	SSSGSL	ALD	I	ALGSGGY	PKGR	73														
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NTDB	id	159	SP	RS09750	WP	001085462.1	.MAKK....PKKLEEISKKFGA	ERE	KALND	ALKL	IEKD	FGKGS	SIMRLGER	.AEQKVQVM	SSSGSL	ALD	I	ALGSGGY	PKGR	73														
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NTDB	id	625	LCA	RS02525	WP	001374196.1	MAK	DERQAALDA	ALKK	IEKN	FGKGS	SIMRMGEK	.VDTQVST	VSSGSL	ALDE	AL	LGVG	GGYP	RGR	60														
NTDB	id	115	BSU	16940	NP	389576.2	MS	DRQAALDM	ALKQ	IEKQ	FGKGS	SIMKLGEK	.TDTRIST	VP	SGSL	ALD	TALGI	GGY	P	RGR	58													
NTDB	id	1313	DR	RS12030	WP	010888966.1	MSK	DATKEISAP	TD	AKERS	KA	IET	AMS	QIEKA	AFGKS	SIMKL	GA	E	SKLD	VQV	ST	GS	SLD	LALGV	GGIP	RGR	72							
NTDB	id	1245	Cj1673c	YP	002345041.1	MD	DNKRKSLDA	AALKS	L	DKT	FGKGT	ILRLG	DK	.EVEQ	IDS	IGT	GS	VGLD	LALGI	GGV	PKGR	59												
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NTDB	id	1334	RAOC	RS04870	WP	004918236.1	MAKT	.ETTSPN	DDKK	KALAA	VLEK	L	DKT	YGKG	TVM	KL	G	D	GTVD	HN	IEV	I	P	S	G	S	L	G	L	D	LALGV	GGY	PKGR	68
NTDB	id	1389	A4U84	RS07560	WP	021112712.1	.MAENK	KSSKN	NTV	.VKQVEP	EQQKQ	ALEA	ALA	QIEKQ	FGKGS	SIMKL	G	QT	.EQLD	IES	V	T	G	S	L	G	L	D	LALGI	GG	L	P	MGR	76
NTDB	id	1130	NGFG	RS03960	WP	003688695.1	MS	DDKS	KALAA	ALA	QIEKS	FGKGA	IMKMD	G	S	Q	Q	EEN	LEV	I	S	T	G	S	L	G	L	D	LALGV	GG	L	P	RGR	60
NTDB	id	1124	NGFG	RS03960	WP	003688695.1	MS	DDKS	KALAA	ALA	QIEKS	FGKGA	IMKMD	G	S	Q	Q	EEN	LEV	I	S	T	G	S	L	G	L	D	LALGV	GG	L	P	RGR	60
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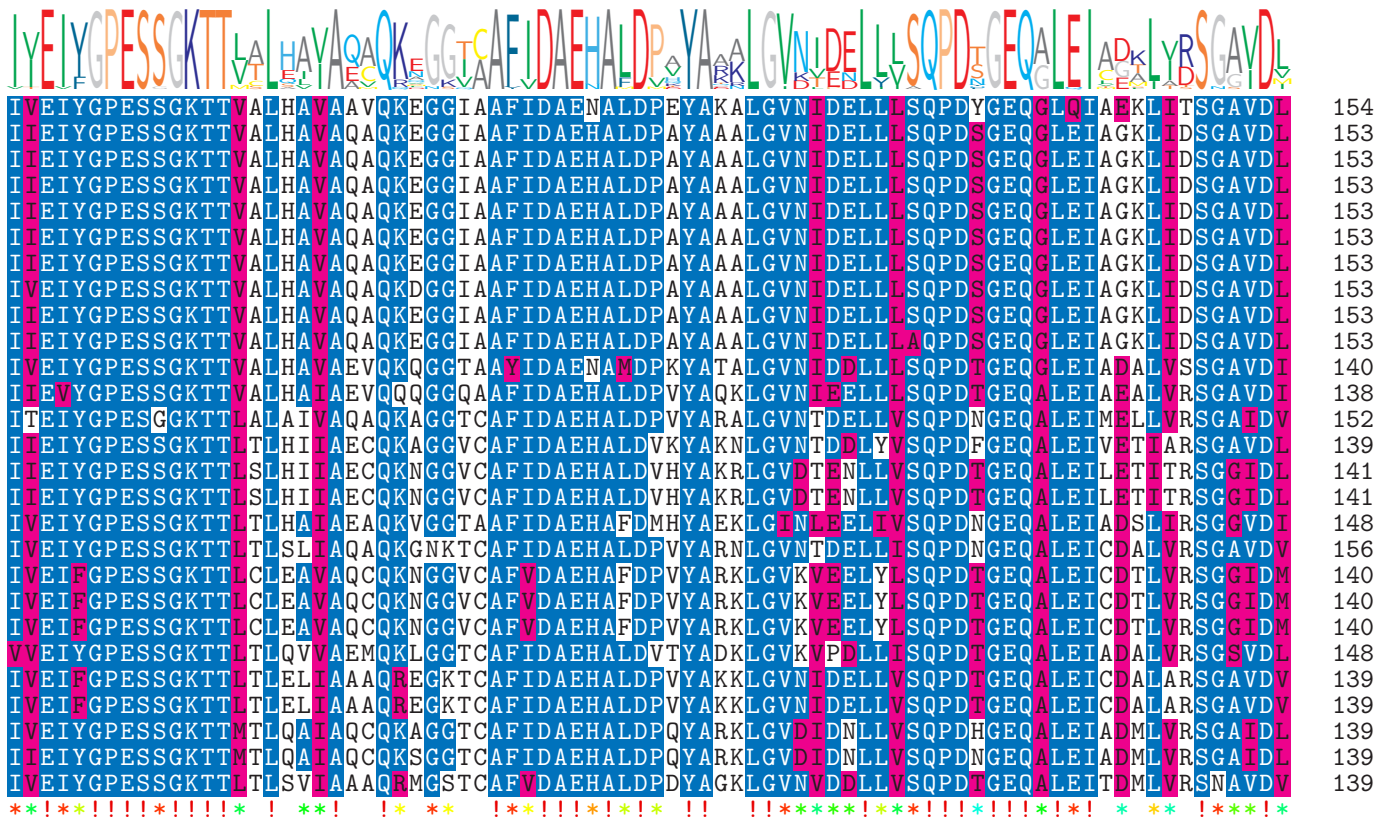
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NTDB id 527 SMSK321 RS11065 WP 001085454.1
NTDB id 408 SMU RS09530 WP 002262392.1
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consensus



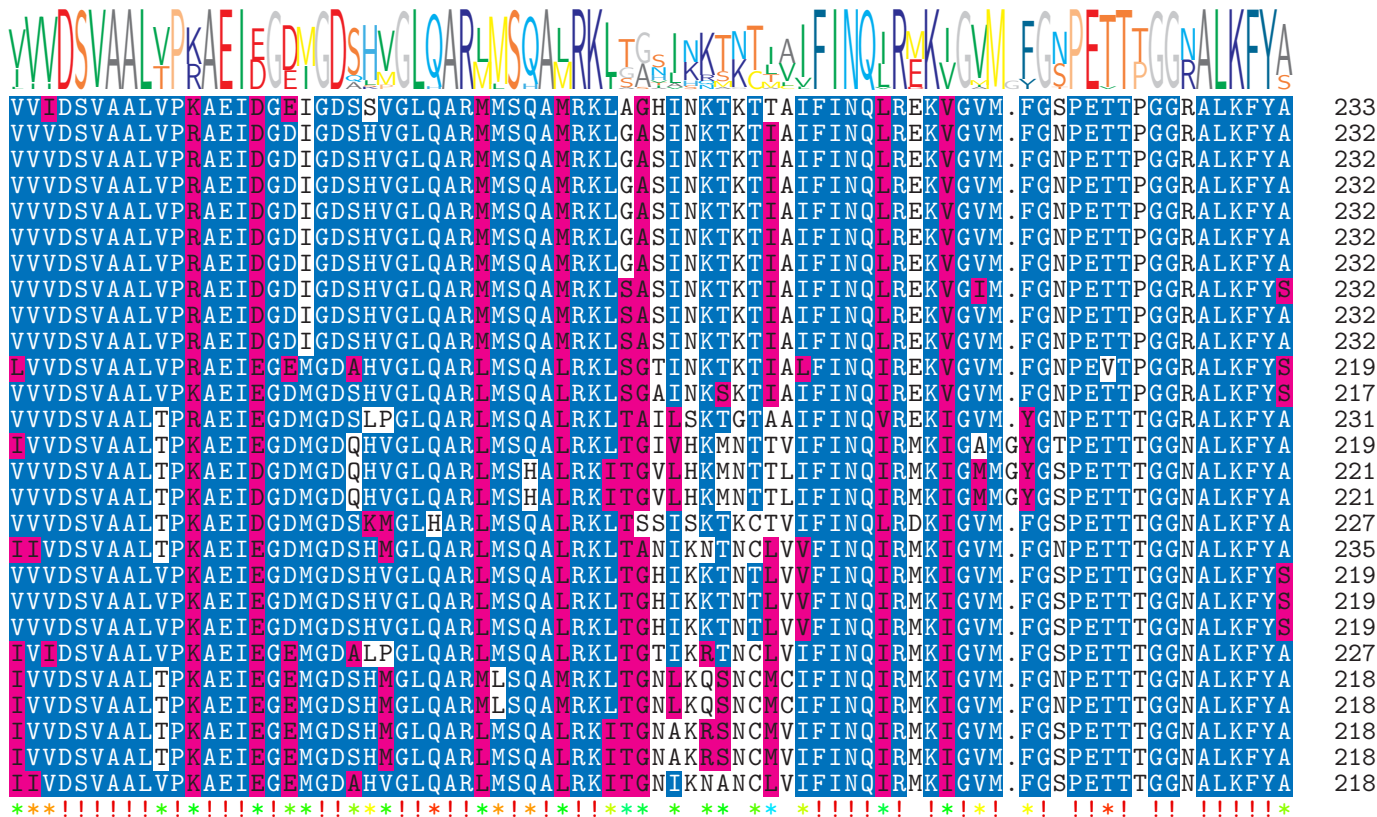
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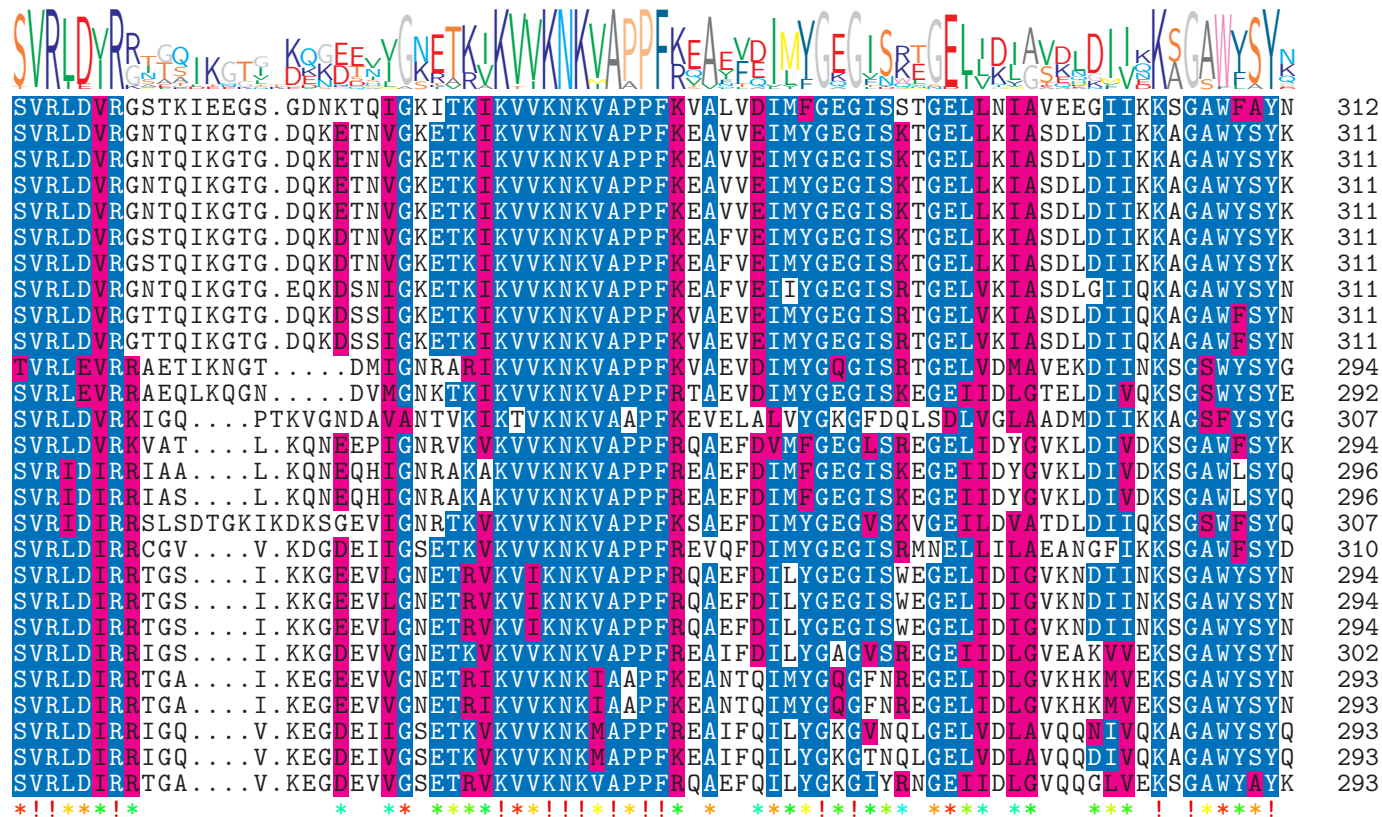
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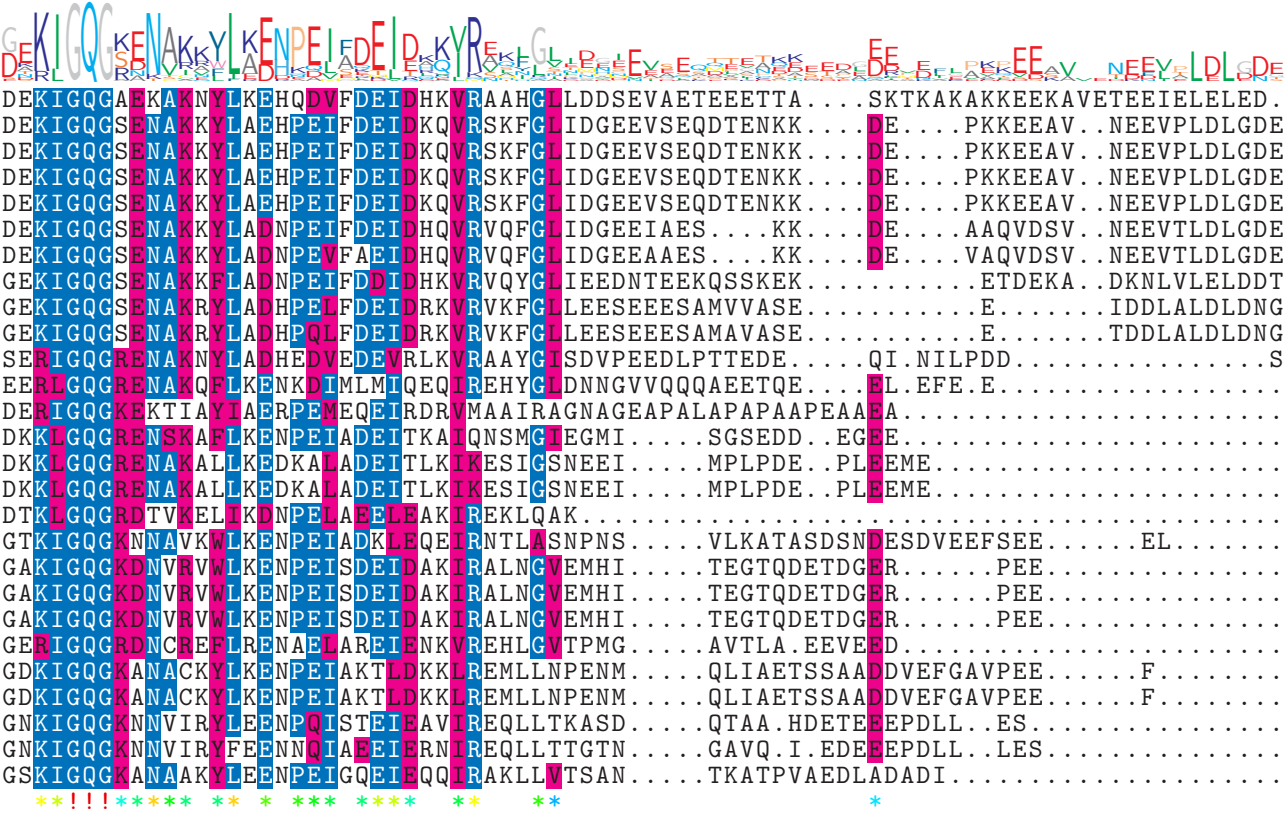
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consensus

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consensus



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consensus			

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