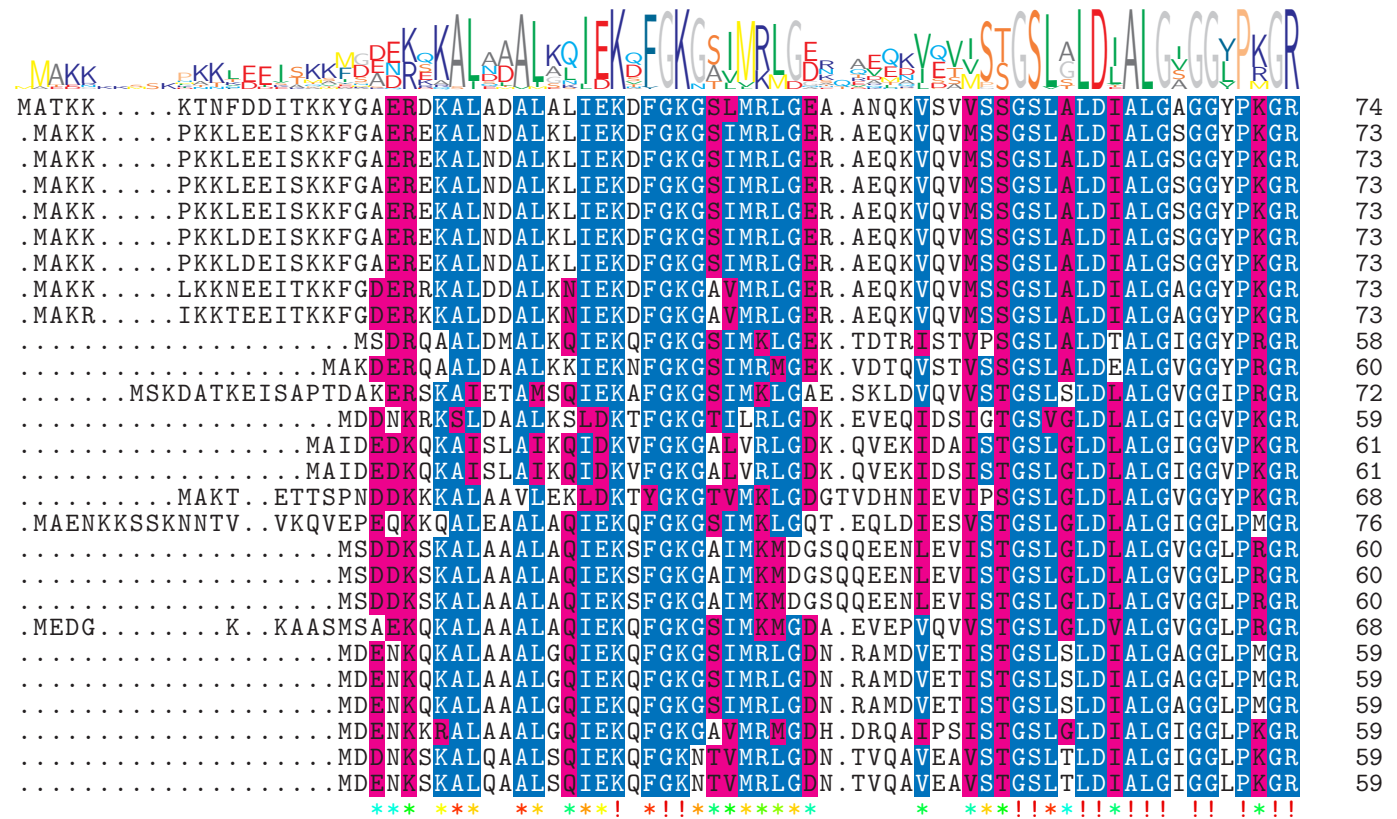


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consensus

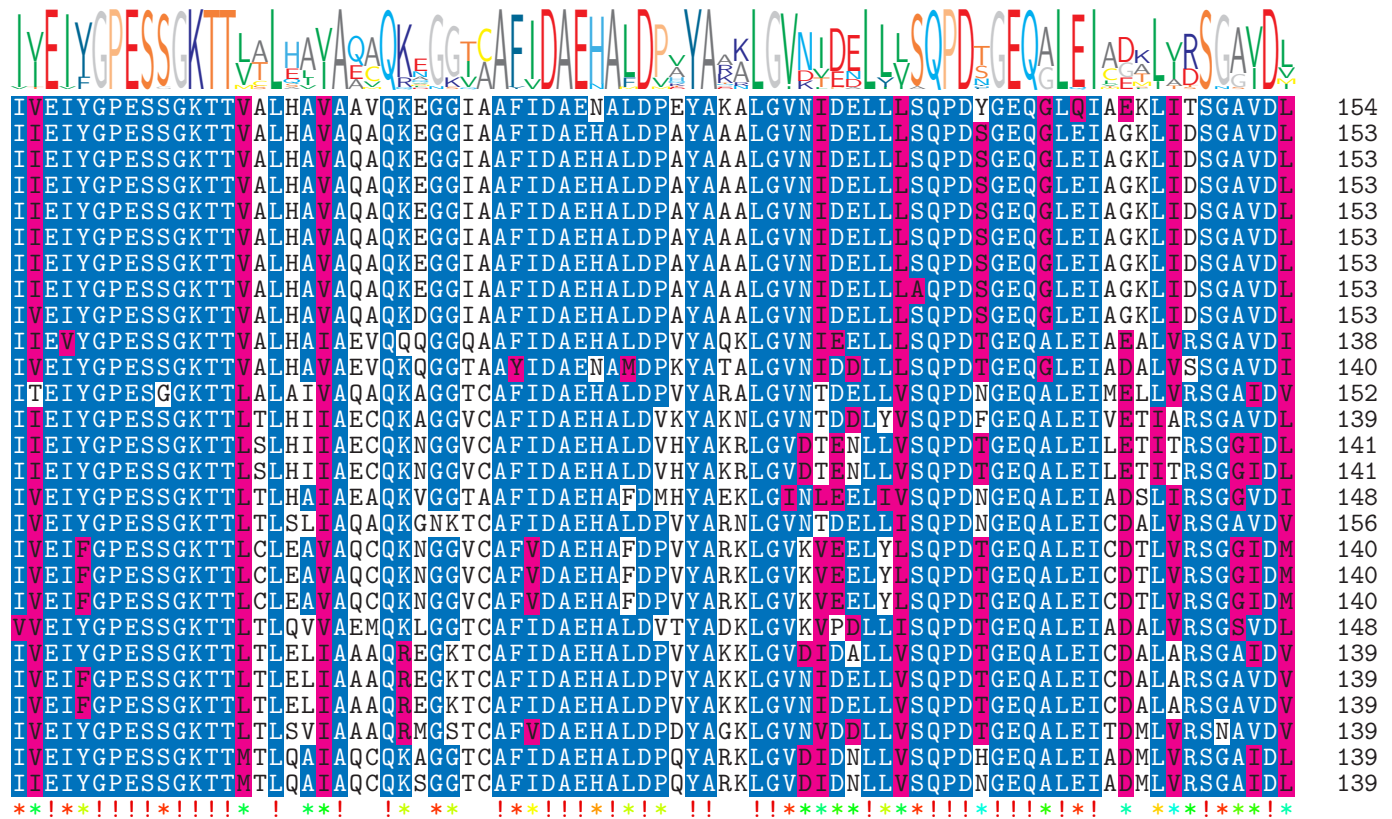
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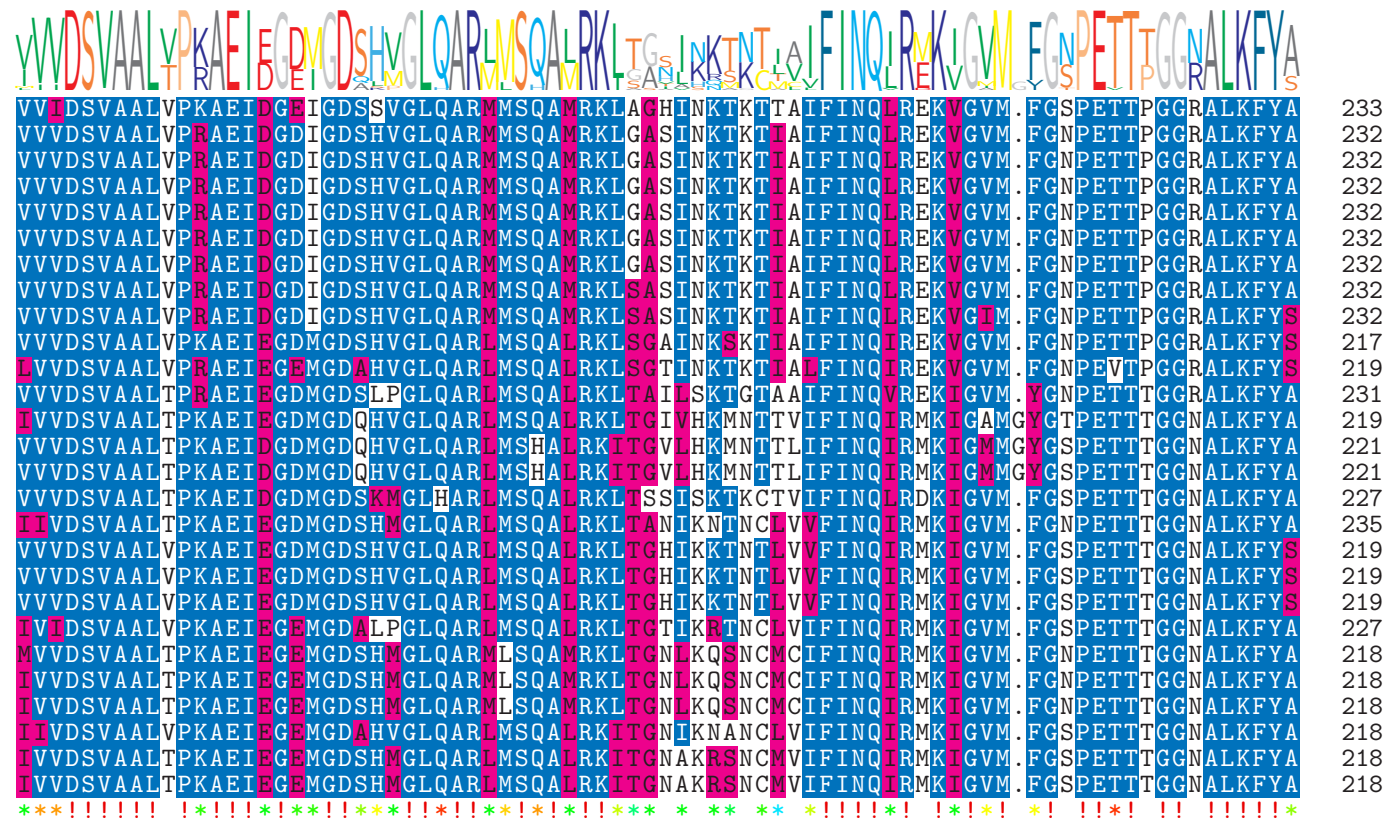
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

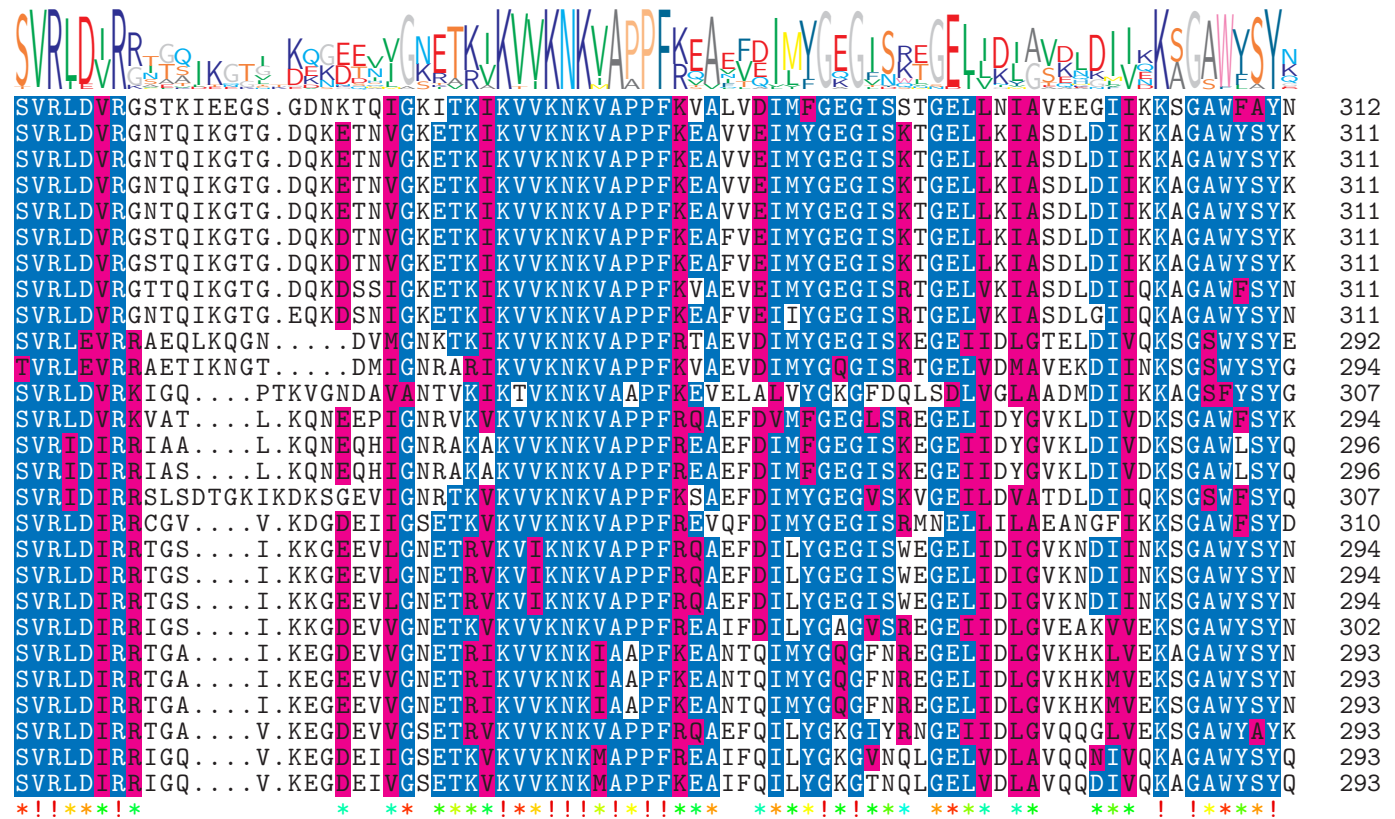
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NTDB id 1022 ACIAD RS06390 WP 0004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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NTDB	id	453	Spy49	1753c	ACI62003.1		GEKIGQGS	ENAKRY	LADHPQ	LFDE	ID	DRKV	RVKF	G	L	LEESEEE	SAMAVASE		E.....	TDDL	LALDL	NGIEIE		377
NTDB	id	408	SMU	RS09530	WP	002262392.1	GEKIGQGS	ENAKKFL	ADNPEI	FFDD	ID	DHKV	RVQY	G	L	IEEDN	TEEKQ	SSKEK		ETDEK	..ADKN	LVLEL	DDTIELE	382
NTDB	id	115	BSU	16940	NP	389576.2	EERL	GGCRENA	KQFL	ENKDI	IMLM	IQE	Q	Q	Q	Q	Q	Q	Q	Q	Q	Q	Q	348
NTDB	id	625	LCA	RS02525	WP	011374196.1	SERIGQGS	ENAKNY	LADHED	VE	EV	RLKV	RAAYG	I	S	DVPEED	LPTTEDE	QI	NILP	DDST		EE		355
NTDB	id	1313	DR	RS12030	WP	010888966.1	DERIGQCK	EKTIA	YIAER	PEME	Q	EIR	DRVMAA	AIRAGN	AGEAP	ALAPAPA	AAPEAAEA							363
NTDB	id	1245	Cj1673c	YP	002345041.1		DKKL	GGCRENS	KAF	LKENPE	IADE	ITKAI	QNSMG	I	E	G	MIS....	GSED....	DEGEE					343
NTDB	id	1214	NCTC11637	00359	SQJ03206.1		DKKL	GGCRENA	KALL	KEDK	ALADE	ITLKIK	ESIGS	N	EEIM....	PLPD....	EPLEE	..ME					347	
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NTDB	id	1123	OK783	RS03845	WP	003688695.1	GAKIGQCK	DNV	RVVLKENPE	ISDE	IDA	KIRAL	NCVEM	HIT....	E.....	GTQDET	DGE	..RPEE					348	
NTDB	id	1427	RS	RS02760	WP	011000510.1	GERIGQCR	DNCREF	LRENAEL	AREI	ENKVR	REHL	G	VTMGA		VTL	AAEEVE	D.....					352	
NTDB	id	404152	GNY17	RS14725	WP	00547855																		

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consensus			

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