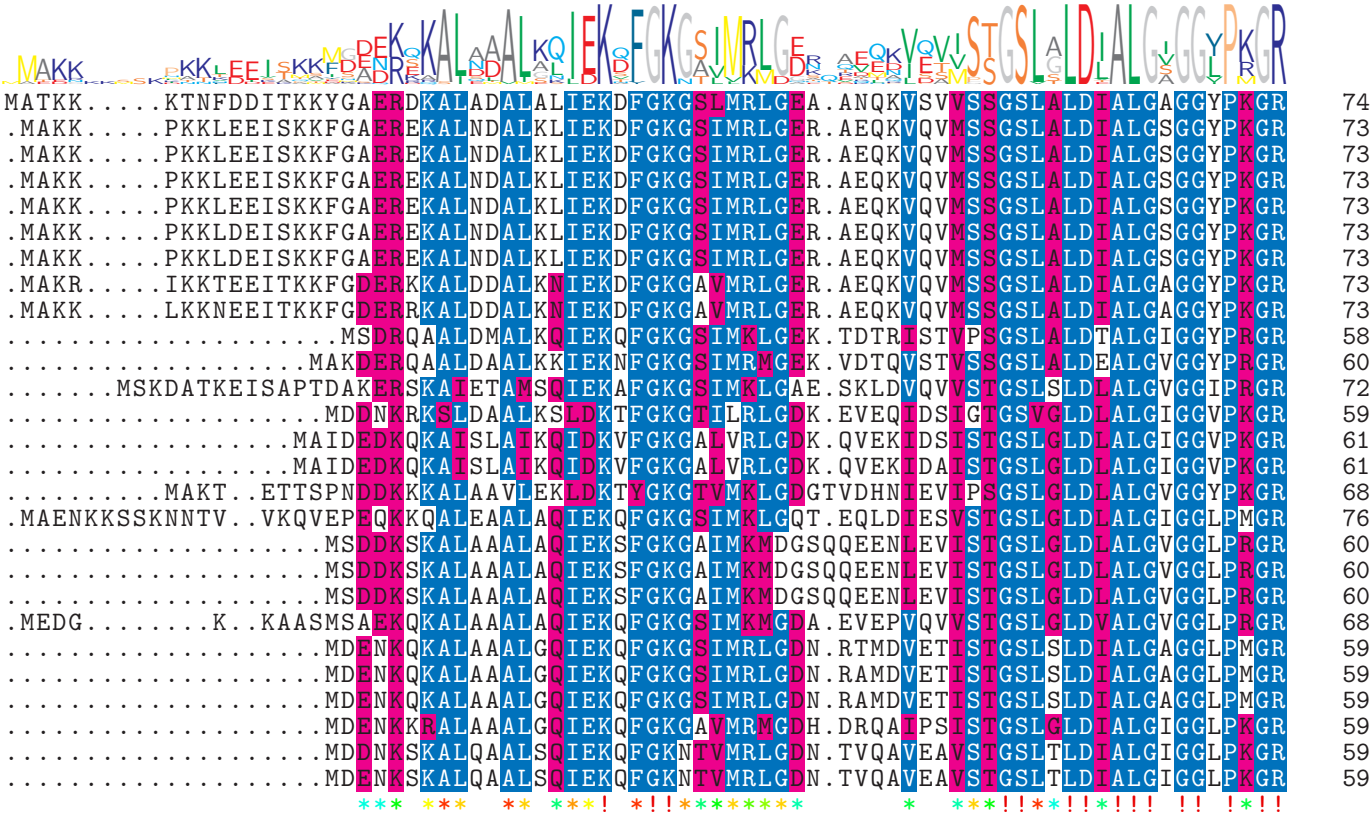


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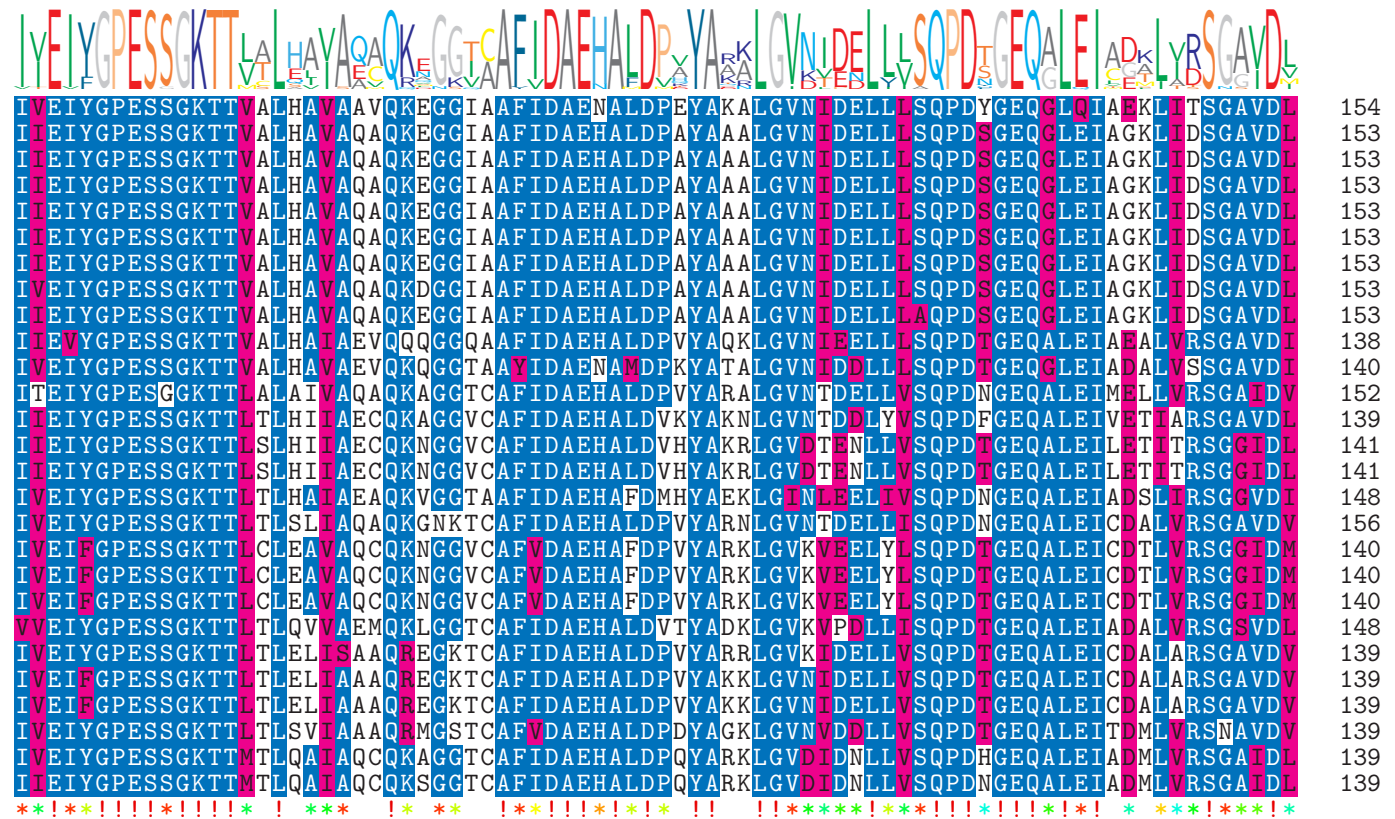
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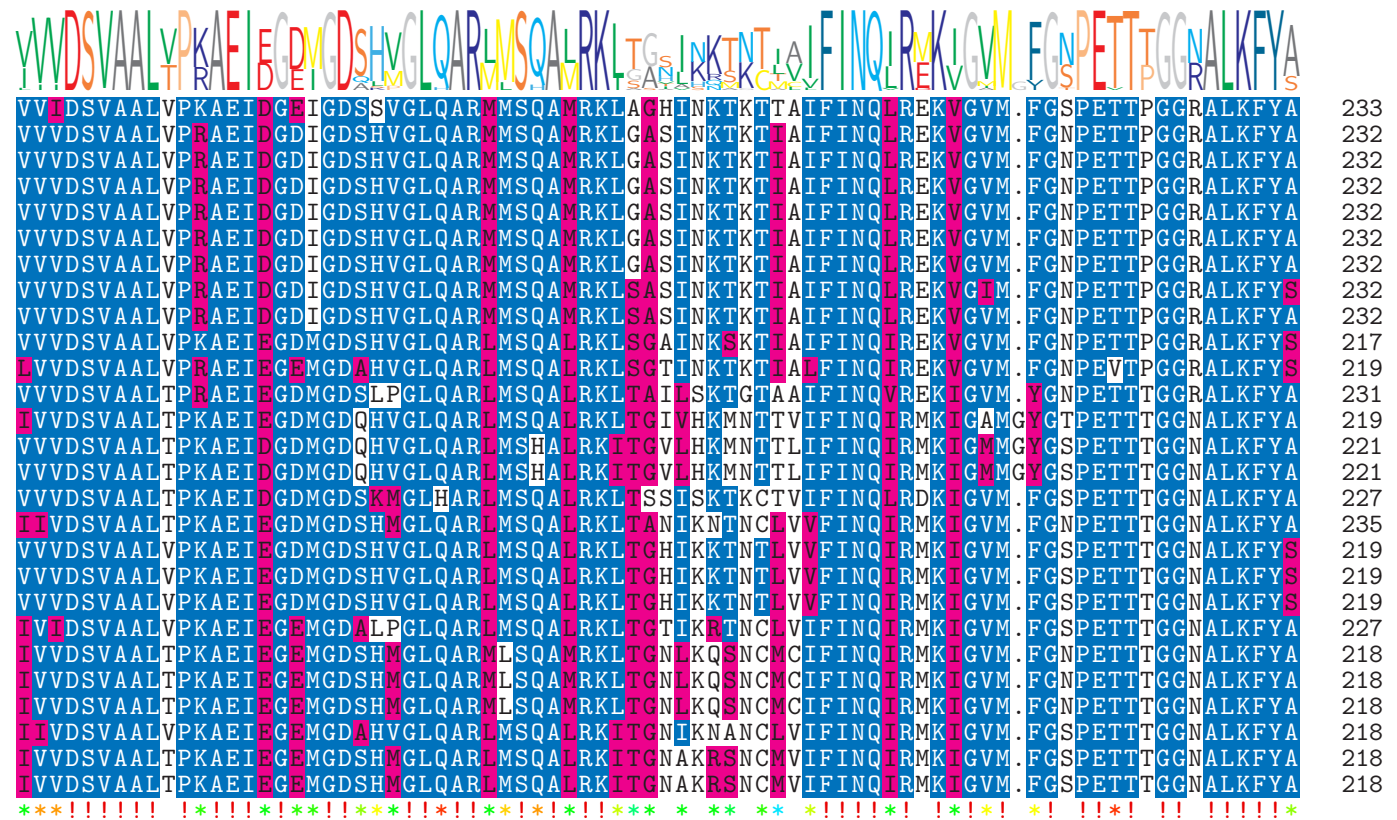
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consensus

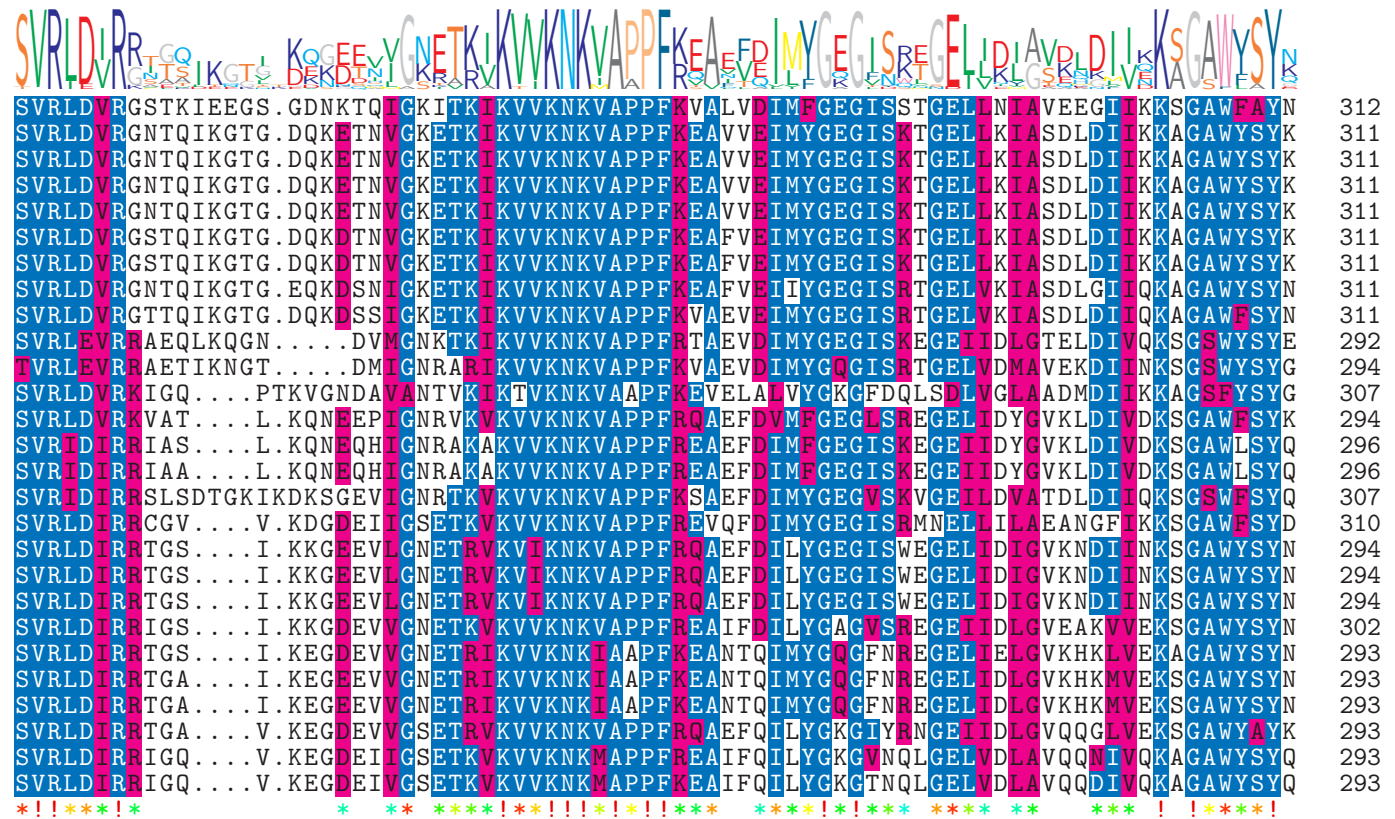
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consensus

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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQ..DTENKKD.E...PKKEEA..VNEEVPLDLGDEL	382
DEKIGQGSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAES.....KKD.E...AAQVDS..VNEEVPLDLGDEL	378
DEKIGQGSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAES.....KKD.E...VAQVDS..VNEEVPLDLGDEL	378
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQFGLIEEDNTEEK..QSSKEK.....ETDEK..ADKNLVLELDDTI	379
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEEESA..MAVASE.....E.....TDDLALDLNGI	374
EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQQ..AEETQEE.L.EFE.E.....	348
SERIGQGRENAKNYLADHEDVEDEVRLKVRVQFGLSDVPEEDLP..TTEDE.Q.I.NILPDDST.....EE.....	355
DERIGQGEKTIATYIAERPMEQETIDRVMAAIRAGNAGEAPALAPAPAPEAAEA..	363
DKKLGGGRENKSAFLKENPEIADEITKATQNSMGTGEMIS..GSEDDEG..EE.....	343
DKKLGGGRENKALLKEDKALADEITLKIKESIGSNEEIM..PLPDEPL..EEME.....	347
DKKLGGGRENKALLKEDKALADEITLKIKESIGSNEEIM..PLPDEPL..EEME.....	347
DTKLGGGRDTPVKELIKDNPELAELEAKIREKLQAK..	343
GTKIGQGGKNNAVKWLKENPEIADKLEQIRNTLASNPNSV..LKATASDSNDESDVEEFSEE.....EL.....	372
GAKIGQGGKNVRVWLKENPEISDEIDAKIRALNGVEMHIT..EGTQDET.....DGERPEE.....	348
GAKIGQGGKNVRVWLKENPEISDEIDAKIRALNGVEMHIT..EGTQDET.....DGERPEE.....	348
GAKIGQGGKNVRVWLKENPEISDEIDAKIRALNGVEMHIT..EGTQDET.....DGERPEE.....	348
GERIGQGGKRDNCREFLRENAELAREIENKVRHGLVTPMGA..VTL.AEEVEED.....	352
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GSKIGQGGKANACKYLEENPEIGQIEEQQIRAKLLVTSANT..KATPVAEDLADADI.....	347
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GNKIGQGGKNNVIRIYFENNQIAEIERNIREQLLTTGTNG..AVQ.I.EDEEEPDLLES.....	349

logo

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consensus								

-  non conserved
-  similar
-  ≥ 50% conserved