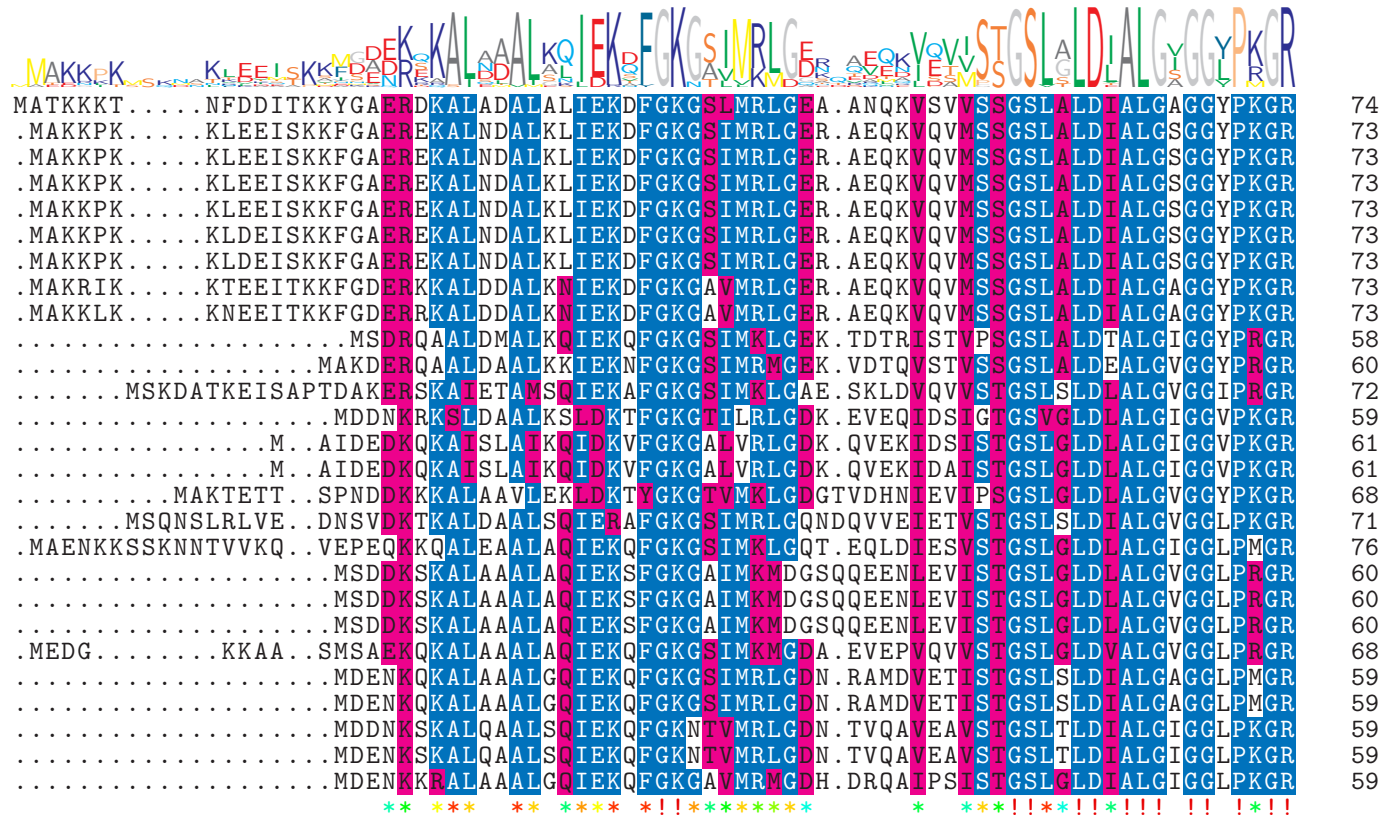


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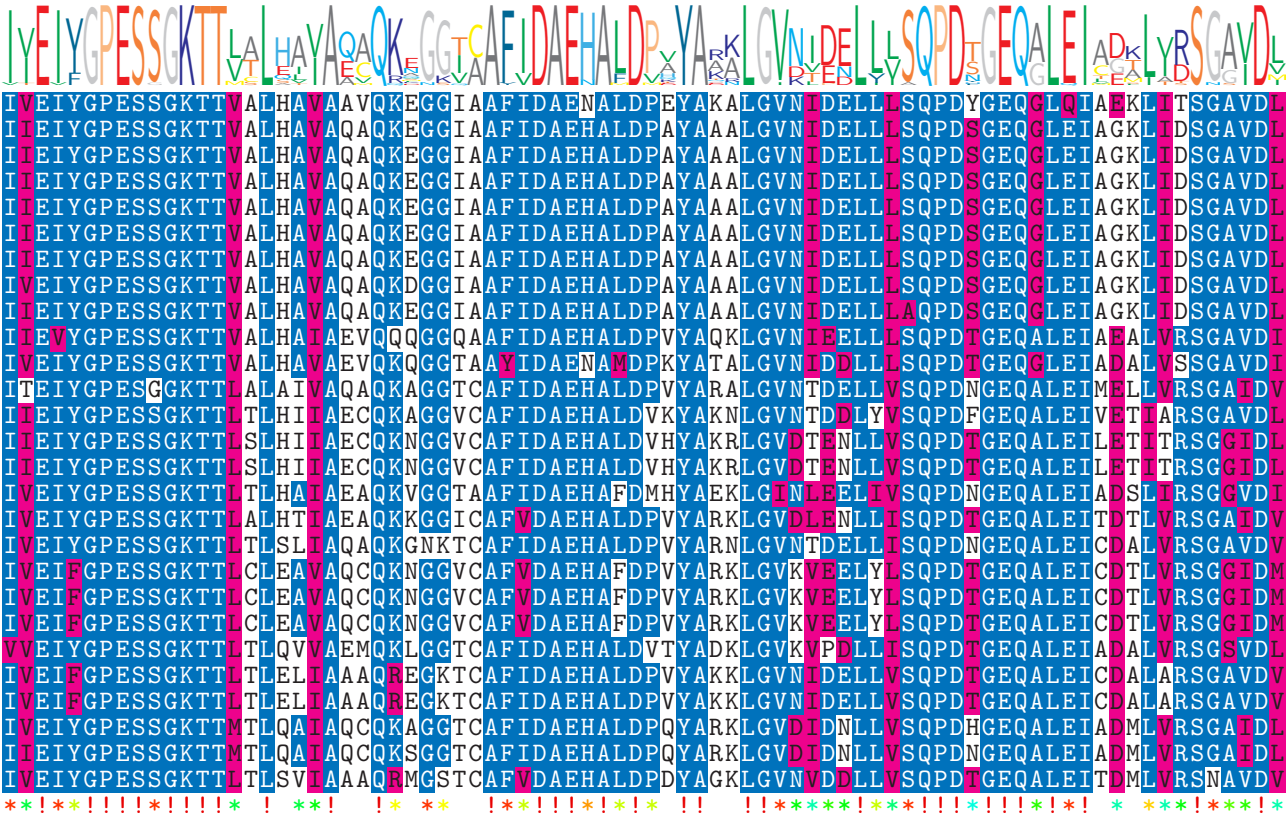
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consensus

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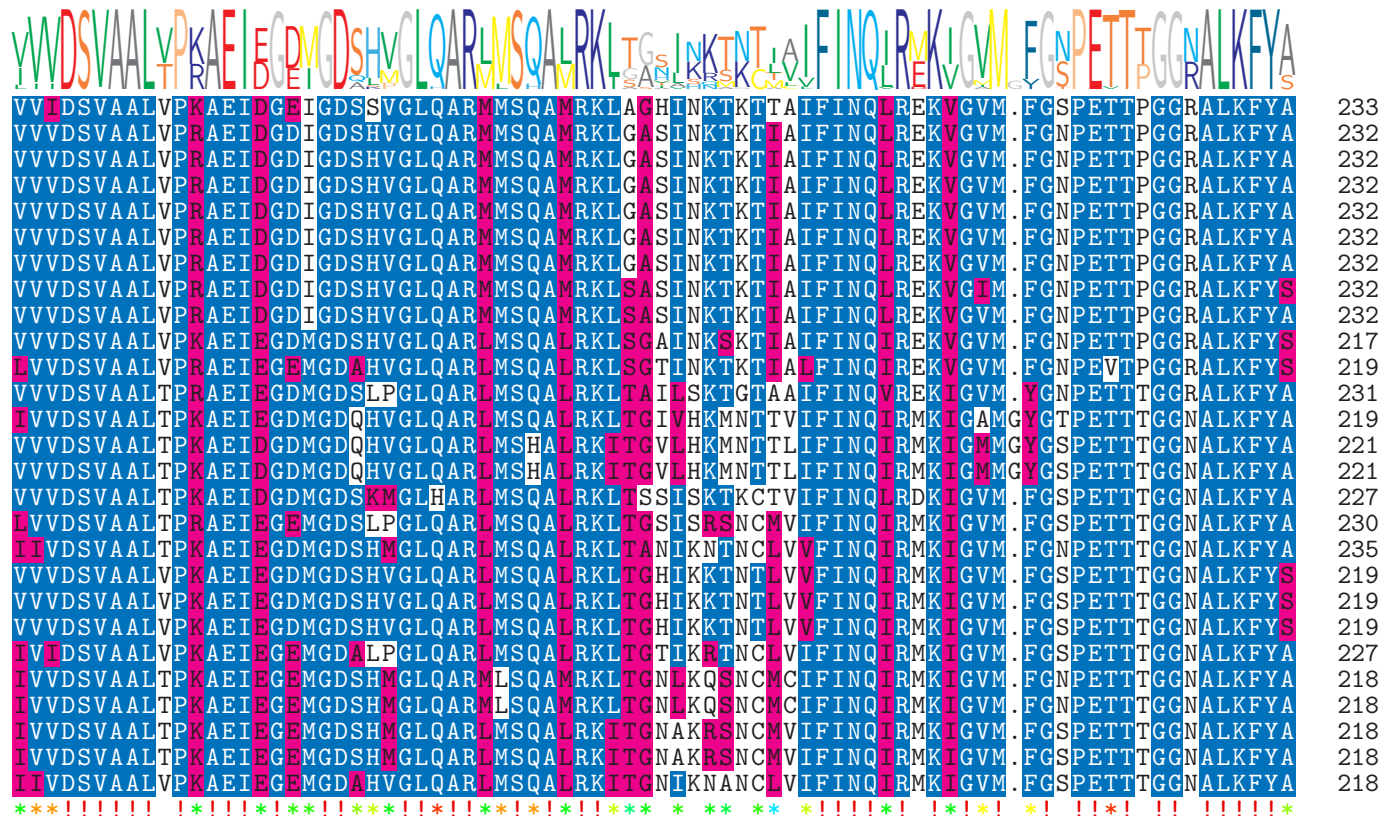


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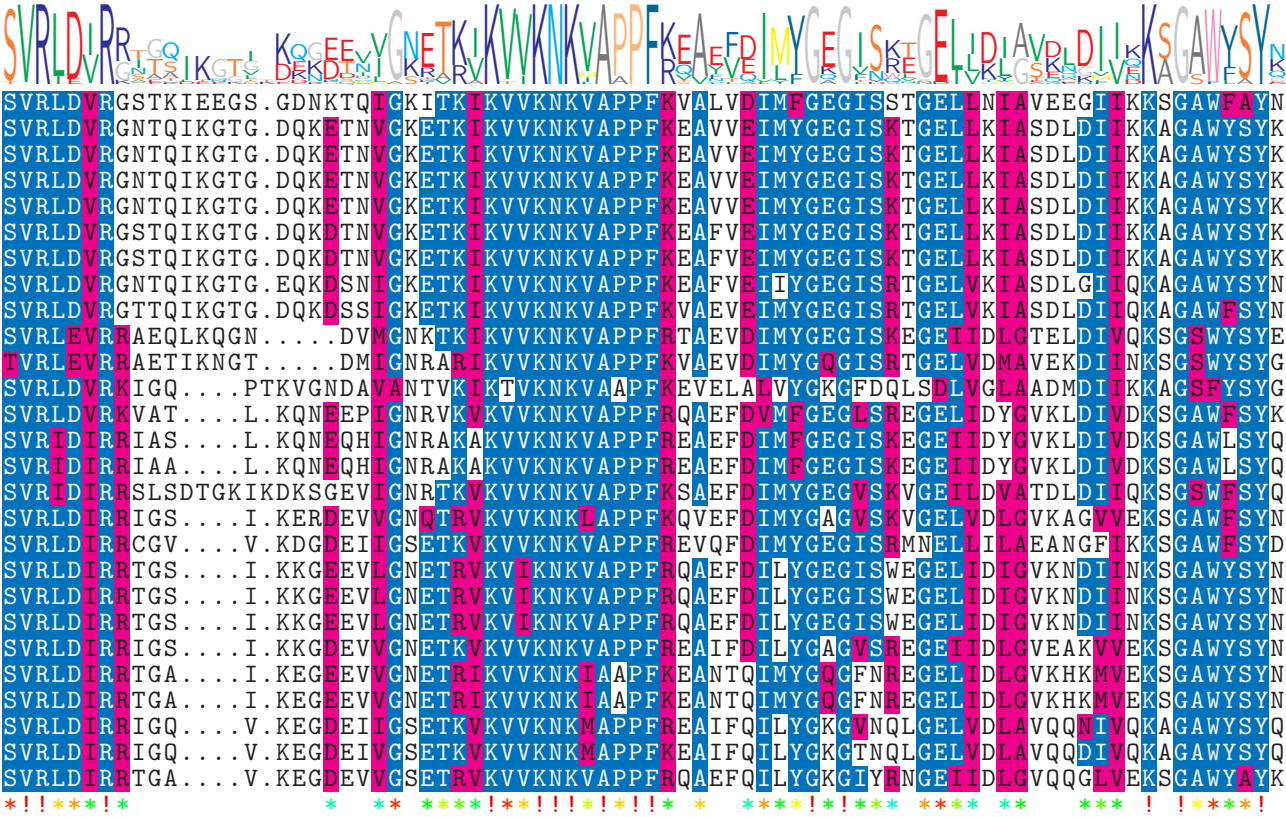
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DEKIGGQSENAAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAE...SKK.DE...AAQVDS..VNEEVPLDLGDELEI	380
GEKIGGQSENAAKFLADNPEIFDDIDHKVRSVQGLIEEDNTEEEKQSSKEK.....ETDEK..ADKNLVLELDDTTIEL	381
GEKIGGQSENAAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNNGIEI	376
EERLGGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNNGVVQQQAETQE.EL.EFE.E.....	348
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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved