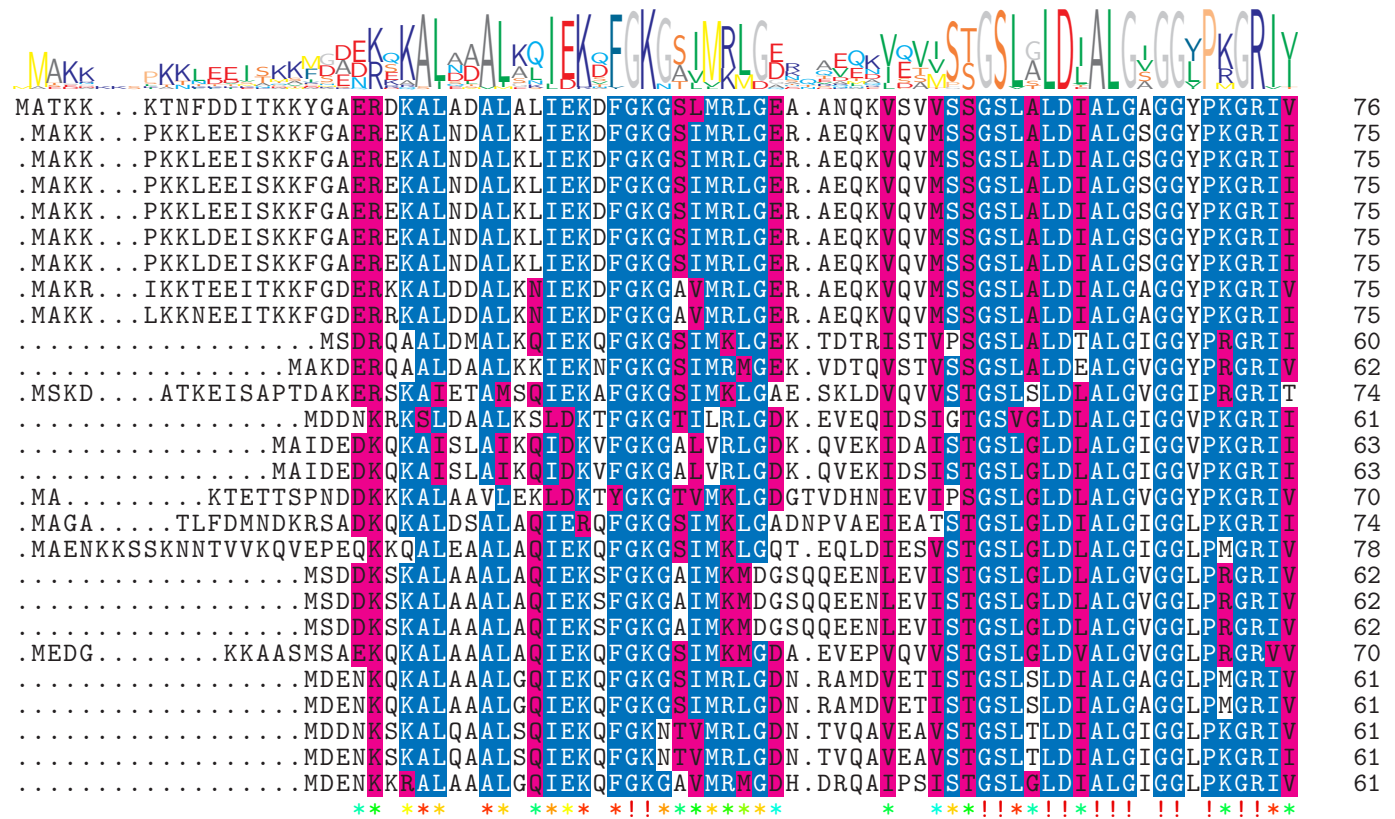


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consensus

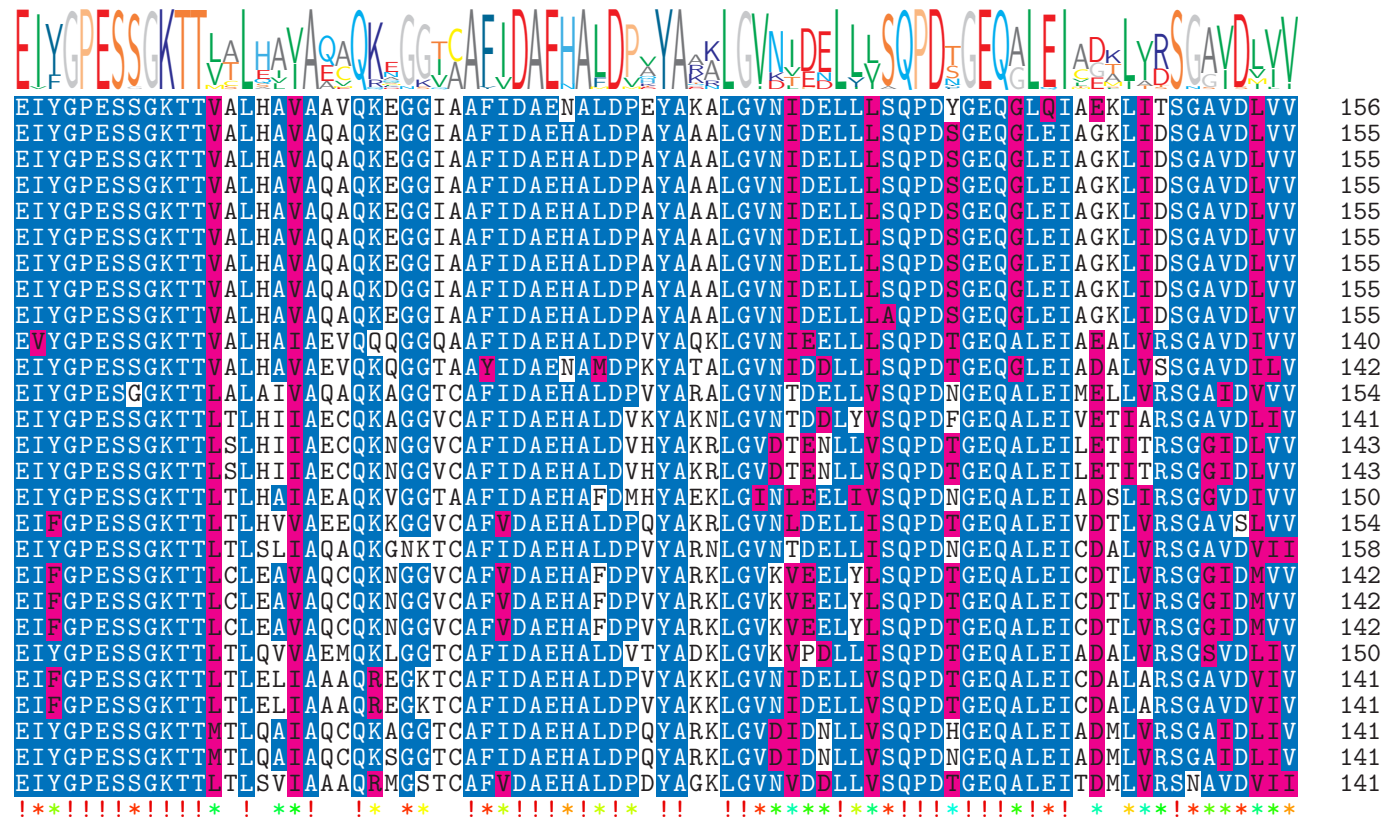
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consensus

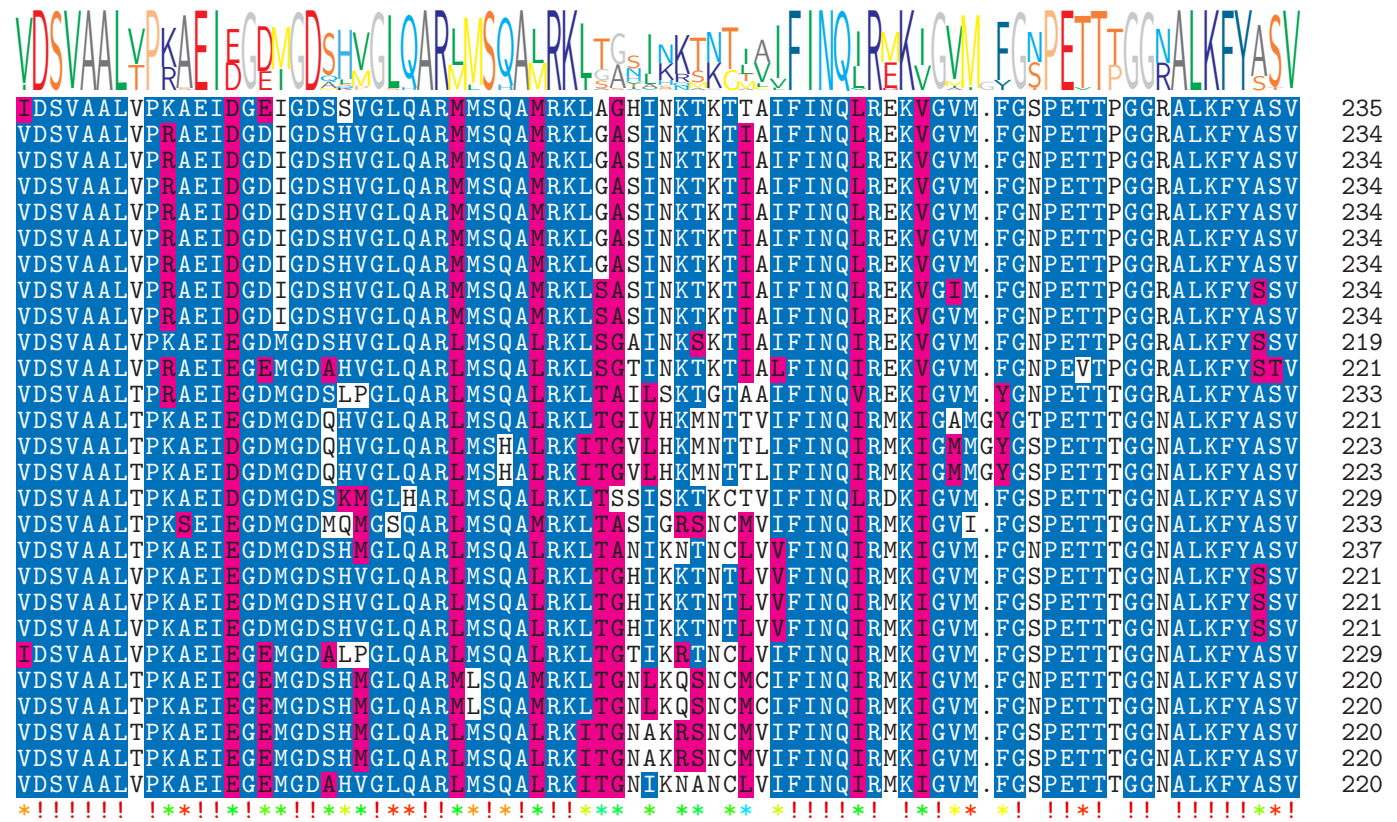
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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logo

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NTDB id 159 SP RS09750 WP 001085462.1	RLDVRG	NTQIKGTG	.DQKETN	VGKETK	IKVVKNKVAPPF	KEAVVE	EIMY	GEGIS	KTGELL	KIASDL	DI	IKKAGAW	YSYKDE	313	
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NTDB id 408 SMU RS09530 WP 002262392.1	RLDVRG	NTQIKGTG	.EQKDSN	IGKETK	IKVVKNKVAPPF	KEAFVE	EIIY	GEGIS	RTGELV	KIASDL	GII	IQKAGAW	YSYNGE	313	
NTDB id 453 Spy49 1753c ACI62003.1	RLDVRG	TTQIKGTG	.DQKDSN	IGKETK	IKVVKNKVAPPF	KVAEVE	EIMY	GEGIS	RTGELV	KIASDL	DI	IQKAGAW	FSYNGE	313	
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	RIDIRRI	A....	ALK.QNE	QHIGNRA	KA	KVVKNKVAPPF	REAEF	DIMF	GEGIS	KEGEI	IDYGV	KLDIV	DKSGAW	LSYQDK	298
NTDB id 1211 C694 RS00795 WP 000963128.1	RIDIRRI	A....	SLK.QNE	QHIGNRA	KA	KVVKNKVAPPF	REAEF	DIMF	GEGIS	KEGEI	IDYGV	KLDIV	DKSGAW	LSYQDK	298
NTDB id 1334 RAOC RS04870 WP 004918236.1	RIDIRRS	LSDTGKIK	DKSGEV	IGNRTK	VKVVKNKVAPPF	KSAEF	DIMY	GEGV	SKVGE	ILDV	ATD	LDI	IQKSG	SWFSYQDT	309
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consensus	!***!*		*	*	*	*****!***!!*!!*!!*!!*	*****!*	*****	*****	*****	!!	*****	! !*****!		

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consensus

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KIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEIE	387
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KIGQGSENAKKYLAHNPEIFDEIDHQVRVQFLIDGEEIAE...SKKDE...AAQVDS..VNEEVPLDLGDELEIEIE	383
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KLGGGRDRTVKELIKDNPELAEELEAKIREKLQAK.....	343
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logo

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consensus

 non conserved
 similar
 ≥ 50% conserved