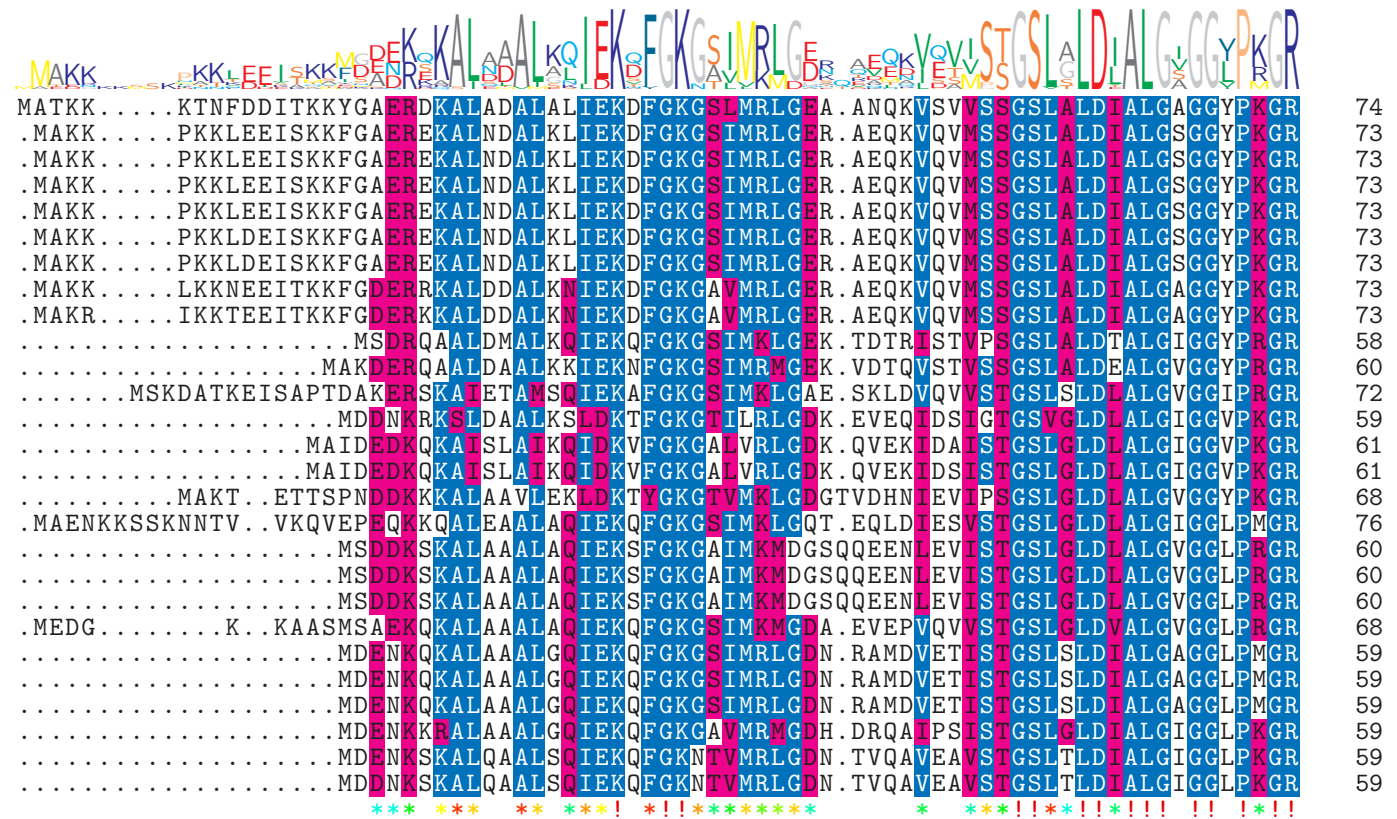


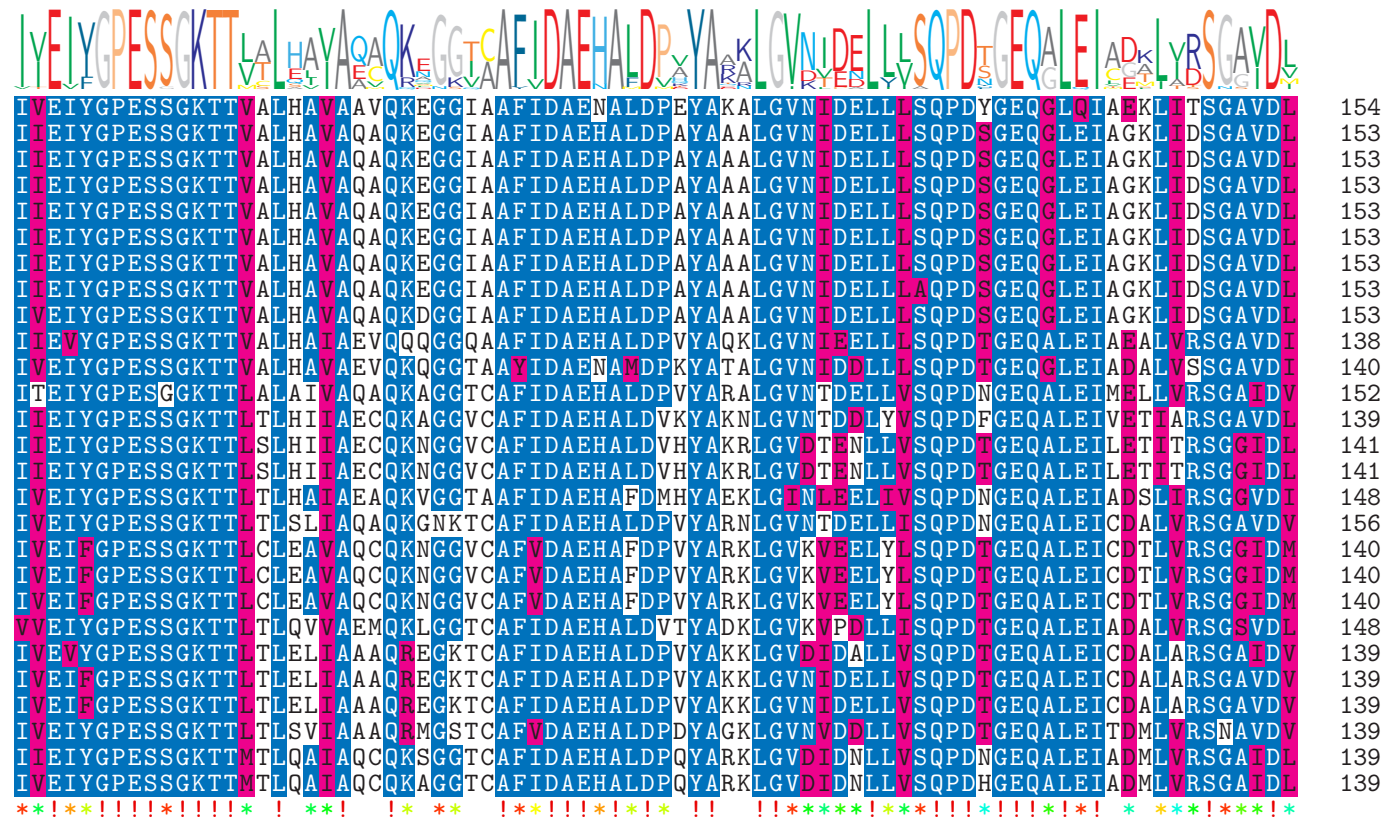
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consensus

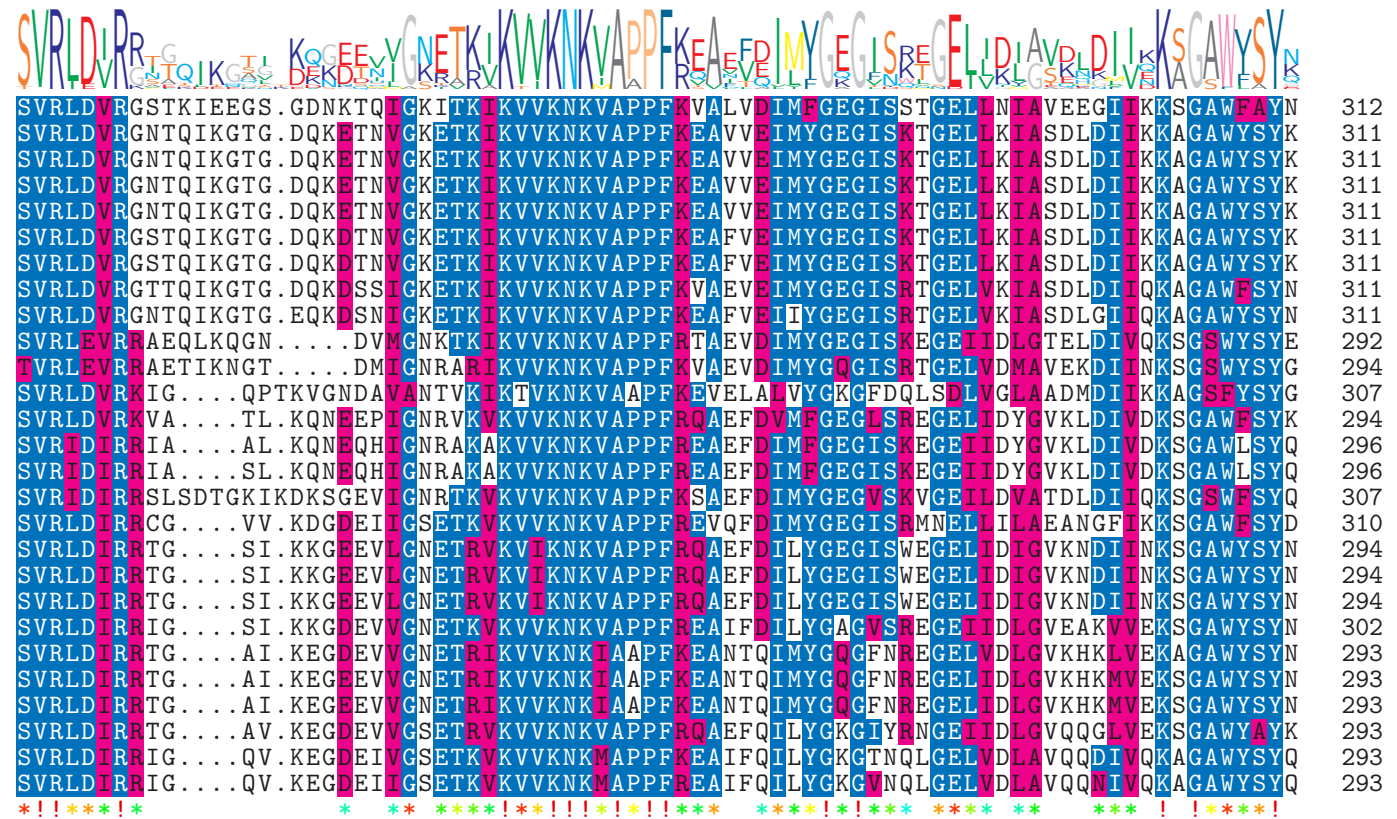
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DEKIGQGSSENAKKYLAHNPEVFAEIDHQRVQVQFLIDGEEAAE...SKKDE...VAQVDS..VNEEVTLDLGDLEIE	381
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GEKIGQGSSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSSENAKKFLADNPEIFDDIDHKKVRVQVGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGGQRENAAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGQGSRENAKNYLADHEDVEDEVRLKVRVQVGLSDVPEEDLPTTEDE.QI.NILPDDST....EF.....	355
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GTKIGQGKNNAVKWLKENPEIADKLEQEI RNTLASNPNSVLKATASDSNDES.....DV.....EFSE...EEL...	372
GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQD.....E.....TD...GERPEE...	348
GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQD.....E.....TD...GERPEE...	348
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GERIGQGRDNCREFLRENAELAREIENKVRHGLVTPMGAVTLAE.EVEED.....	352
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consensus

 non conserved
 similar
 ≥ 50% conserved