

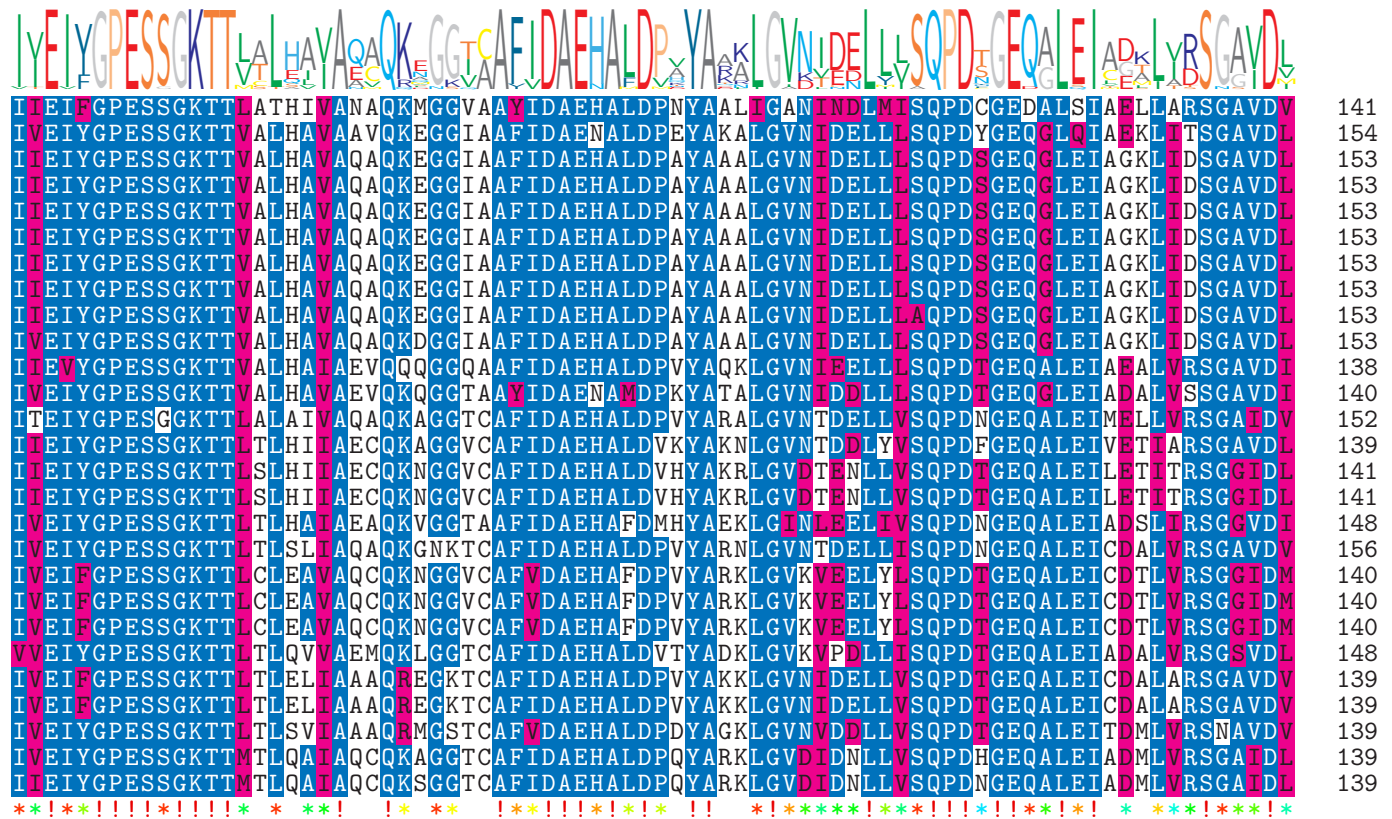
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NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTD	AKERSKATETAMSQIEKAFGKGSIMKLGAE	.SKLDVQV	VSTGSLSLDIALGVGGI	PKGR	72	
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDEN	KSKALQAALSQIEKQFGKNTVMRLGDN	.TVQAVE	AVSTGSLTLDIALGIGGL	PKGR	59	
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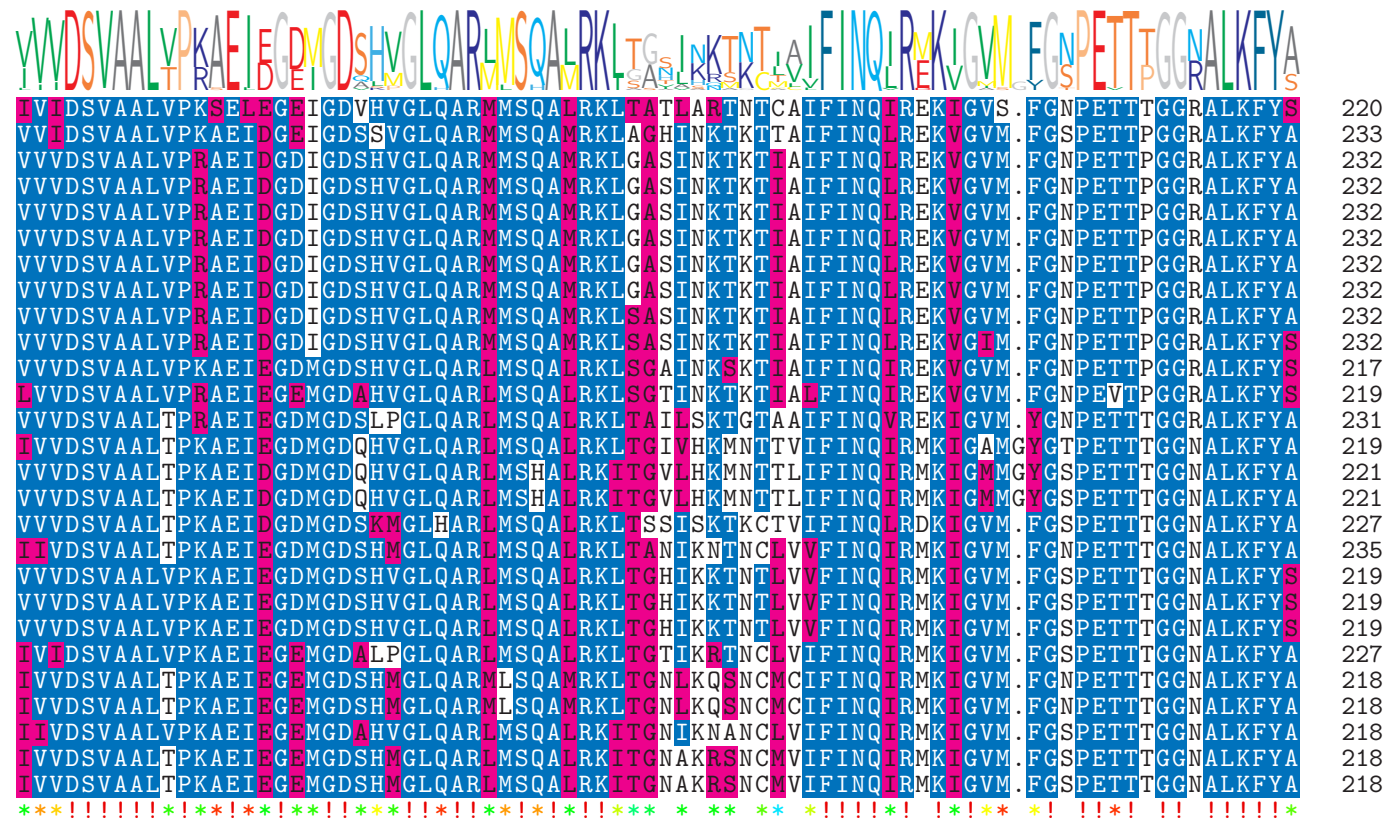
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consensus

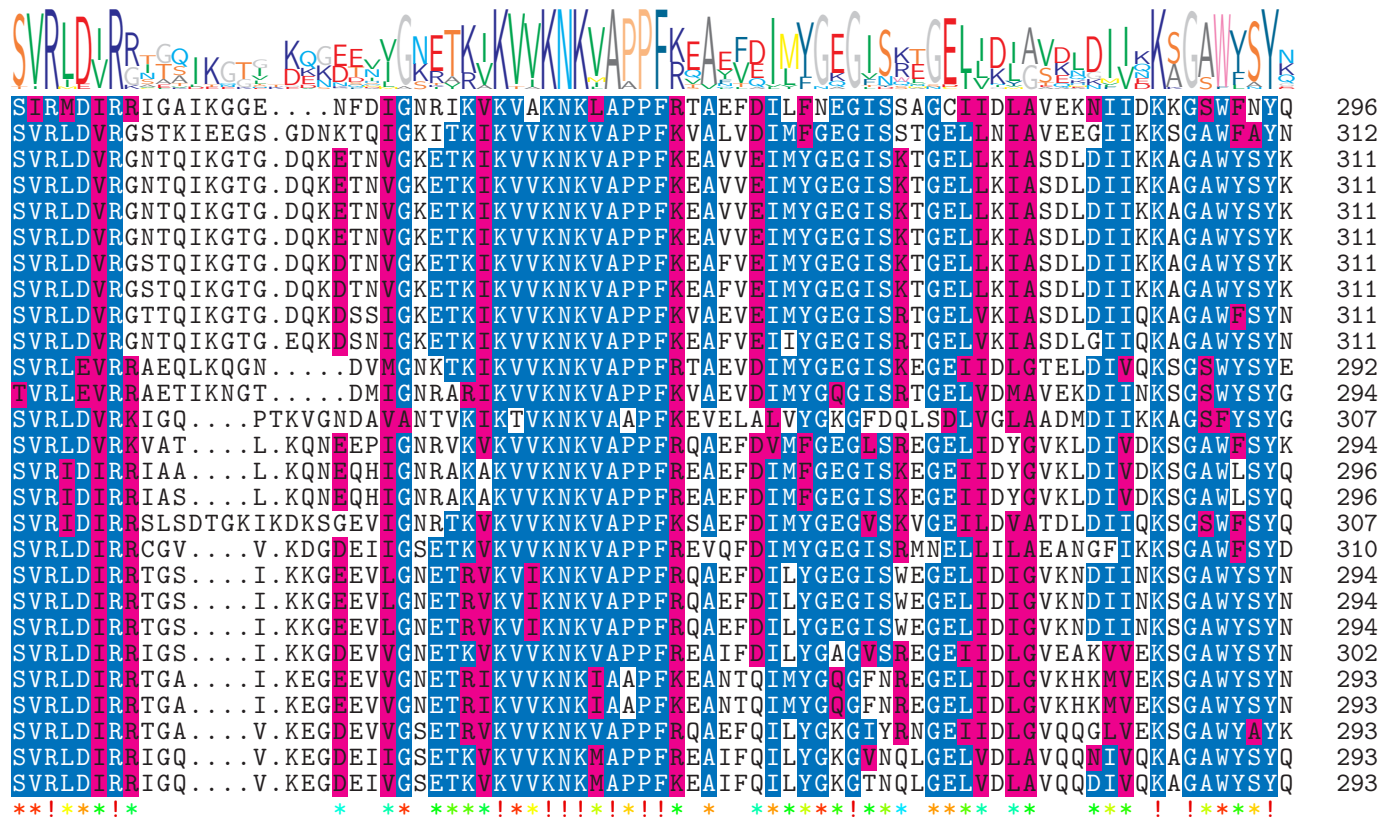
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consensus

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Sequence	Score
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DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP..LE..EME	347
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logo

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consensus			

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