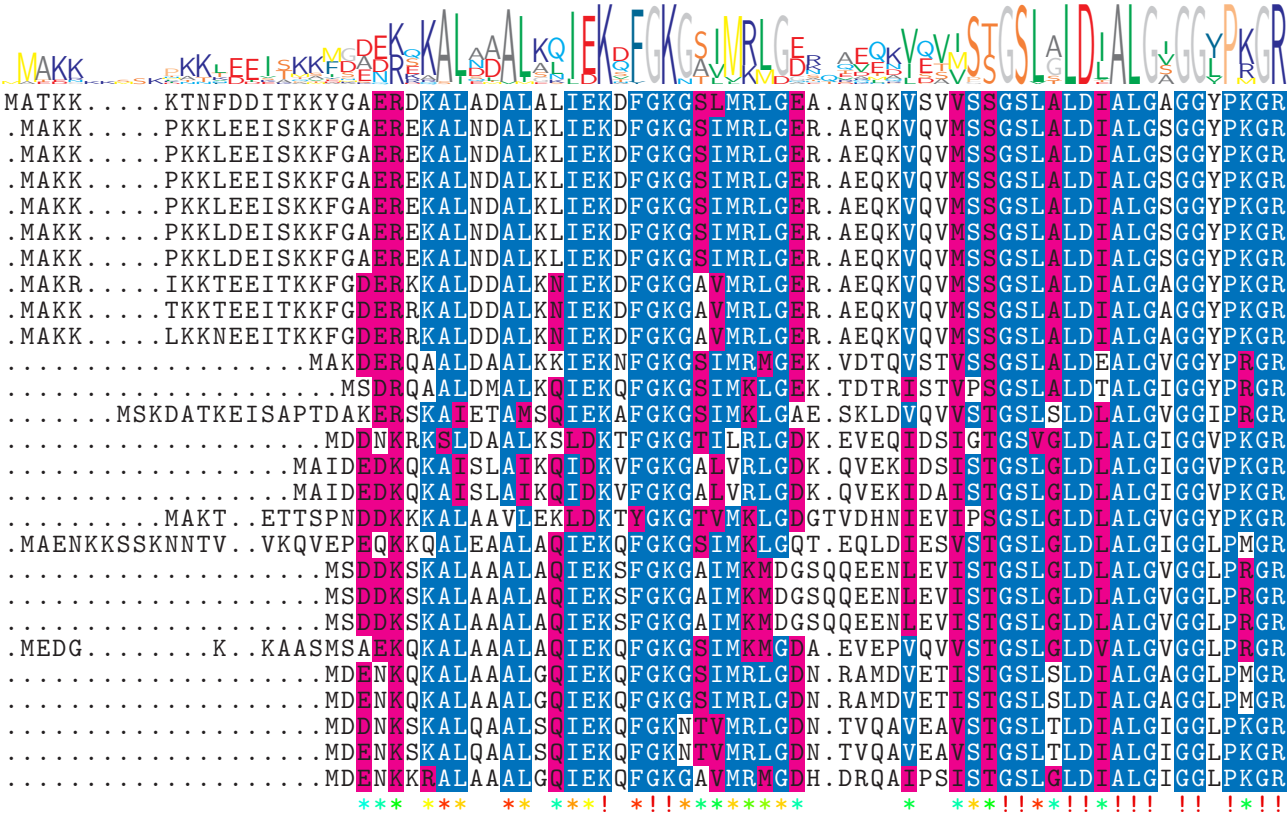


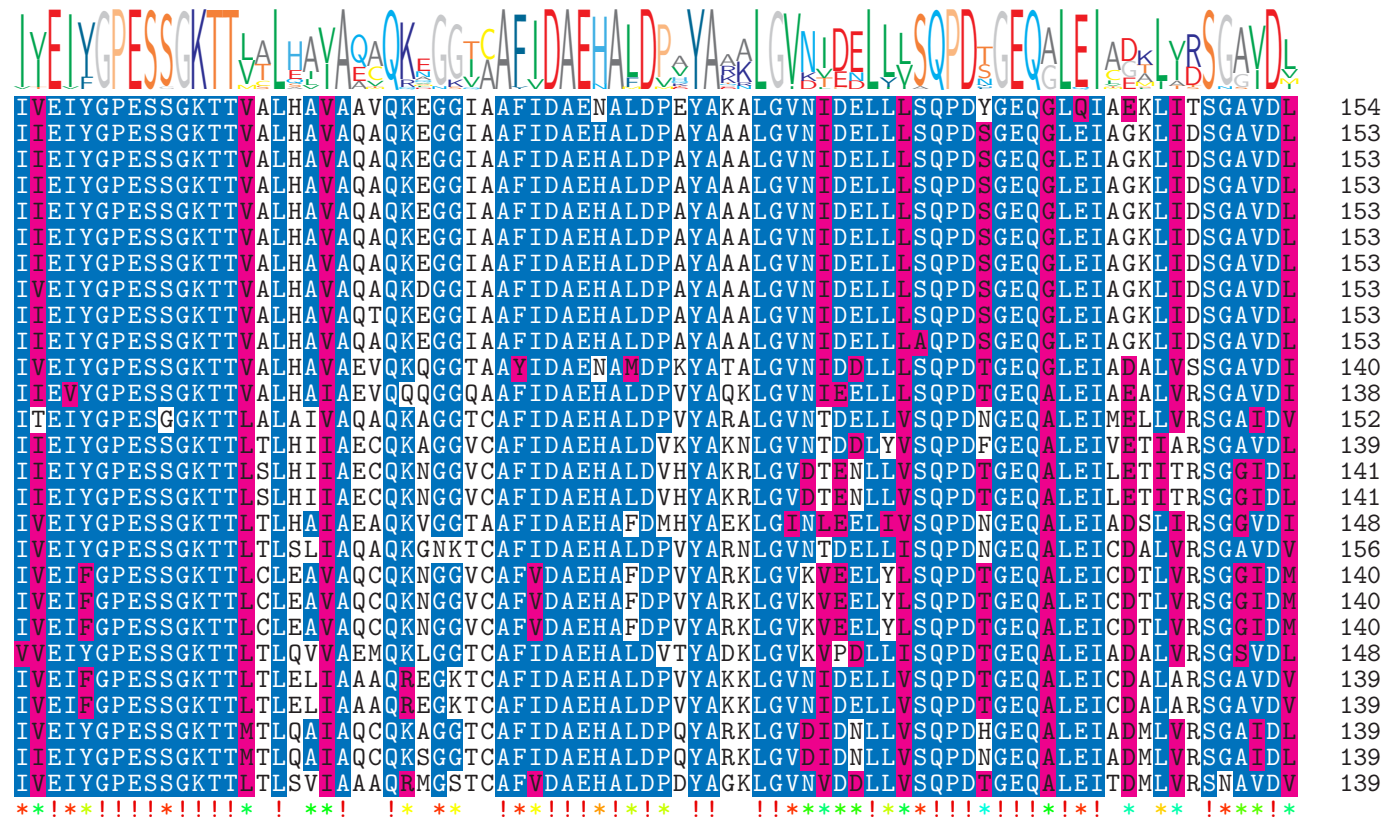
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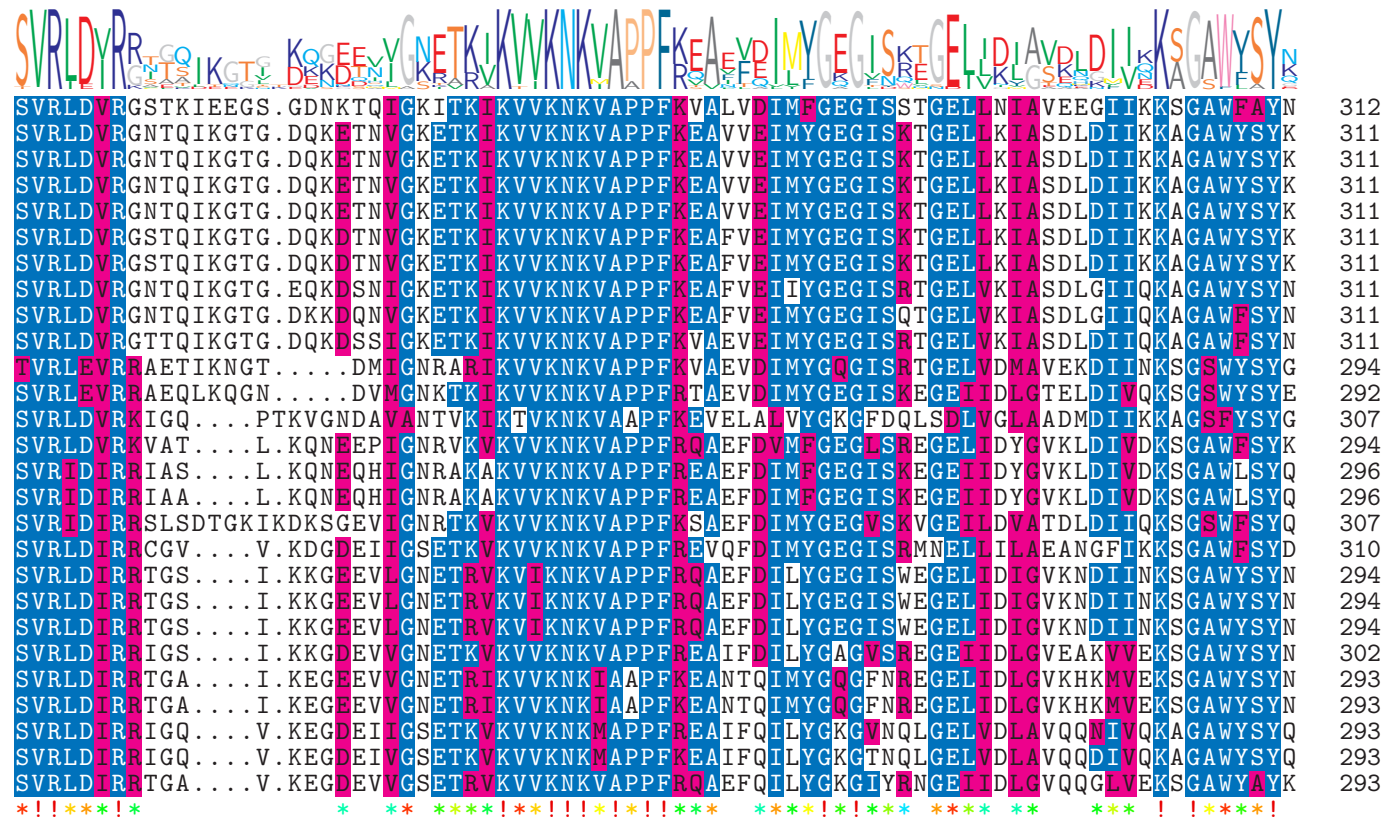
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consensus

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DEKIGQGSSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAES...KKDEAAQ...V..DSVNEEVTLDLGDELEIE	381
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GEKIGQGSSENAKKFLADNPEIFDDIDHKVRVQFGLIEEDNTTEKKQSSKEK...ET...D..EKADKNLVLELDDTIELE	382
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GEKIGQGSSENAKRYLADHPQLFDEIDRKVRVVKFGLLEESEESAMAVA.....SEETDDLALDLNNGIEIE	377
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DKKLGQGSRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDE..PLE.....EME	347
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GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI....TEGTQDETGE.....R.....PEE.....	348
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logo

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved