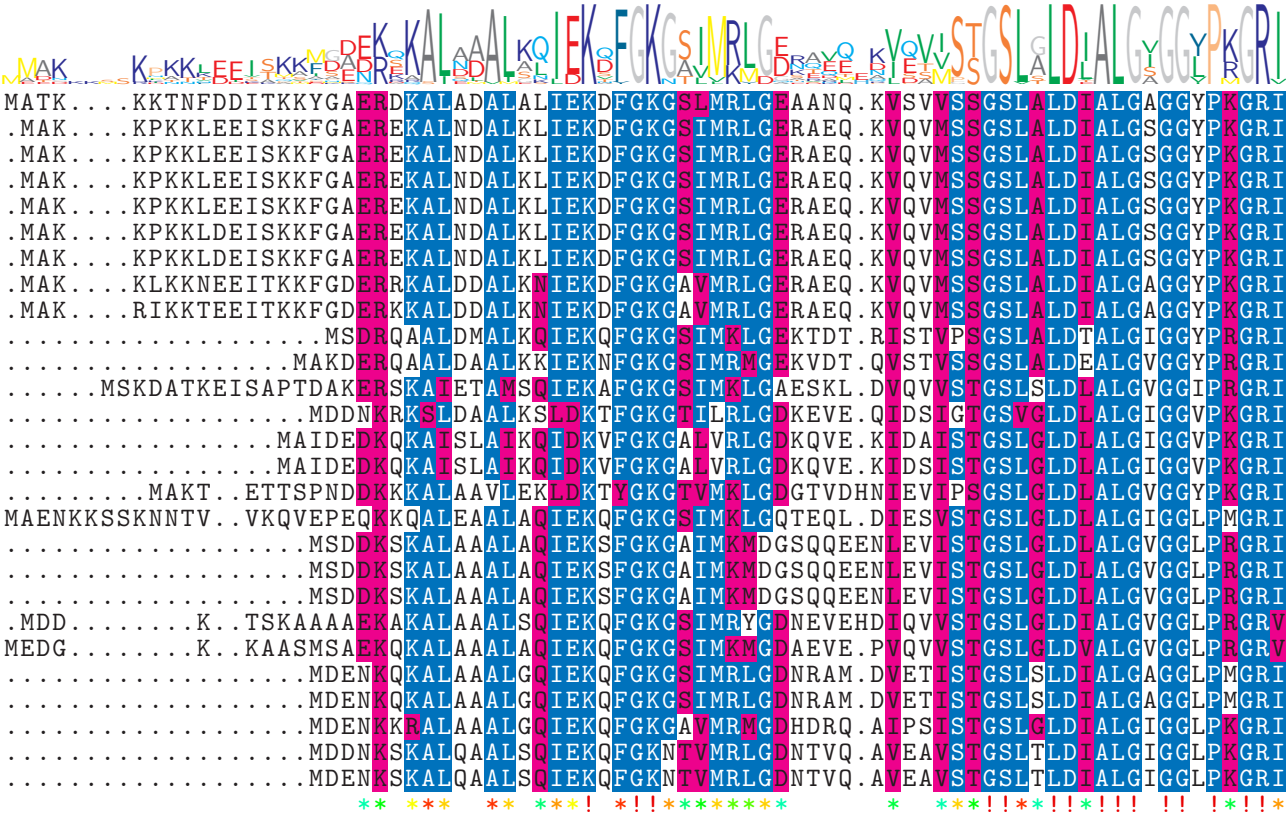


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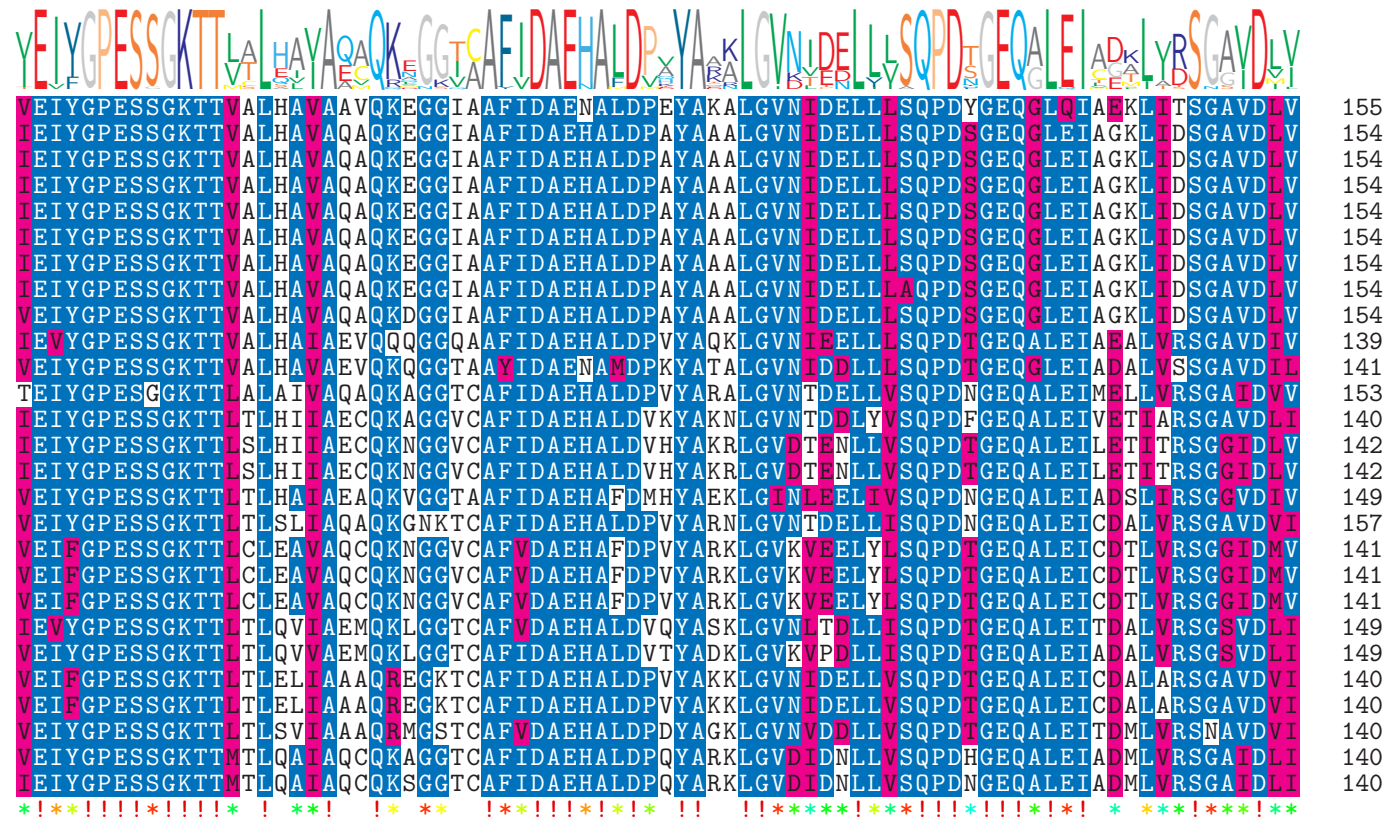


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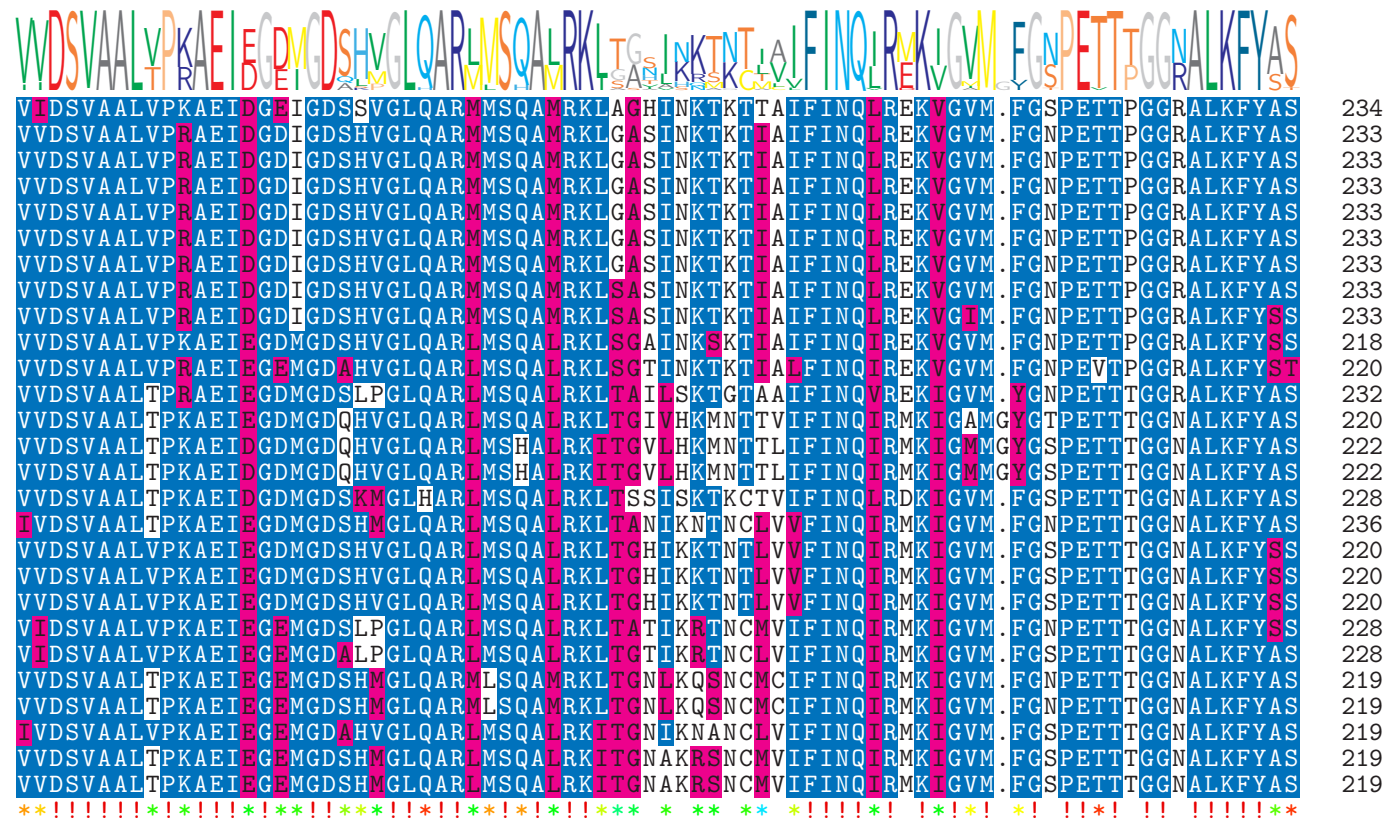
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EKIGQGSENAKKFLADNPEI FDDIDHKVRVQYGLIEEDNTEE..KQSSKEK.....ETDEK..ADKNLVLELDDTIEL	381
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ERIGGGREN AKNYLADHEDVEDEVRLKVRAAYGISDVPEEDL..PTTEDE.QI.NILPDDST.....EE.....	355
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KKLGGGREN AKALLKEDKALADEITLKIKESIGSNEEIM..PLP....DEPLE.EME.....	347
KKLGGGREN AKALLKEDKALADEITLKIKESIGSNEEIM..PLP....DEPLE.EME.....	347
TKLGGGRD TVKELIKDNPELAELEAKIREKLQAK.....	343
TKIGQGNNAVKWLKENPEIADKLEQEI RNTLASNPNSV..LKATASDSND ESDVEEF.....SEEL.....	372
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-  non conserved
-  similar
-  ≥ 50% conserved