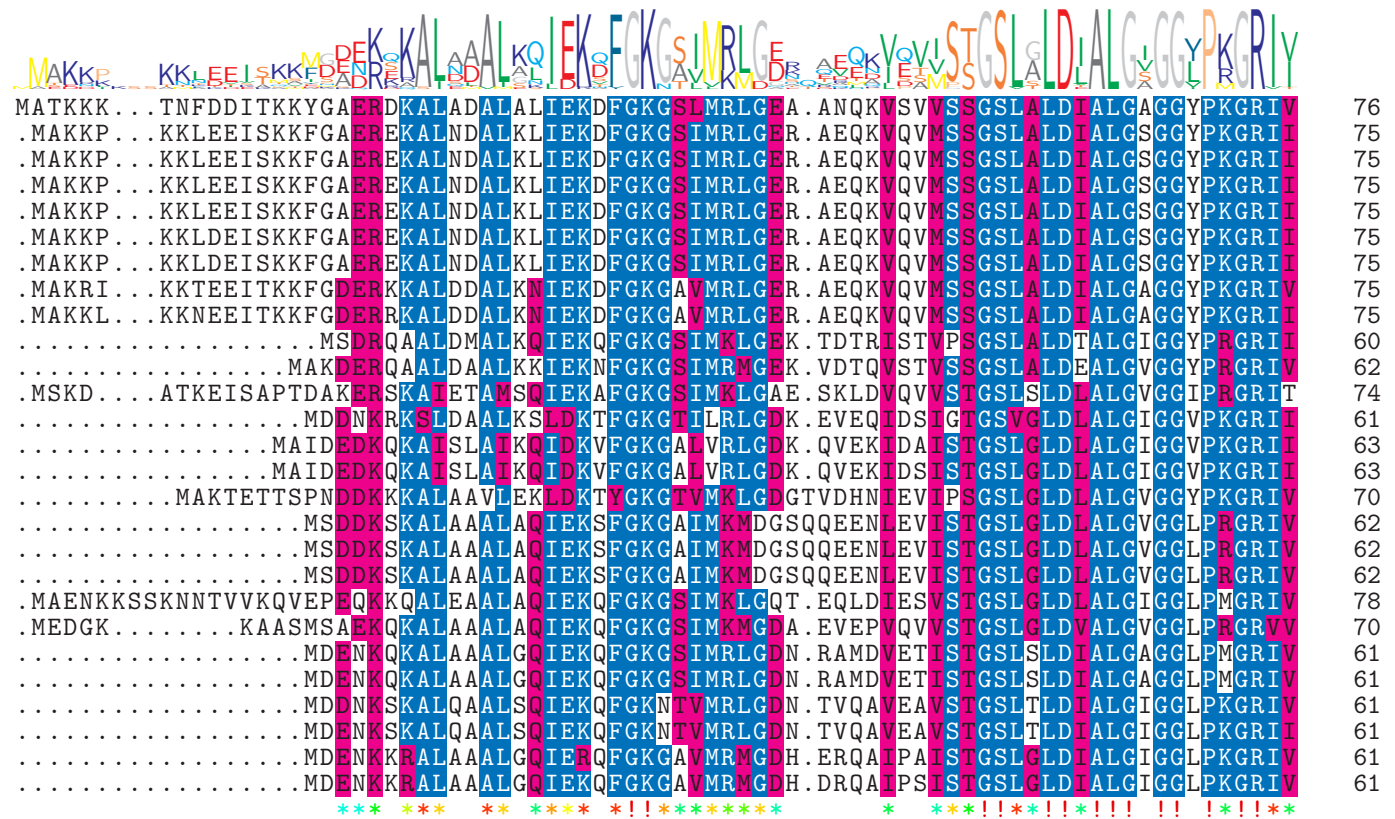


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consensus

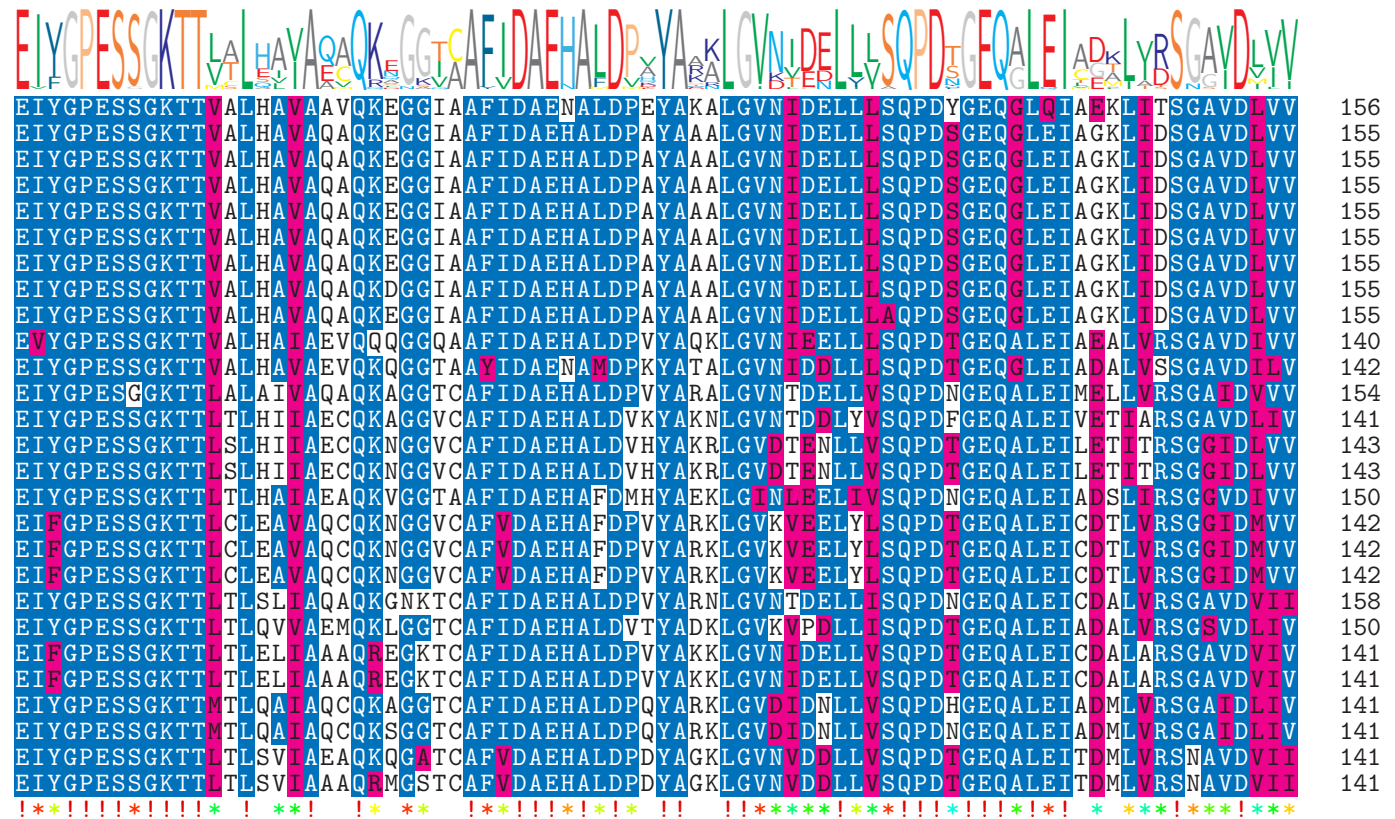
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consensus

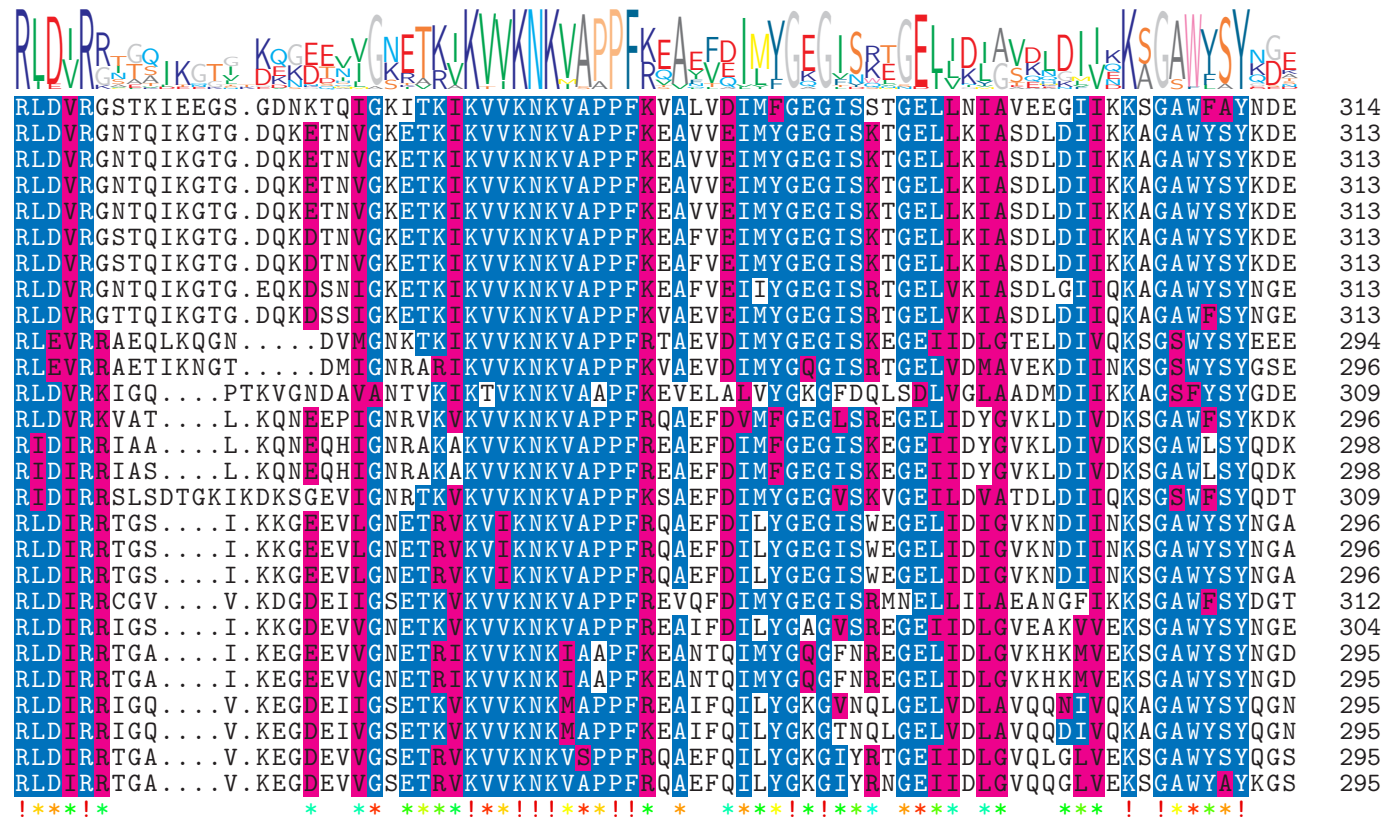
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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NTDB id 600 KW2 RS01765 WP 011675413.1	KIGQGA	EKAKNYLKEHQDV	DEIDHKVRAAHG	LDDSEVAETEEETT	...	ASKTKAKAKKEEKAVETEEIELELED	...	387					
NTDB id 269 KZH43 RS08665 WP 001085462.1	KIGQGS	ENAKKYLA	EHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENK	...	KDE	...	PKKEEA	..VNEEVPLDLGDELE	383		
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NTDB id 453 Spy49 1753c ACI62003.1	KIGQGS	ENAKRYLA	DHPQLF	DEIDRKVRVKFGL	LEESEEEESAMAVAS	...	E	...	E	...	TDDLALDLDNGIE	375	
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	KLGGGR	ENAKALLKEDKALADEITLK	IKESIGSNEEIM	...	PLPDEP	LEEME	...					347	
NTDB id 1211 C694 RS00795 WP 000963128.1	KLGGGR	ENAKALLKEDKALADEITLK	IKESIGSNEEIM	...	PLPDEP	LEEME	...					347	
NTDB id 1334 RAOC RS04870 WP 004918236.1	KLGGGR	DTVKELIKDNPELAELEAKI	REKLQAK	...								343	
NTDB id 1130 NGFG RS03960 WP 003688695.1	KIGQGD	NVRVWLKENPEISDEIDAK	IRALNGVEMHIT	...	EGTQDET	DGE	...	RP	EE	...		348	
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NTDB id 1123 OK783 RS03845 WP 003688695.1	KIGQGD	NVRVWLKENPEISDEIDAK	IRALNGVEMHIT	...	EGTQDET	DGE	...	RP	EE	...		348	
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NTDB id 290485 CSC26 RS29990 WP 003092260.1	KIGQGD	KANAAYLEDNPEIGSVLEKT	IRDQLLAKSGPV	...	KADAE	EVADAE	AD	...				346	
NTDB id 1205 PSJM300 05545 AFN77183.1	KIGQGD	KANAAYLEENPEIGQIE	EQQIRAKLLVTSANT	...	KATPVAED	LADADI	...					347	
consensus	***!!!	*****	**	*****	*****	**	**						

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NTDB id 600 KW2 RS01765 WP 011675413.1		387
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved