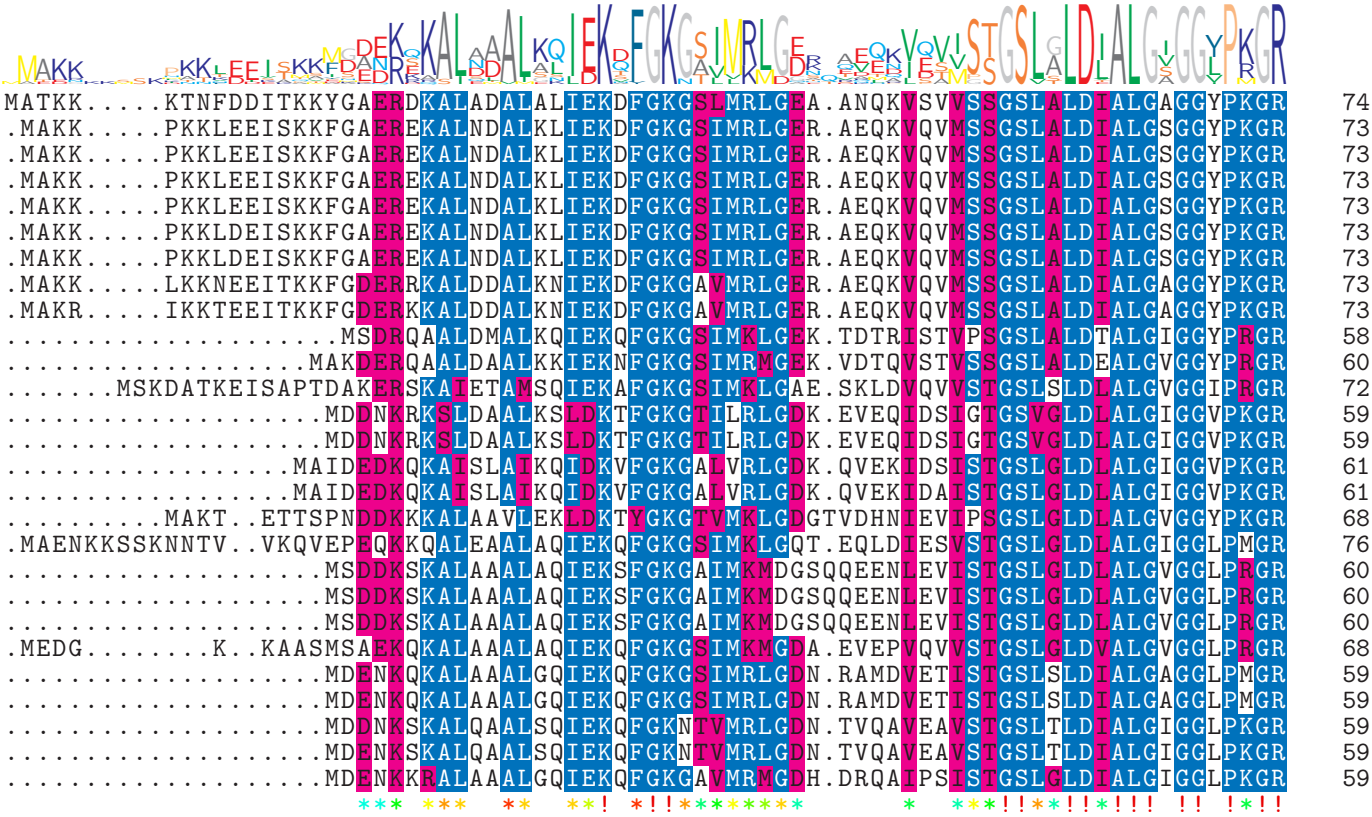


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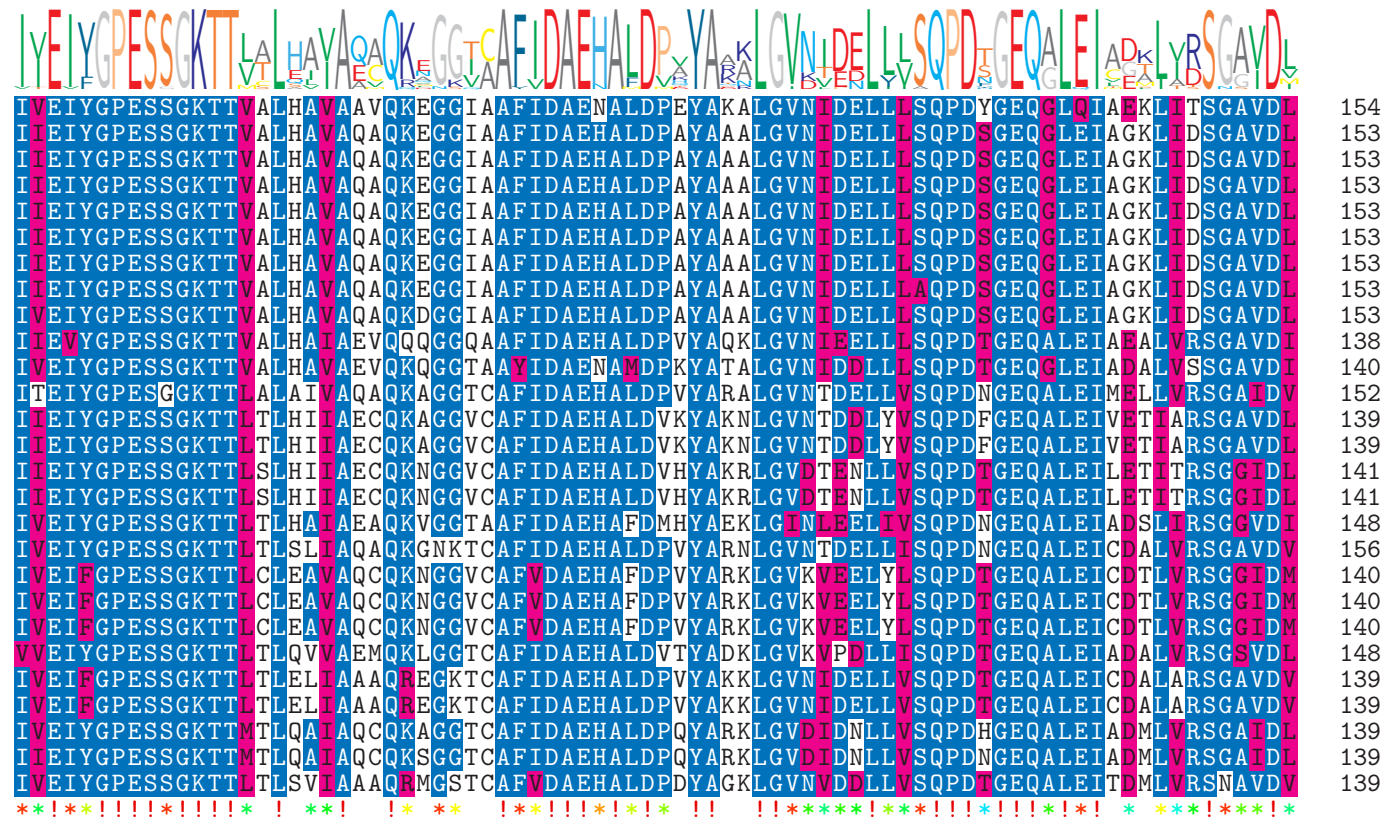
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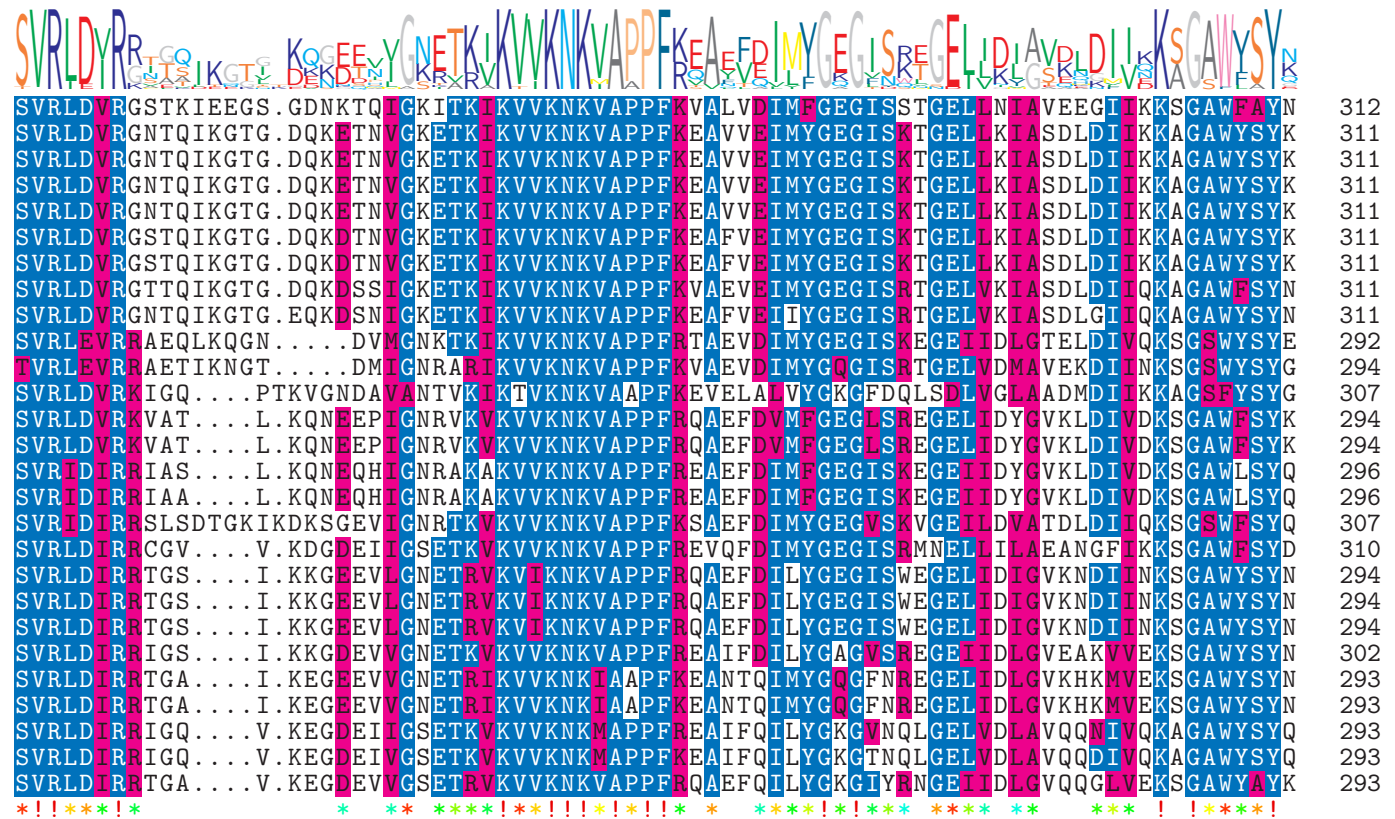
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GEKIGQGSENAKKFLADNPFIIDHQRVVFGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
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logo

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consensus

 non conserved
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 ≥ 50% conserved