

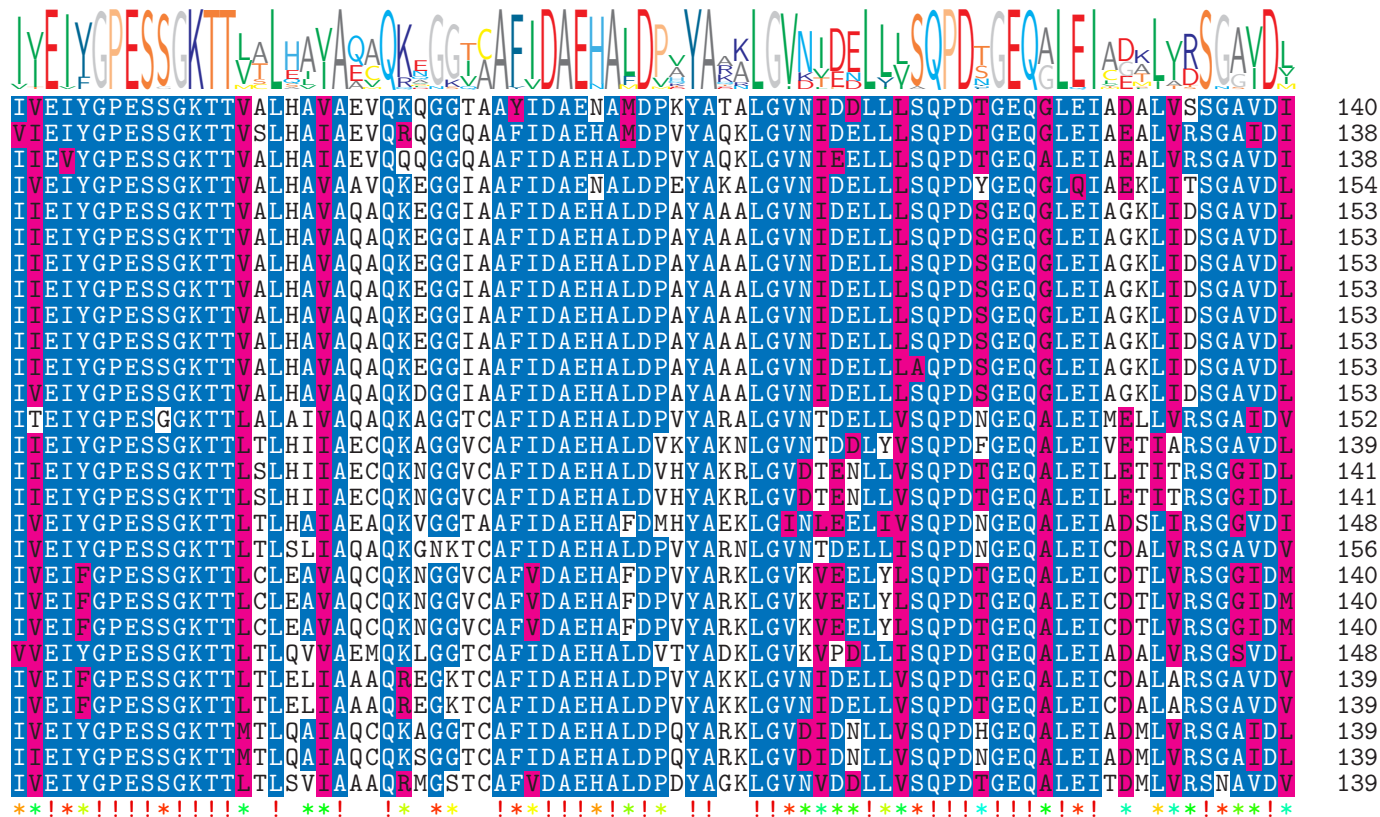
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NTDB id 1205 PSJM300 05545 AFN77183.1	MDEN	KKRALAAALG	QIEKQFGKGAVMRMGDH	.DRQAIPSISTGSLGLDIALGIGGL	PKGR	59				
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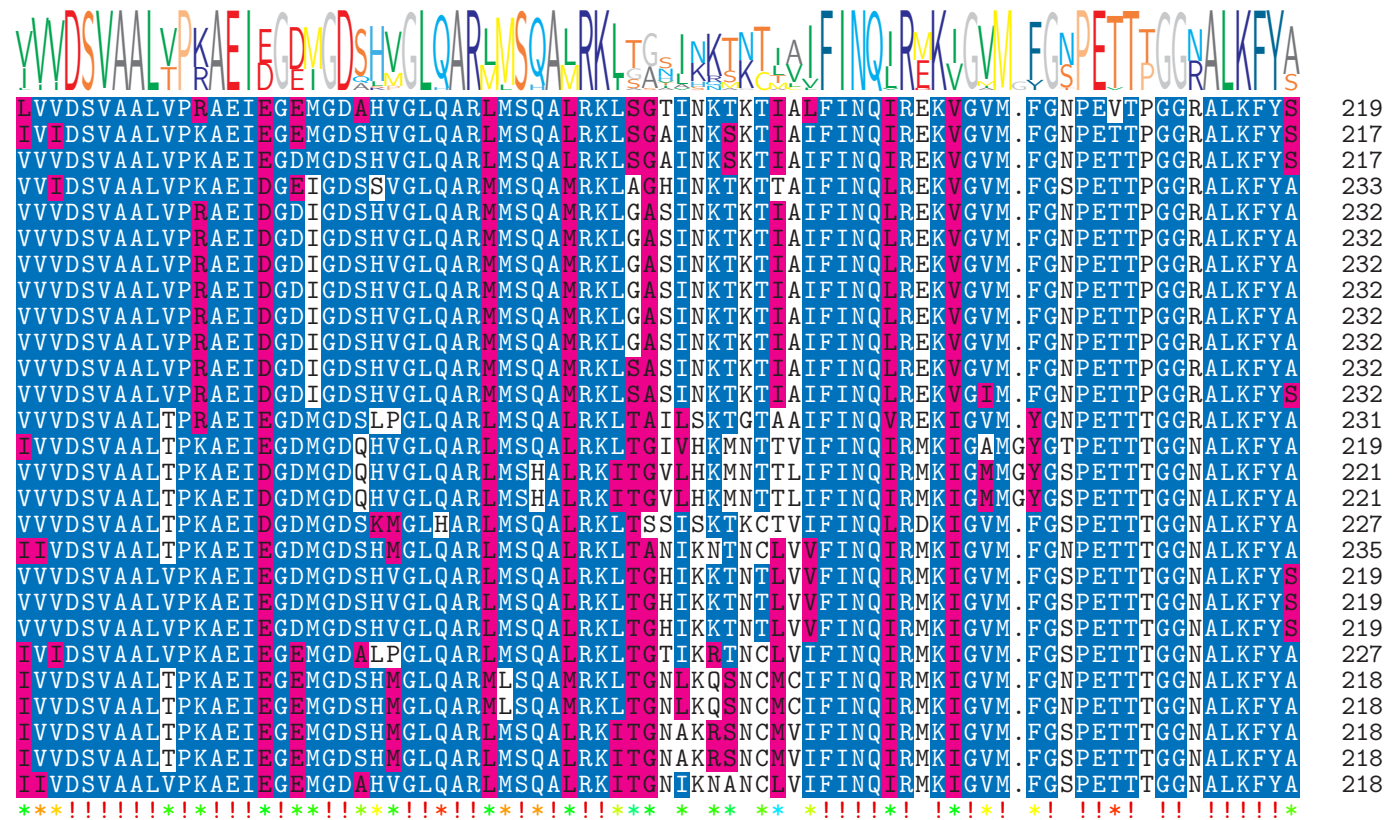
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consensus

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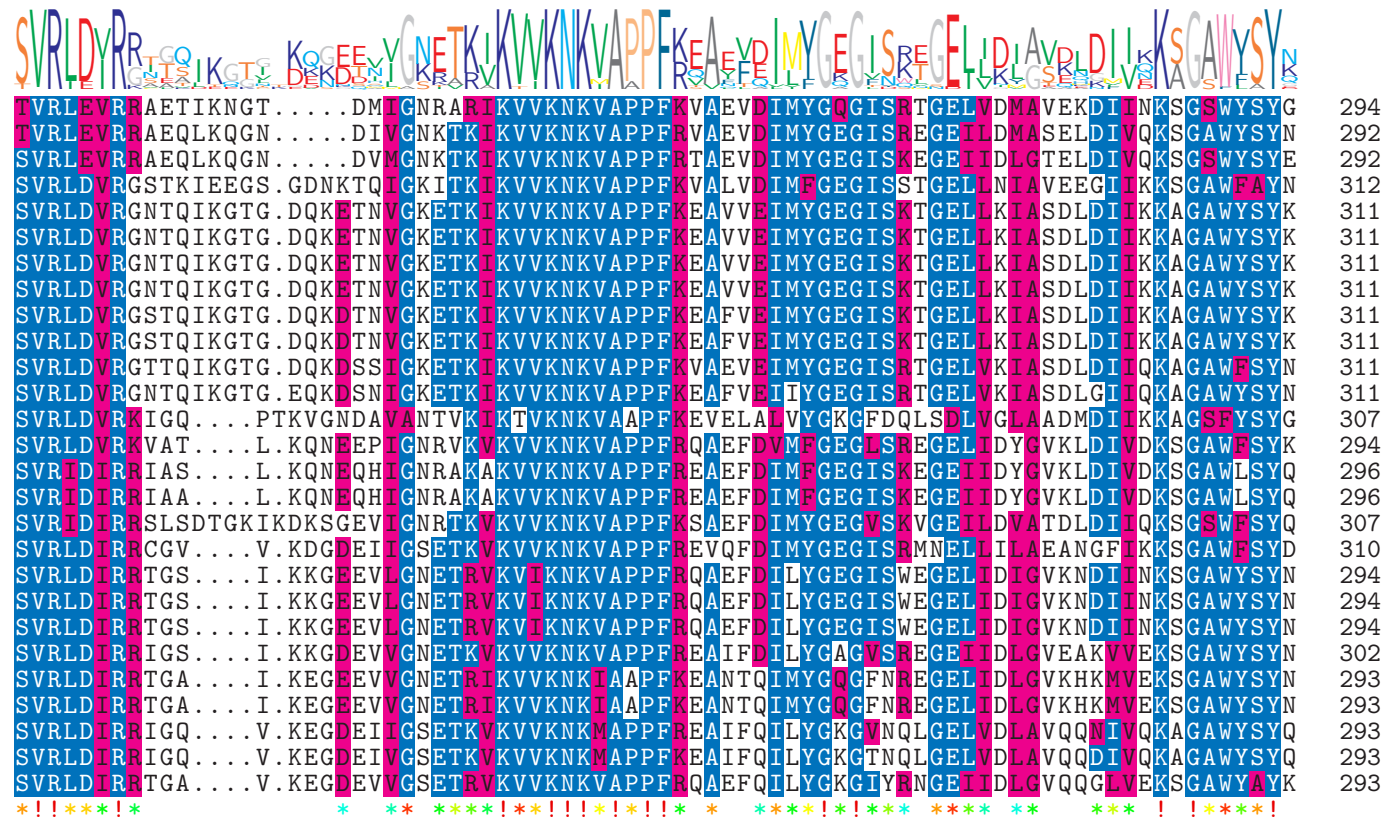
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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Sequence	Score
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EERLGGQRENAKQFLKENPKDMLMIQEIQIREHYGLDNGGVVQQAEETQEEL.....EFEE.....	348
DEKIGQGAEKAKNYLKEHQDVFEIDHKKVRAAHGLDDSEVAETEEETTASKTAKAKKEEKAVETEEIELELED....	387
DEKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQD TENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
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GEKIGQGSSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKE.....ETDEK..ADKNLVLELDDTIELE	382
DERIGQGGKEKTIAYIAIERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAPEA.....AFA.....	363
DKKLGQGRNSKAFLEKENPEIAD EITKAIQNSMGI EGMI.....SGSEDD..E.....GEE.....	343
DKKLGQGRNENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDE..P.....LEEME.....	347
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DTKLGGQGRDITVKELIKDNPELAEELEAKIREKLQAK.....	343
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logo

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved