

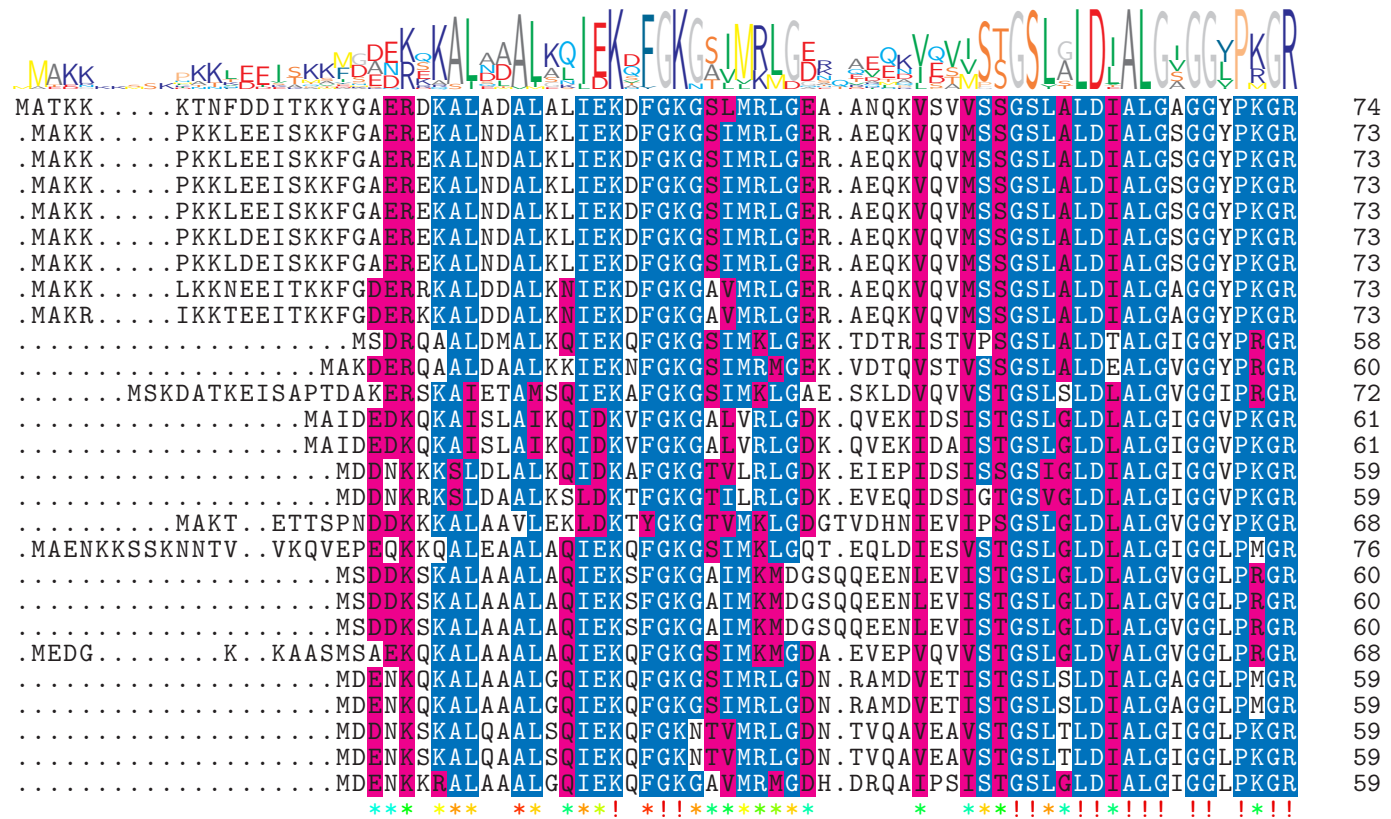
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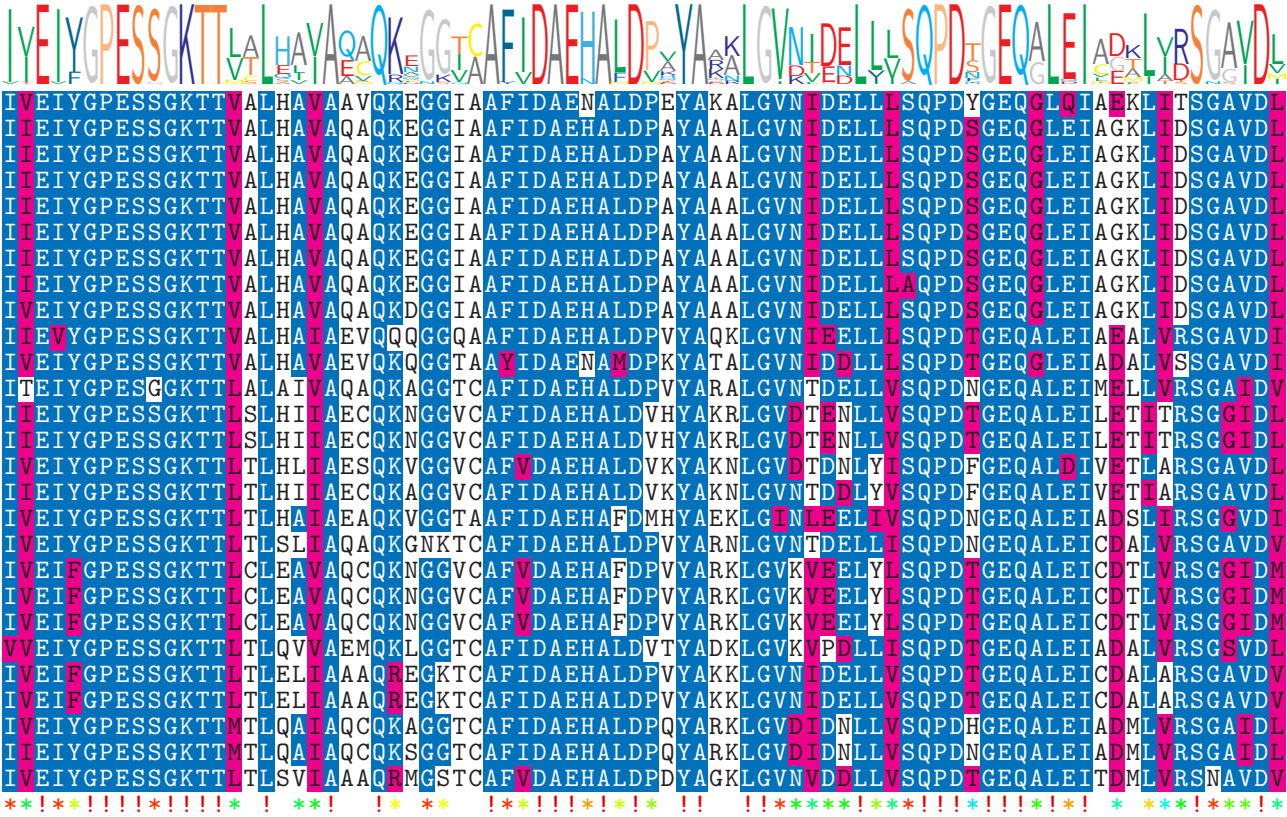
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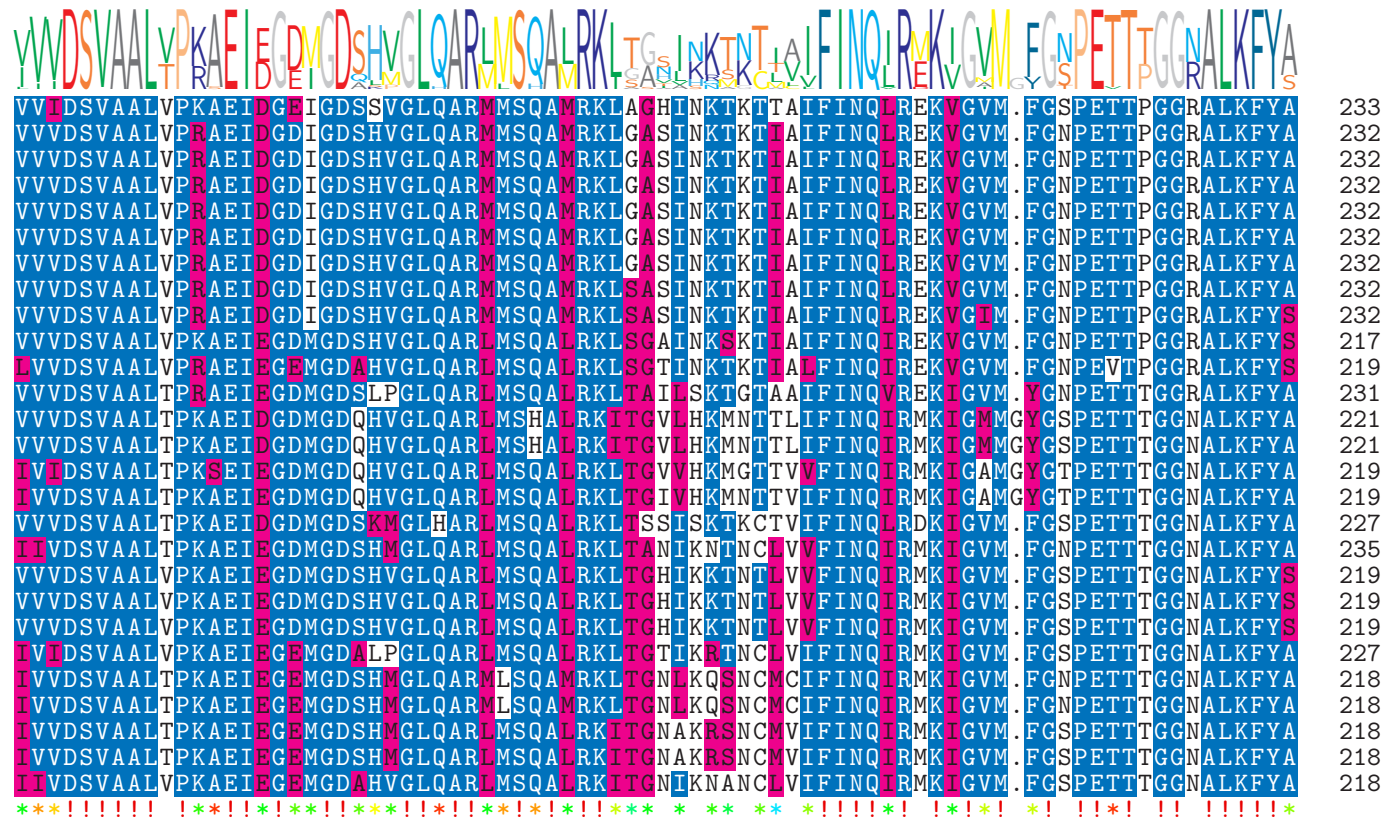
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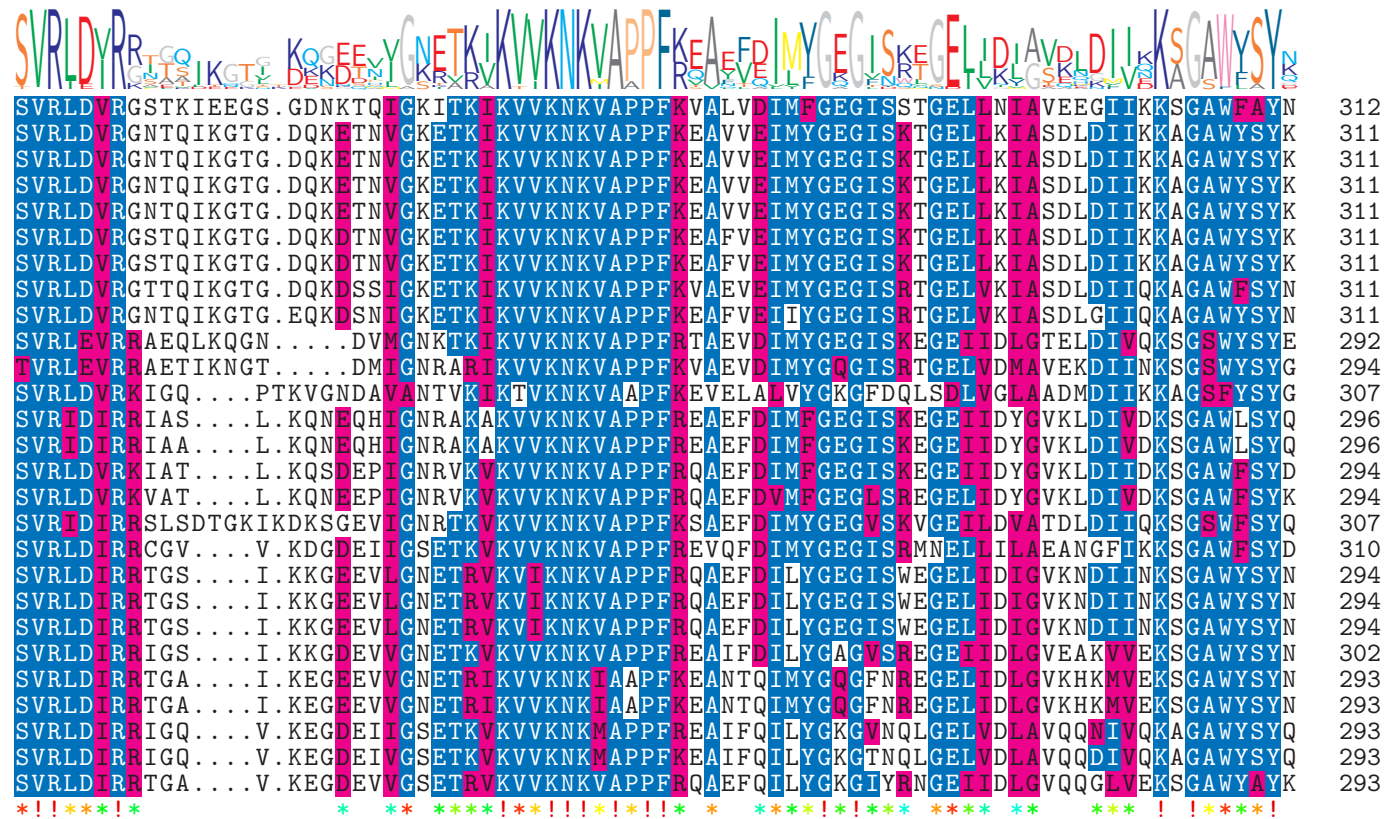


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NTDB id 1205 PSJM300 05545 AFN77183.1
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NTDB id 527 SMSK321 RS11065 WP 001085454.1	DEKIGQGS	ENAKKYLA	DNPEV	FAEID	DHQVRV	VQFGL	IDGEEA	AES....	KKDE....	VAQVDS	..VNEEV	TLDLG	DELEIE	381	
NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGS	ENAKRYLA	DHP	QLFDEID	DRKVRV	VKFG	L	LEESEES	AMAVASE		E		TDDLALD	LDNGIEIE	377
NTDB id 408 SMU RS09530 WP 002262392.1	GEKIGQGS	ENAKKFLA	DNPEIF	DDID	DHKVRV	VQYGL	IEEDNTE	EKQSSKEK		ETDEK	..ADKNL	VLELDD	TIELE	382	
NTDB id 115 BSU 16940 NP 389576.2	EERLGQGR	ENAKQFL	KENKD	IMLM	IQEQI	REHYGL	DNNNGV	VQQQAE	ETQEEL	EFE	E			348	
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	DKKLGQGR	ENAKALL	KEDK	ALADEI	ITLKIK	ESIGS	NEEI		MPLP	DEPLEEME				347	
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consensus			

- non conserved
- similar
- ≥ 50% conserved