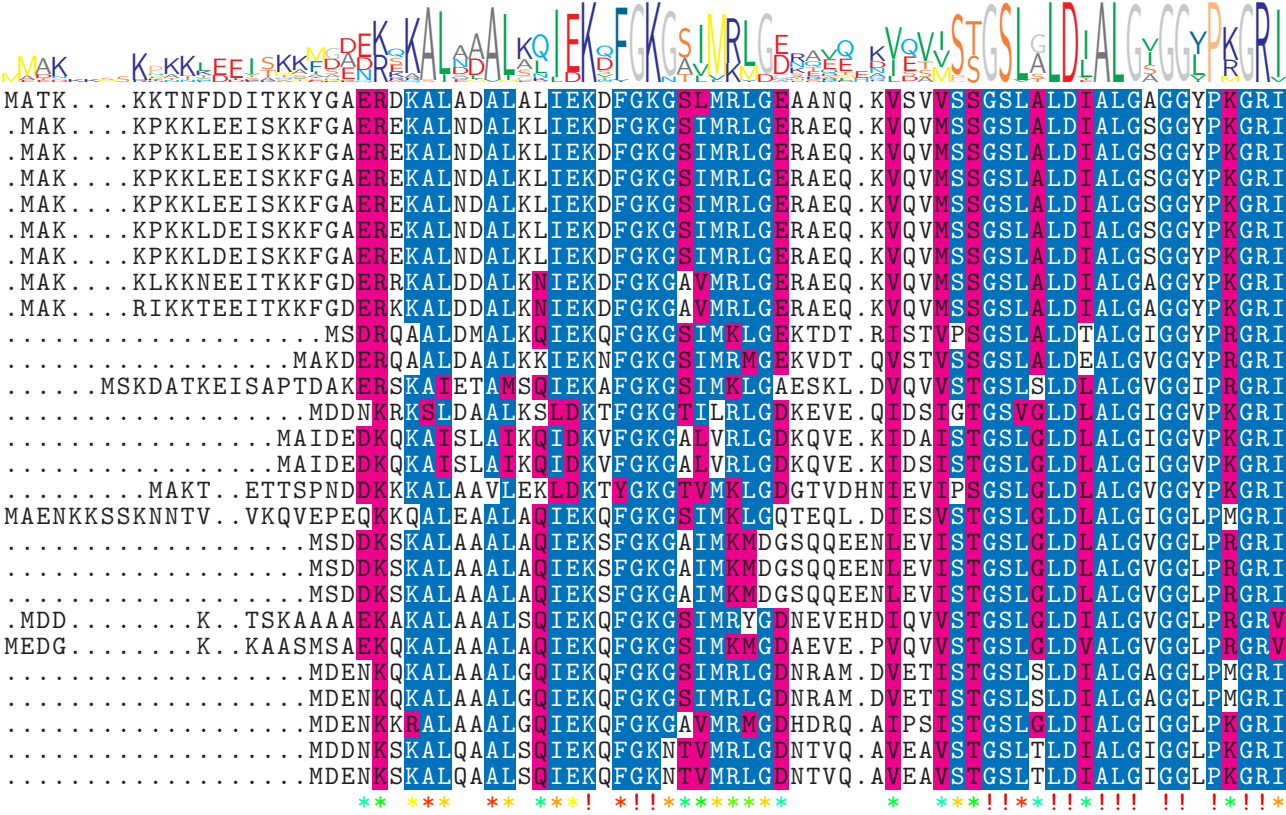


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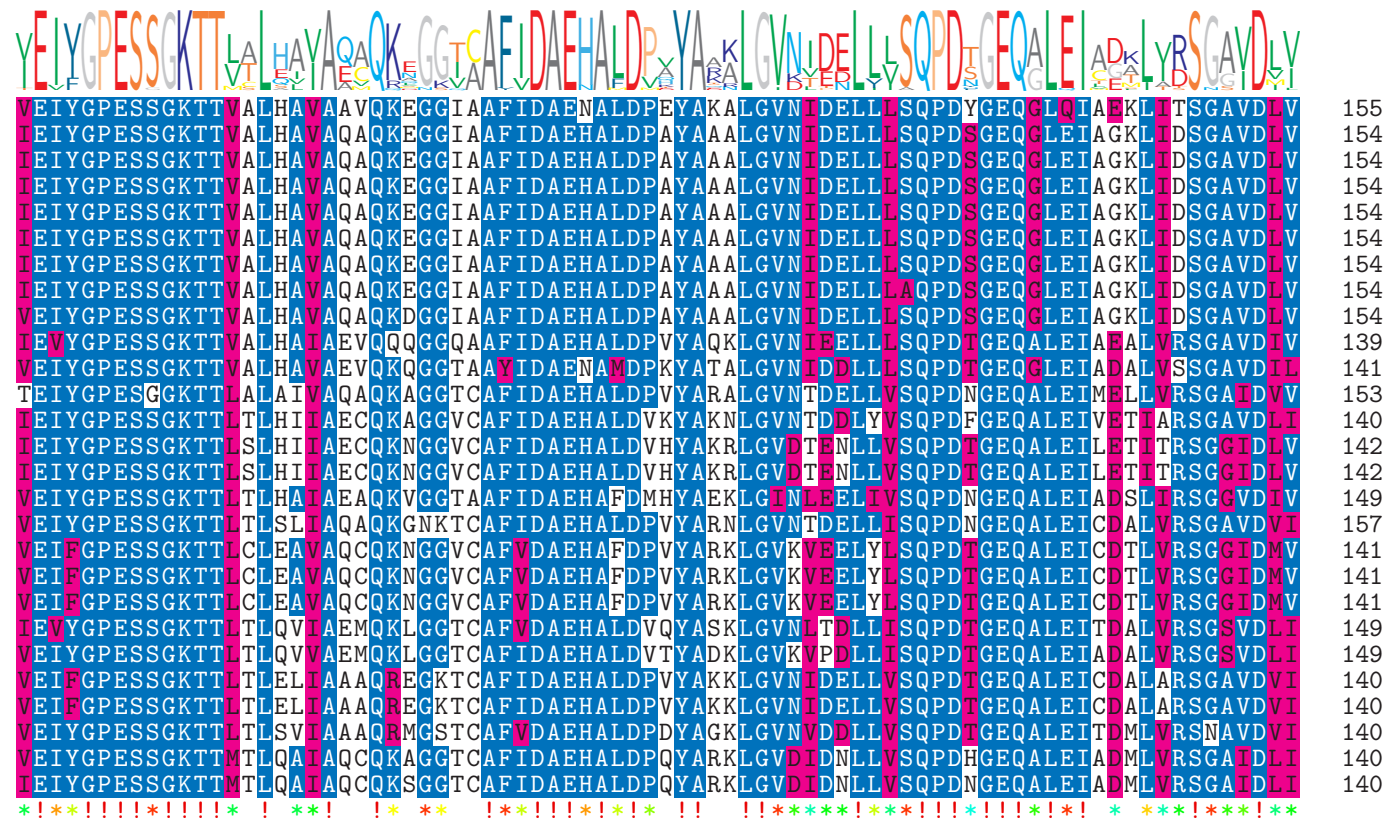


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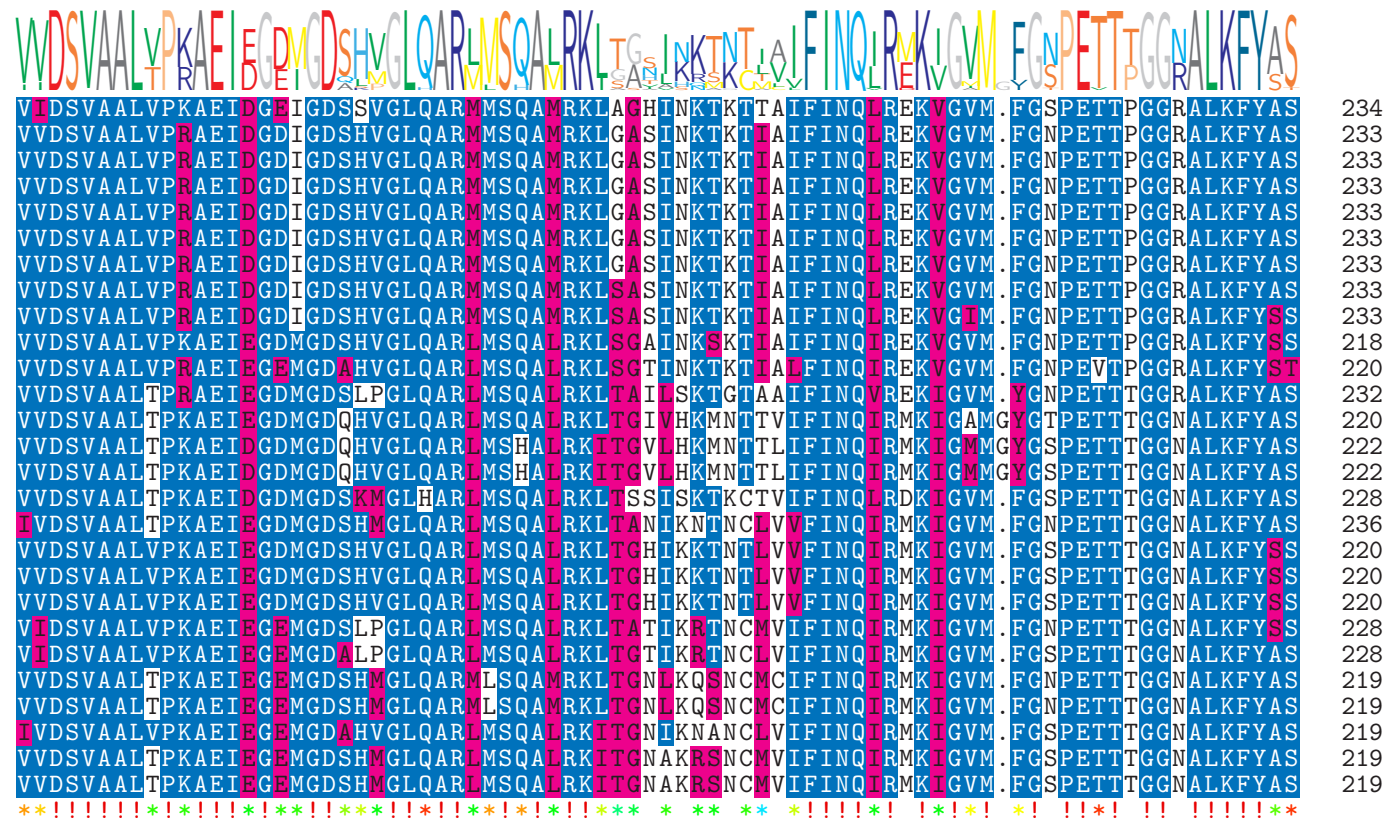
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consensus

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EKIGQGSENAKKFLADNPEI FDDIDHKVRVQYGLIEEDNTEE.. KQSSKEK..... ETDEK.. ADKNLVLELDDTIEL	381
ERLGGGR ENAKQFLKENKDIMLMIQEQIREHYGLDNGGVVQQ.. QAEETQEEL.EFE.E.....	348
ERIGGGREN AKNYLA DHEDVEDEVRLKVR AAYGISDVPEEDL.. PTTEDE.QI.NILPDDST..... EE.....	355
ERIGGQKEKTIAYIA ERPMEQEIRDRVM AAI RAGNAGEAPALAPAPAPEAAEA.EA.....	363
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KKLGGGR ENAKALLKEDKALADEITLKIKESIGSN E EIM.. PLP... DEPLE.EME.....	347
KKLGGGR ENAKALLKEDKALADEITLKIKESIGSN E EIM.. PLP... DEPLE.EME.....	347
TKLGGGRD TVKELIKDNPELAEELEAKIREKLQAK.....	343
TKIGQGKNNAVKWLKENPEIADKLEQEI RNTLASNPNSV.. LKATASDSNDES.DVEEF..... SEEEL.....	372
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consensus						

-  non conserved
-  similar
-  ≥ 50% conserved