

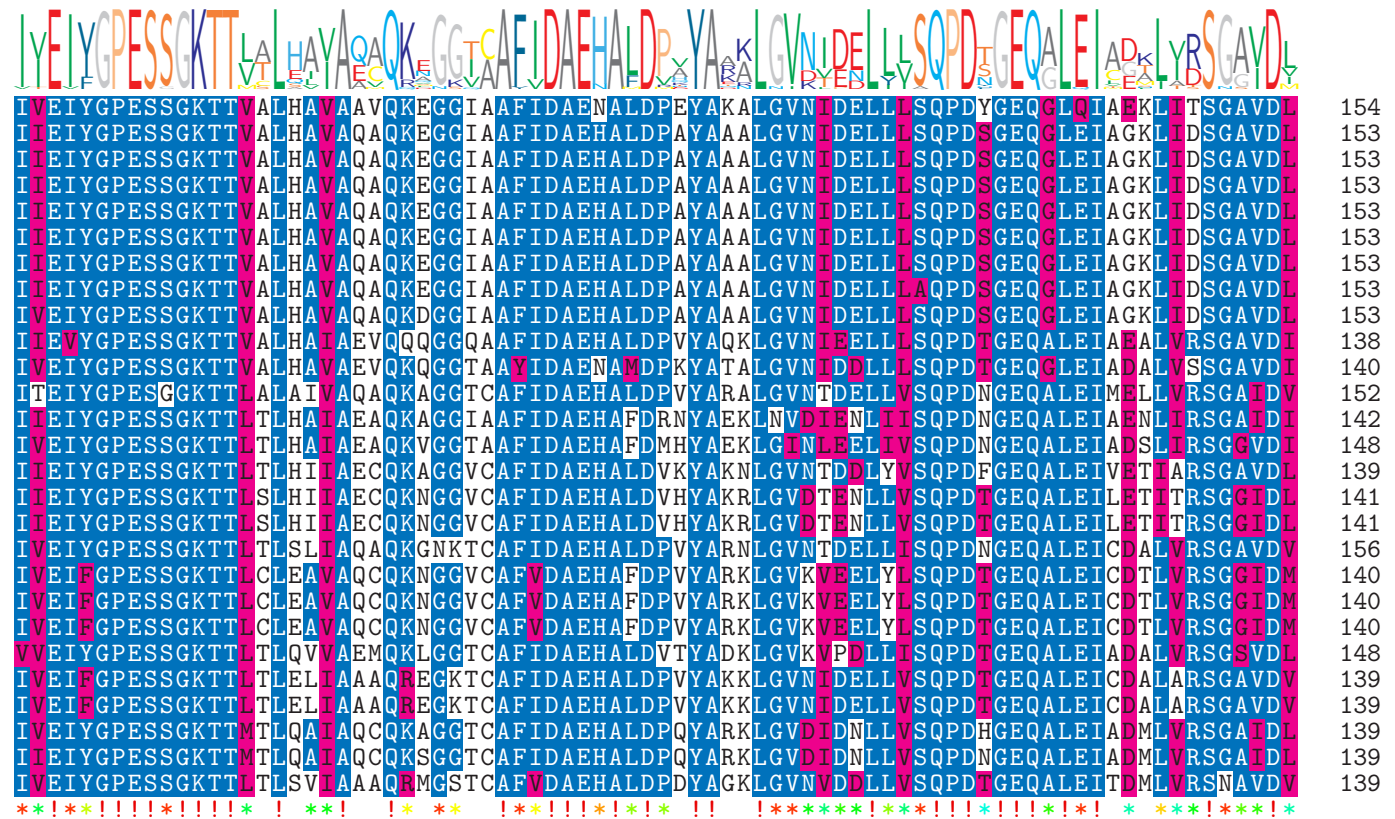
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NTDB id 408 SMU RS09530 WP 002262392.1	.MAKRIK.....KTEEITKKFG	DERRKALDDALKN	IEKD	FGKGAVMRLGER	.AEQKVQV	MSSGSLALD	I	ALGAGGY	PKGR	73
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NTDB id 300459 HYN86 RS04445 WP 113676957.1	MSSDK	EAKLKALQLTL	LDKTY	GKGTVMKMGDKAIV	.EVETISSGSLGV	DL	ALGVNGY	PKGR	62	
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NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKK	RALAAALGQIEKQ	FGKGAVMRMGDH	.DRQAIPS	ISTGSLGLD	I	ALGIGGL	PKGR	59	
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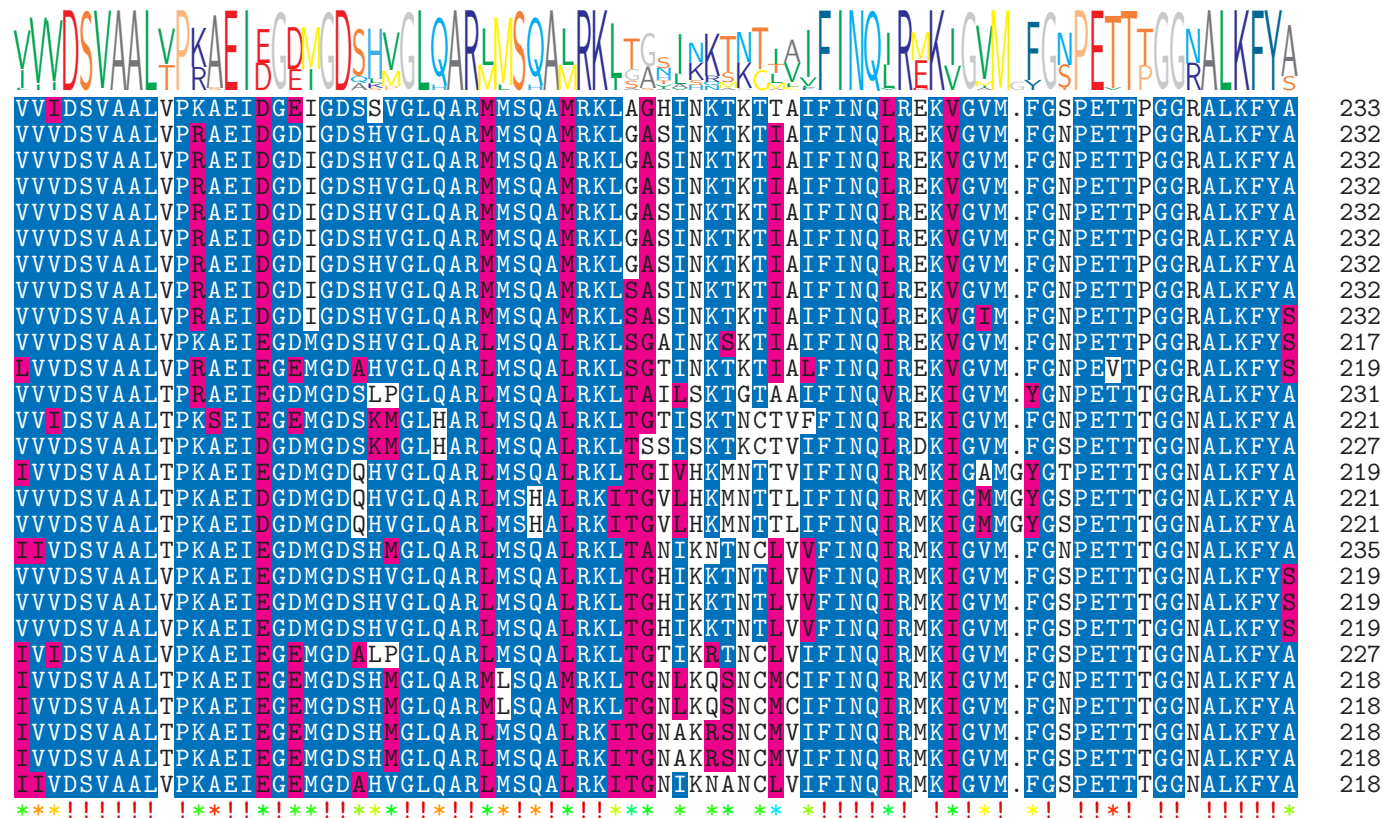
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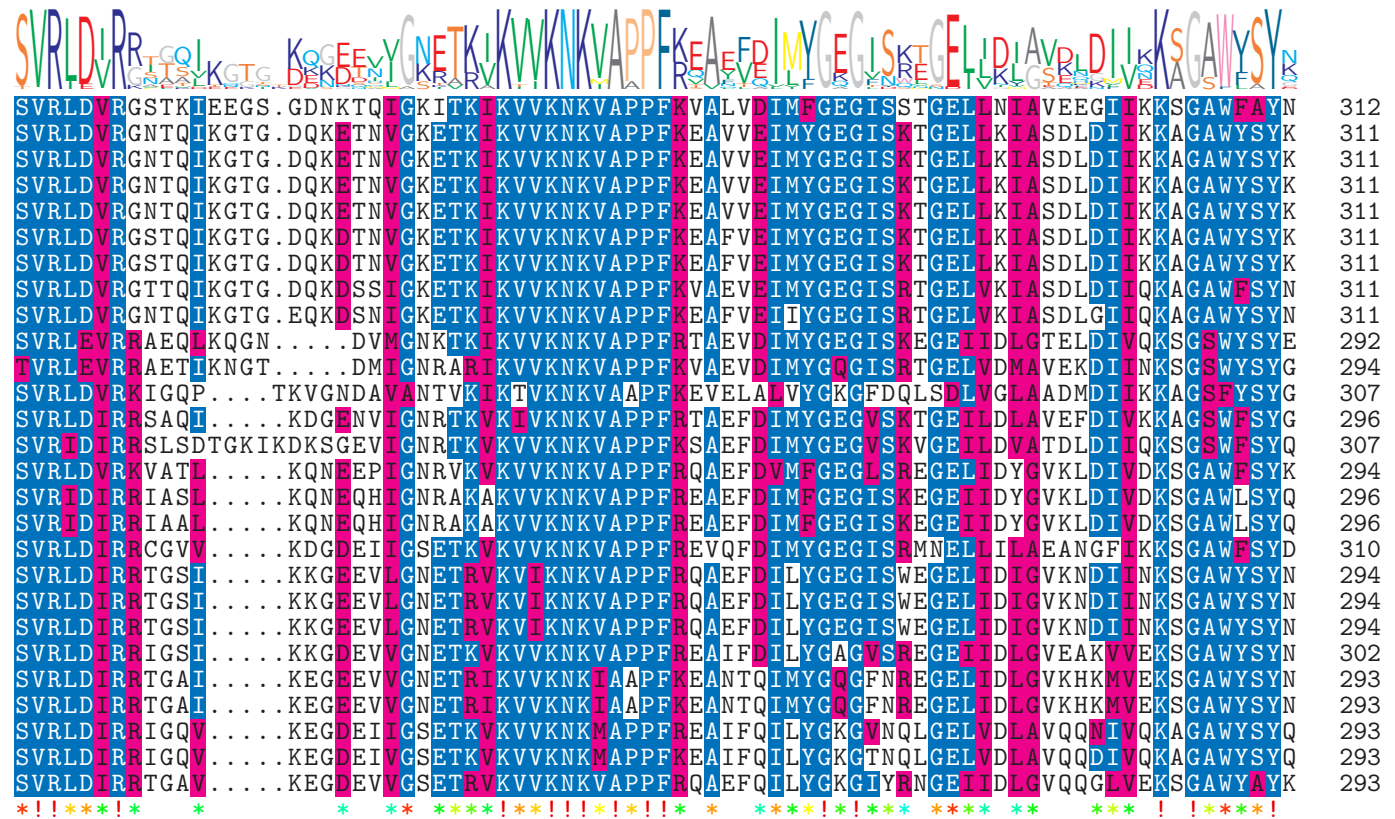
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GEKIGQGS ENAKKFLADNP EIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK..... ETDEK..ADKNLVLELDDTIELE 382

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DKKLGGGREN AKALLKEDKALADEITLKIKESIGSNEE.... IMPLPDEP..LE..EME..... 347

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consensus			

- X non conserved
- X similar
- X ≥ 50% conserved