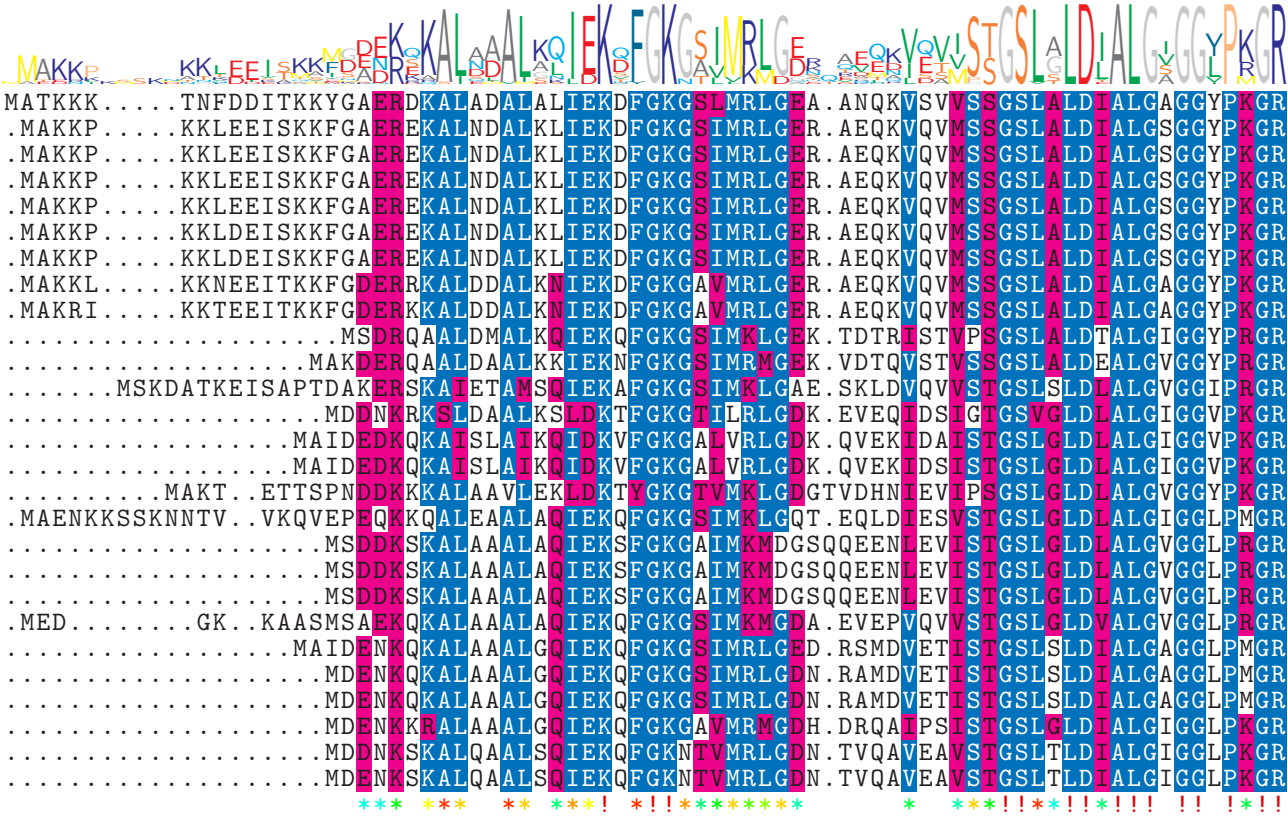


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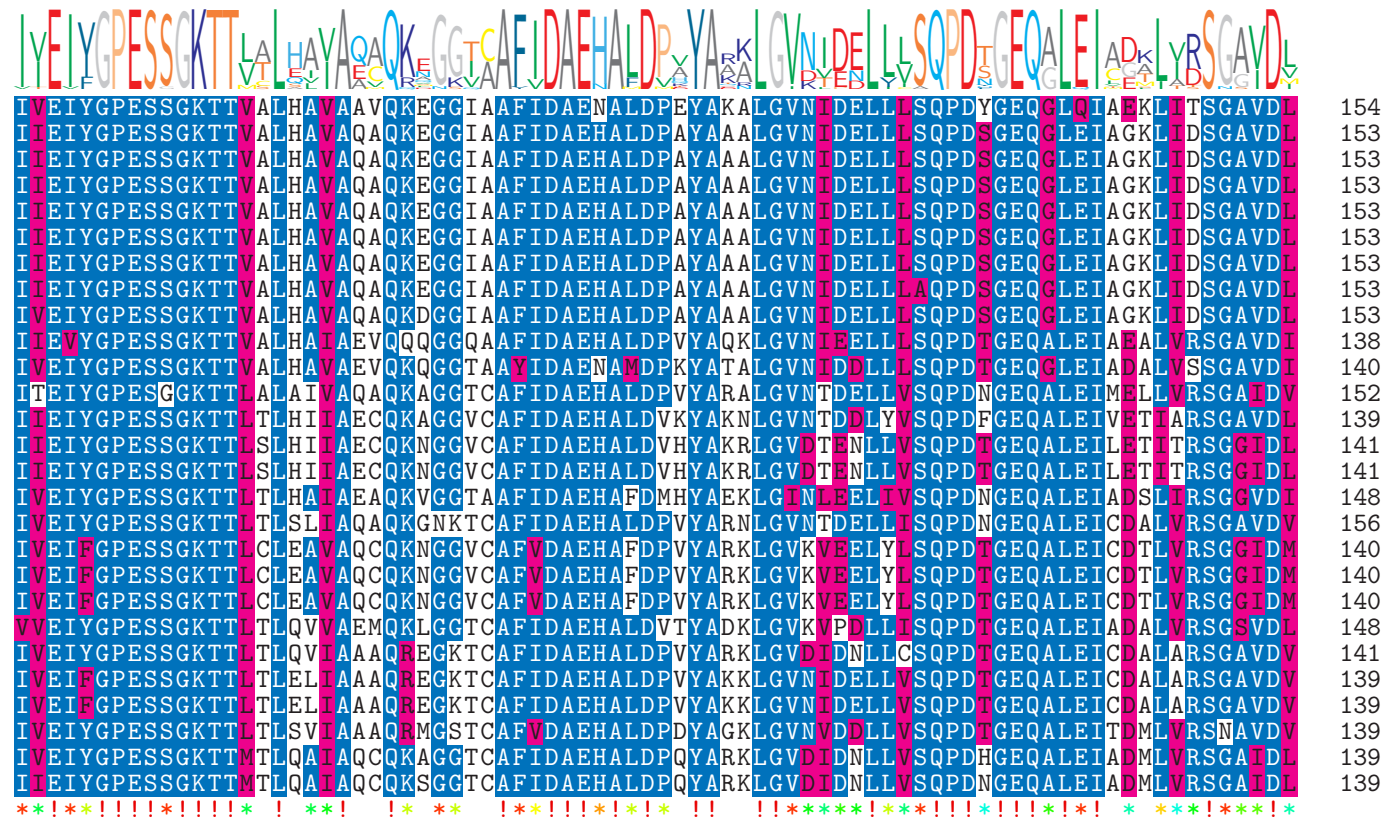


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consensus

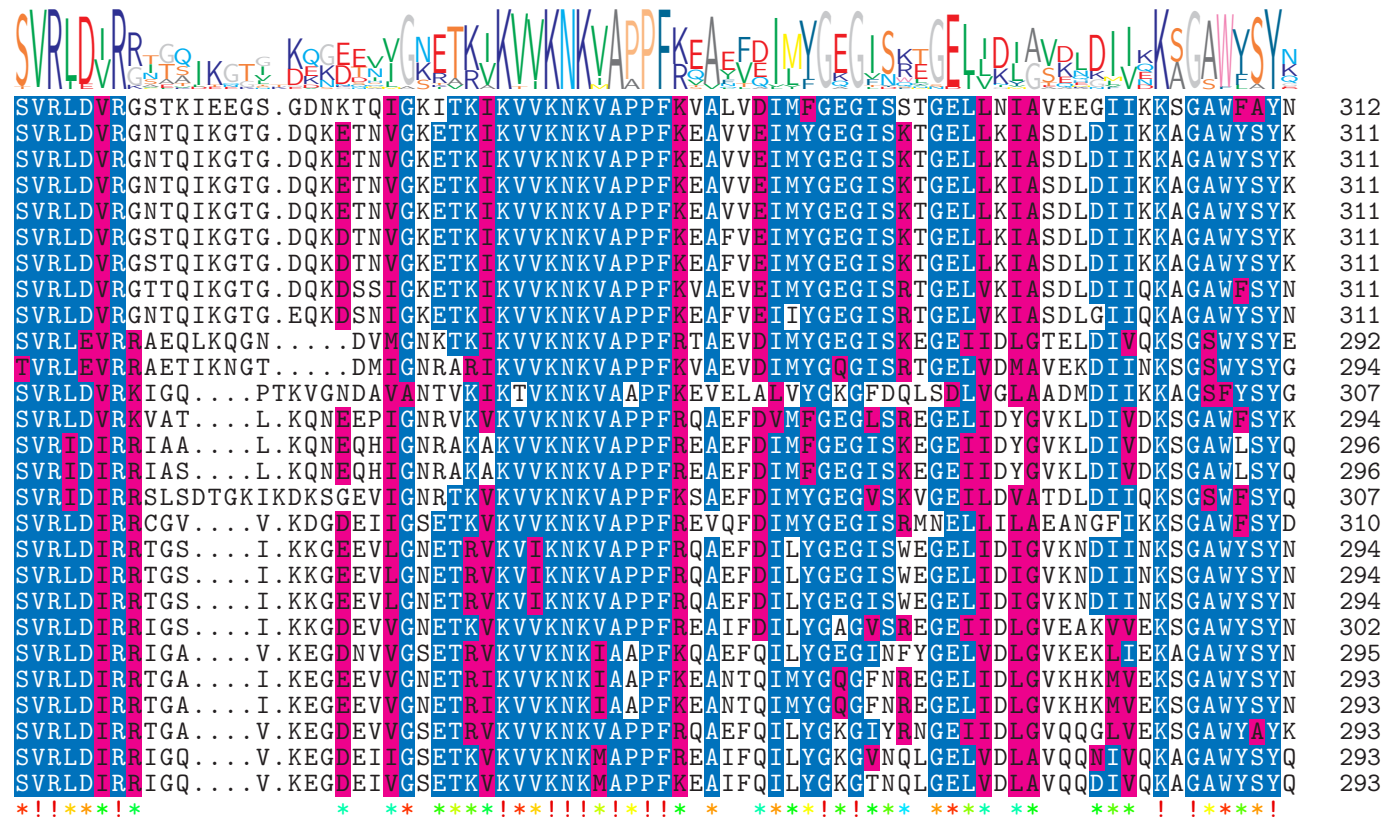
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consensus

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DEKIGQGSSENACKKYLAENPEVFAEIDHQVRVQFGLIDGEEAAES...KKDE...VAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSSENACKRYLADHPQLFDEIDRKVRVKFGLLEESEEEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSSENACKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGGGRENACKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGGGRENACKNYLADHEDVEDEVRLKVRAAAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
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GERIGGGRDNCREFLRENAELAREIENKVRHGLVTPM....GAVTLAE.EVE...ED.....	352
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GNKIGQGGKNNVIRIFYENNQIAEIEIRNIREQLLTGT....NGAVQ.I.EDE...EEPDLLES.....	349

logo

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consensus			

- X non conserved
- X similar
- X ≥ 50% conserved