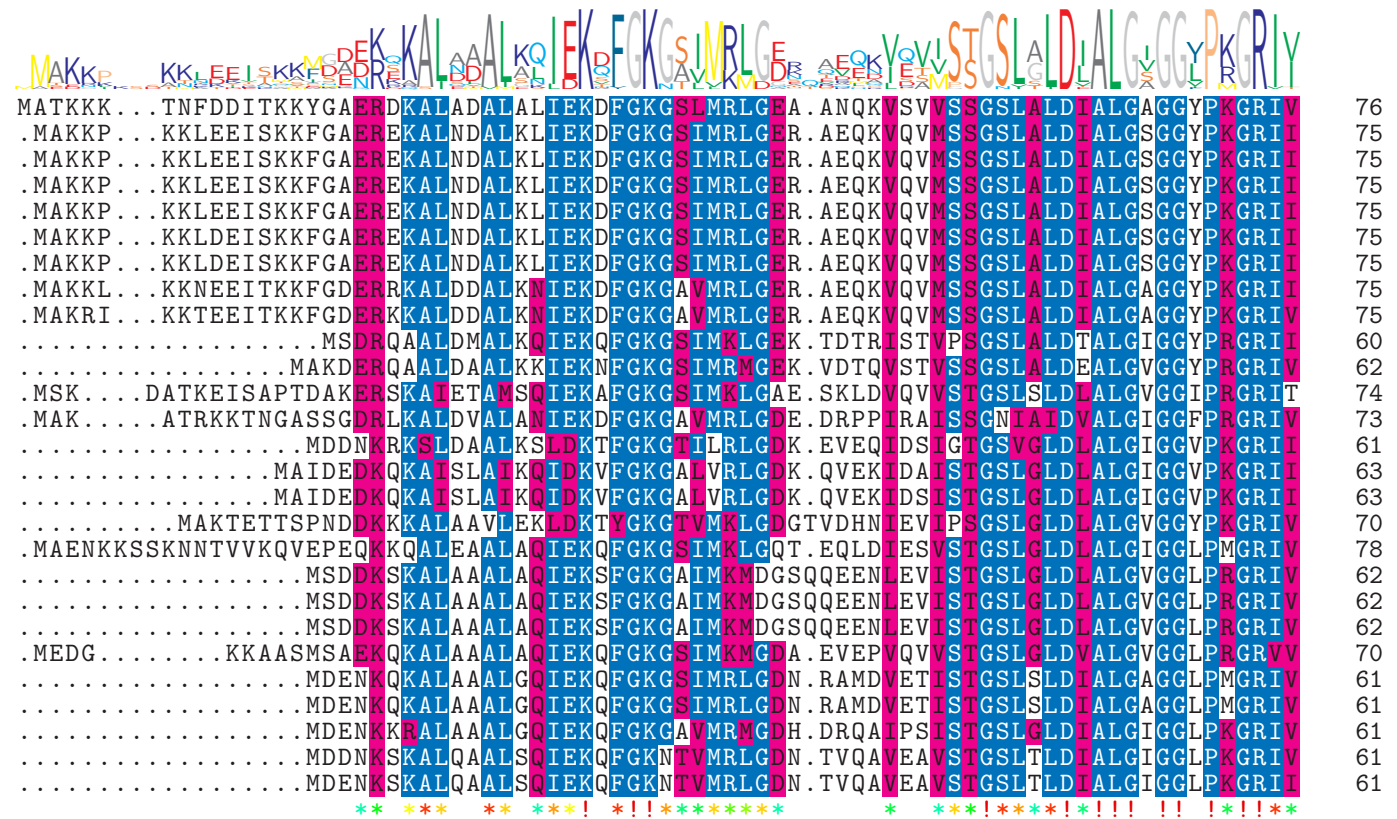


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consensus

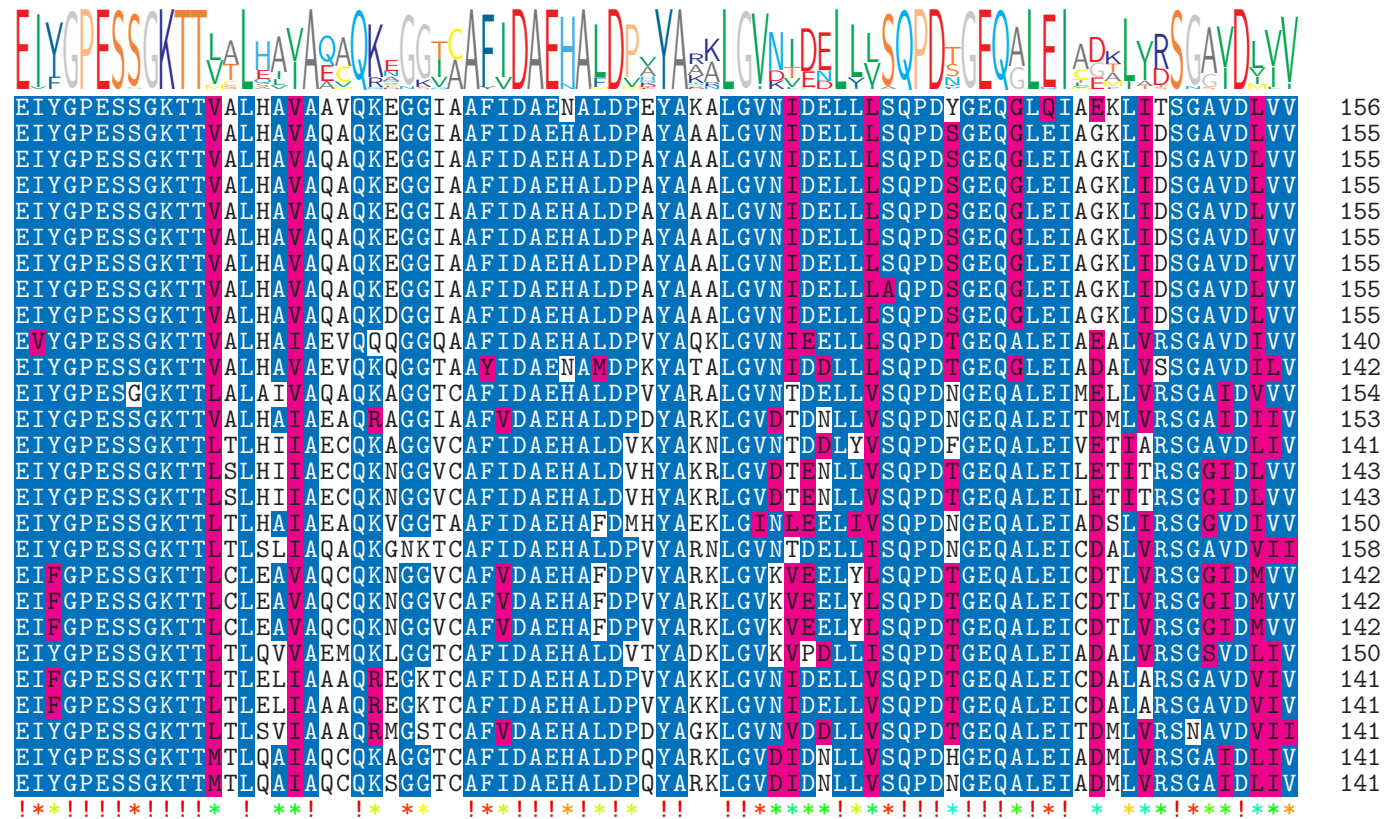
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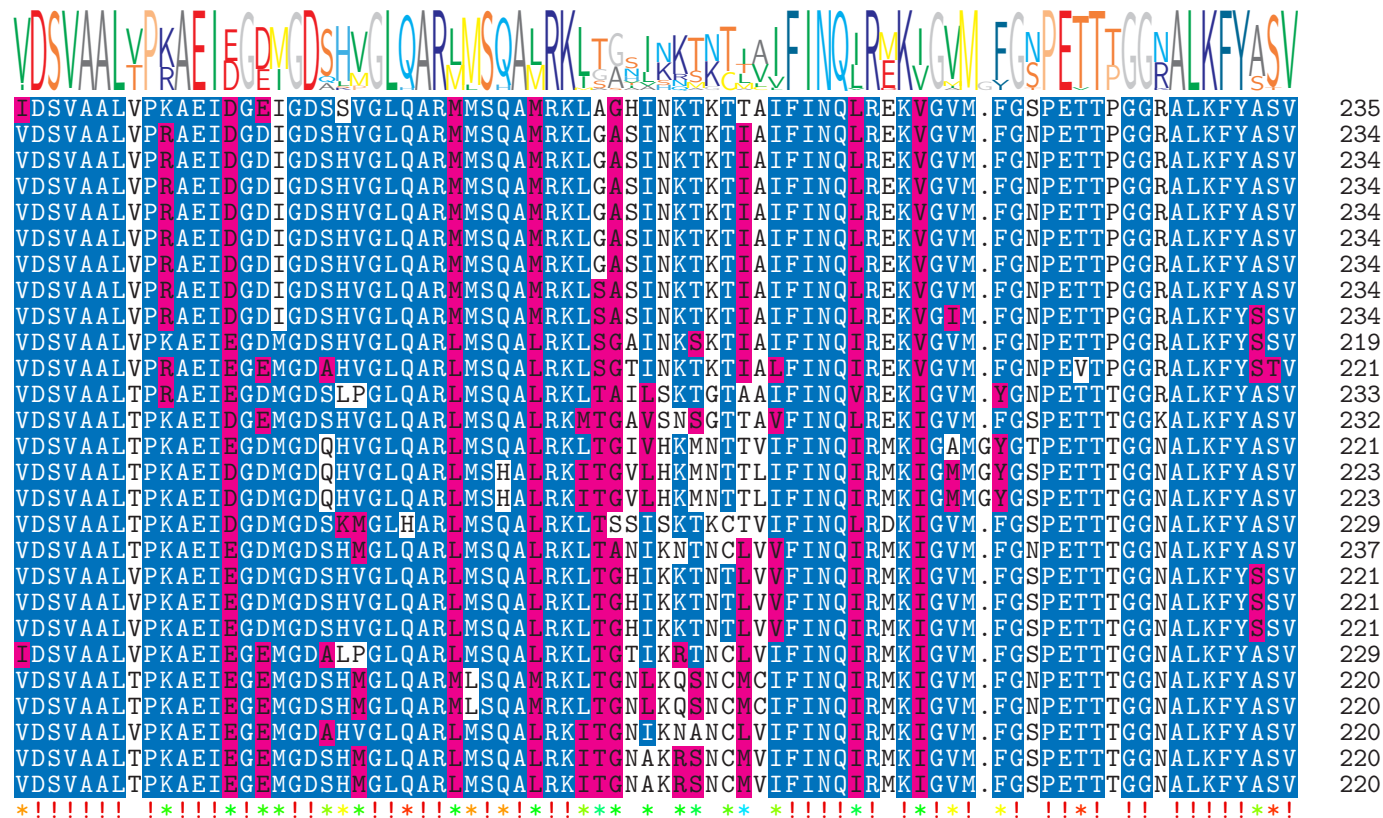
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consensus

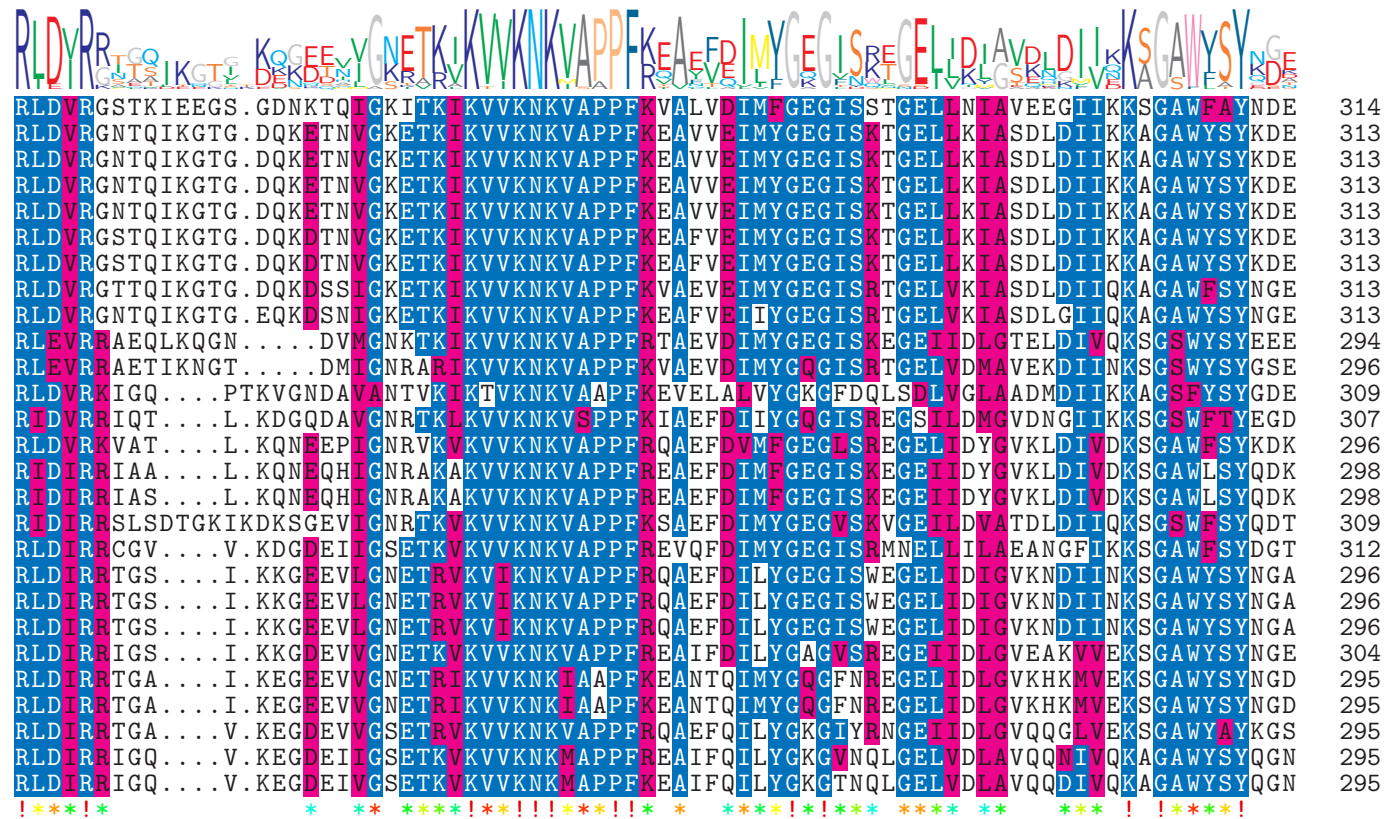
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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logo

NTDB id 600 KW2 RS01765 WP 011675413.1	KIGQGA	EKAKNYLKEHQDV	DEIDHKVRAAHGL	LDDSEVAETEEETT	ASKTKAKAKKEEKAVET	EEIELELED		387
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NTDB id 453 Spy49 1753c ACI62003.1	KIGQGS	ENAKRYLADHPQLF	DEIDRKVRVKFGL	LEESEESAMAVASE	E.....TDD	LALDLNGLIEIED		378
NTDB id 408 SMU RS09530 WP 002262392.1	KIGQGS	ENAKKFLADNPEIF	DDIDHKVRVQYGL	IEEDNTEEKQSSKEK	ETDEK..ADKN	LVLELDDTIELED		383
NTDB id 115 BSU 16940 NP 389576.2	RLGQG	RENAKQFLKENKDIM	LMIQEQIREHYGL	DNNGVVQQAEETQEEL	EFEE.....			348
NTDB id 625 LCA RS02525 WP 011374196.1	RIGQG	RENAKNYLADHEDVE	DEVRLKVRAAYGIS	DVPEEDLPTTEDE	.QI.NILPDDST	EE		355
NTDB id 1313 DR RS12030 WP 010888966.1	RIGQG	KEKTIAYIAERP	EMEQEIRDRVMAA	IRAGNAGEAPALAP	APAAPE.....AA	EA		363
NTDB id 318633 D4R08 RS03570 WP 046650758.1	QLGQG	KEKSRDYLKANPDL	AAELEDKIKRKL	GVGEYANAT....DEEPD	AD.....A	PVDVVPDIDFDDE		368
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	KIGQG	KNNAVKWLKENPEI	ADKLEQEI	RNTLASNPNSVLK....ATASD	SN.....DESD	VEEFSEEL...		372
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NTDB id 1205 PSJM300 05545 AFN77183.1	KIGQG	KANAACKYLEENPEI	GIEEQIRAKLLV	TSANTKA....TPVAE	DL.....ADADI			347
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	KIGQG	KNNVIRYFEENNQI	AEEIERNIREQLLT	TGTNGAV.....Q.I.E	DE.....EED	PDLL..LES.....		349
consensus	**!!!	*****	**	*****	*****	**	**	

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved