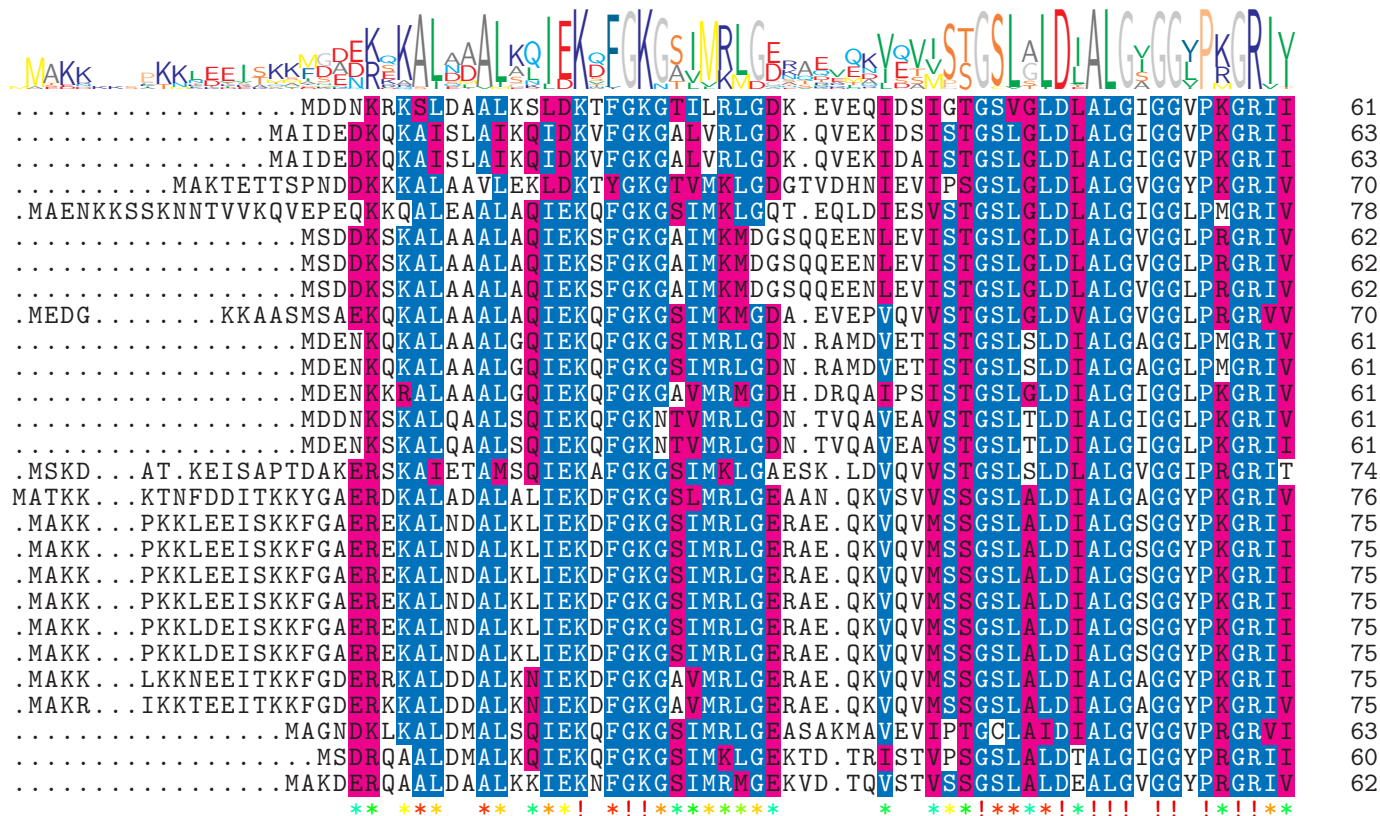


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consensus

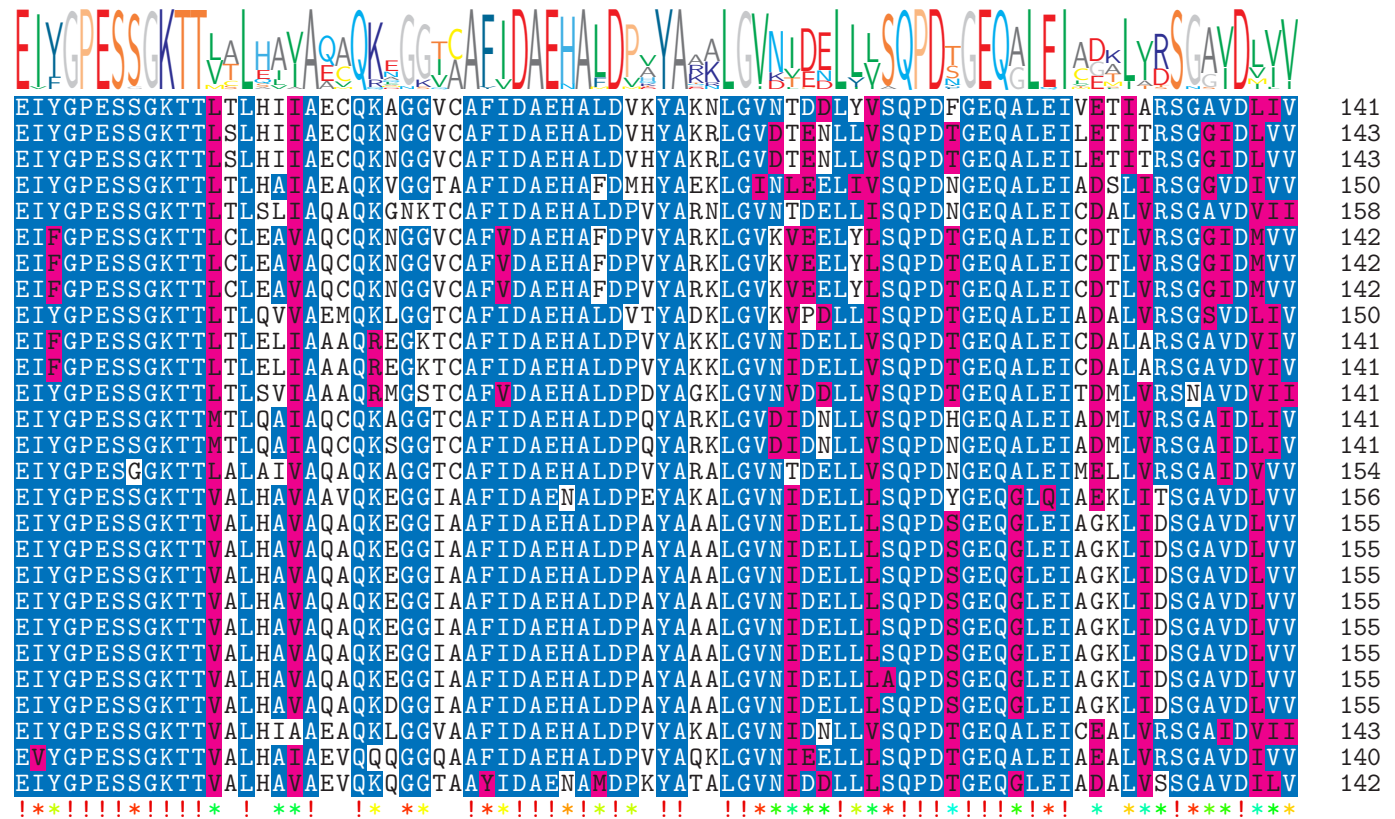
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consensus

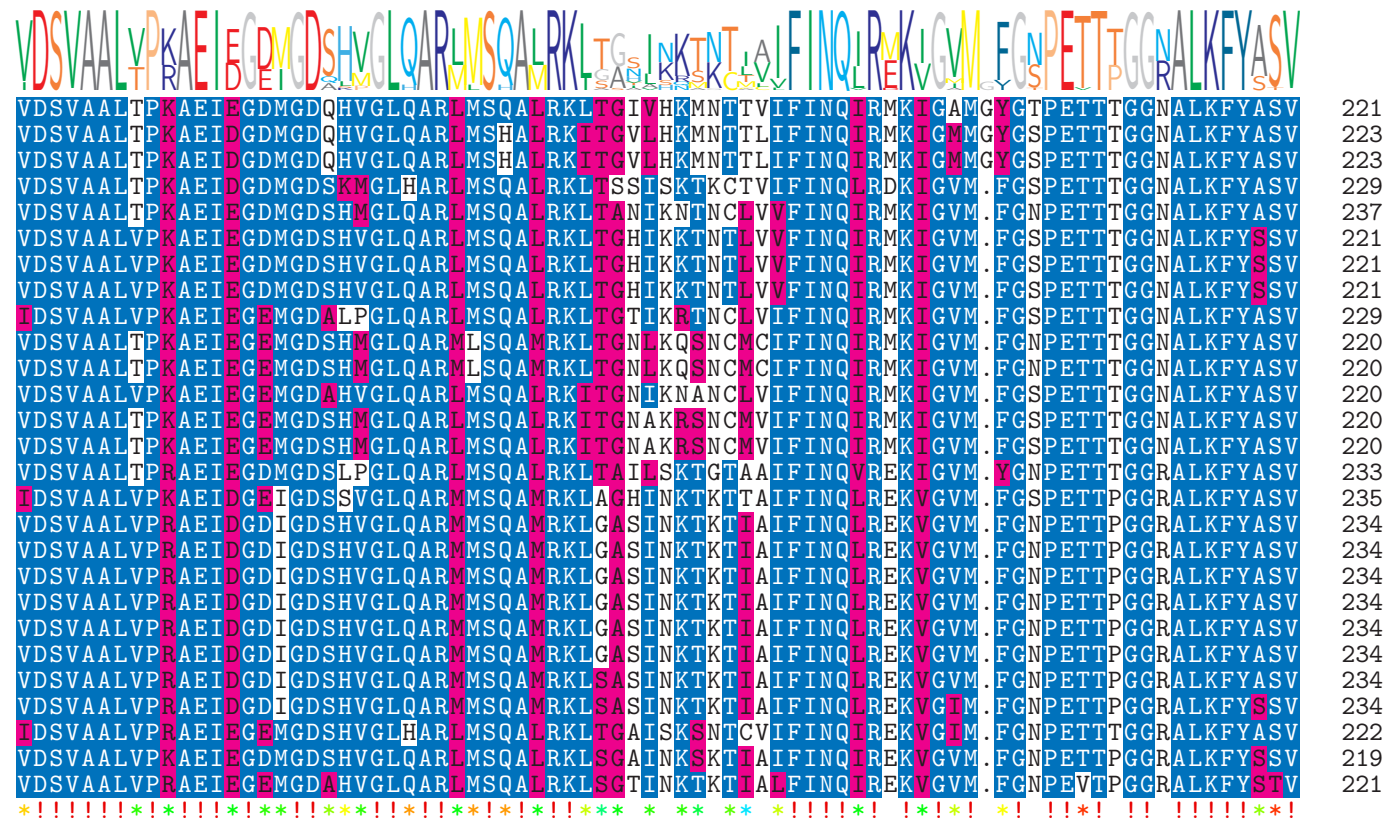
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consensus

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NTDB id 1124 NGFG RS03960 WP 003688695.1	RLDIRRTGS....I.KKGEV	LGNETRVKVI	KNKVAPPF	RQAEFDI	DILYGEGISWE	GELIDIG	VKNDI	IINKSG	AWSYNGA	296	
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NTDB id 269 KZH43 RS08665 WP 001085462.1	RLDVRGNTQIKGTG.DQKETN	VGKETKIKV	KNKVAPPF	KEAVVEI	EIMYGEGISKT	GELLKIAS	SDL	DII	IKKAGAW	SYKDE	313
NTDB id 228 SPD RS09265 WP 001085462.1	RLDVRGNTQIKGTG.DQKETN	VGKETKIKV	KNKVAPPF	KEAVVEI	EIMYGEGISKT	GELLKIAS	SDL	DII	IKKAGAW	SYKDE	313
NTDB id 194 SPR RS08825 WP 001085462.1	RLDVRGNTQIKGTG.DQKETN	VGKETKIKV	KNKVAPPF	KEAVVEI	EIMYGEGISKT	GELLKIAS	SDL	DII	IKKAGAW	SYKDE	313
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NTDB id 499 SM12261 RS08315 WP 078228268.1	RLDVRGSTQIKGTG.DQKDTN	VGKETKIKV	KNKVAPPF	KEAFVEI	EIMYGEGISKT	GELLKIAS	SDL	DII	IKKAGAW	SYKDE	313
NTDB id 453 Spy49 1753c ACI62003.1	RLDVRGTTQIKGTG.DQKDS	IGKETKIKV	KNKVAPPF	KVAEVEI	EIMYGEGISRT	GELVKIAS	SDL	DII	QKAGAW	FSYNGE	313
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consensus	!***!*	*	**	*****!***!!*!!*!!*	*	*****!***!***	*****	*	***	! !*****!	

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NTDB id 1245 Cj1673c YP 002345041.1	KLGGGRENKAFLENPEIADEITKATQNSMGI	IEGMISGSEDDE..GEE.....	343
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NTDB id 1334 RAO C RS04870 WP 004918236.1	KLGGGRDITVKELIKDNPELAEELEAKIREKLQAK		343
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NTDB id 499 SM12261 RS08315 WP 078228268.1	KIGQGSSENAKKYLAENPEIFDEIDHQVRVQFGL	IDGEEIAE....SKKDE....AAQVDS..VNEEV	TLDLGDELEIEIE
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consensus

 non conserved
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 ≥ 50% conserved