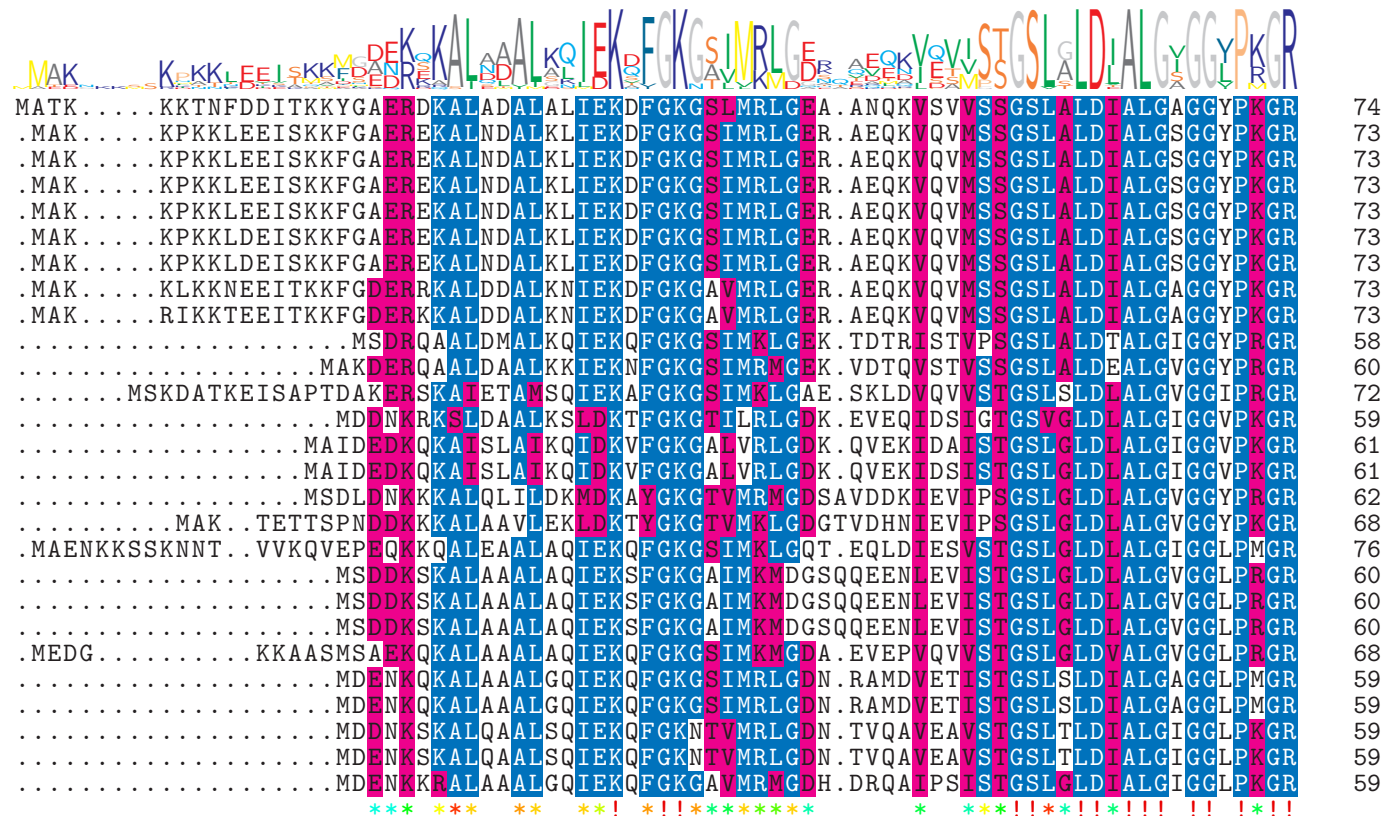


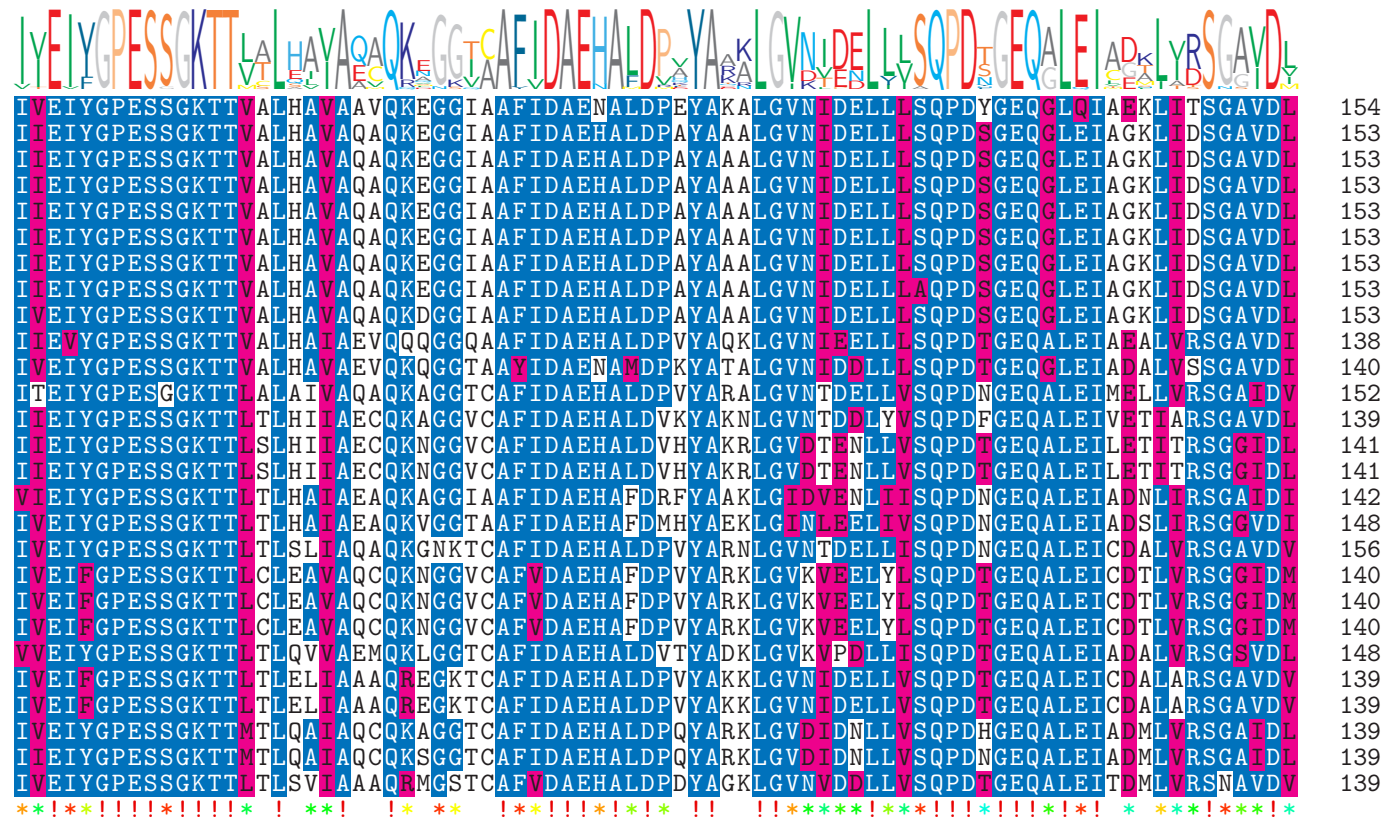
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consensus



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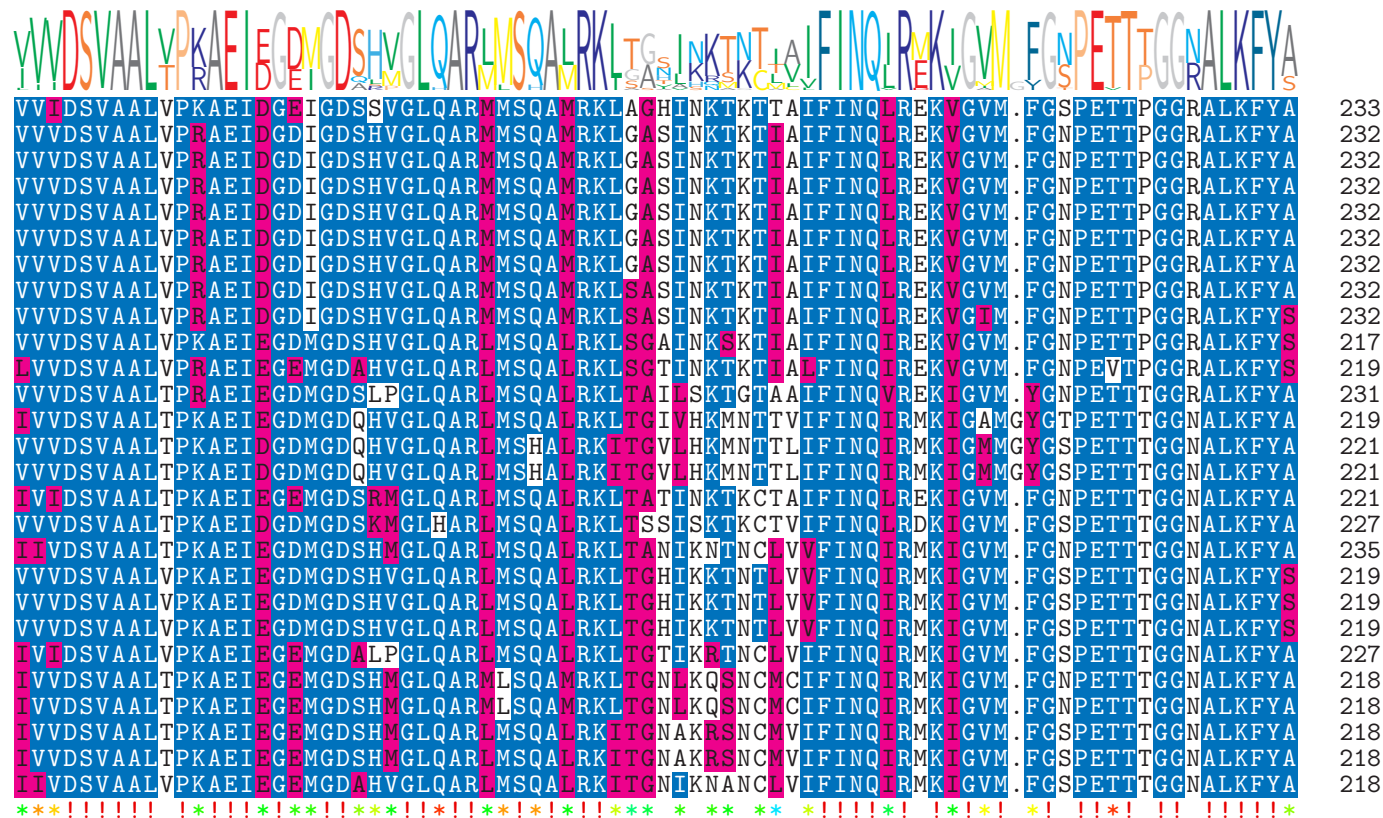
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consensus

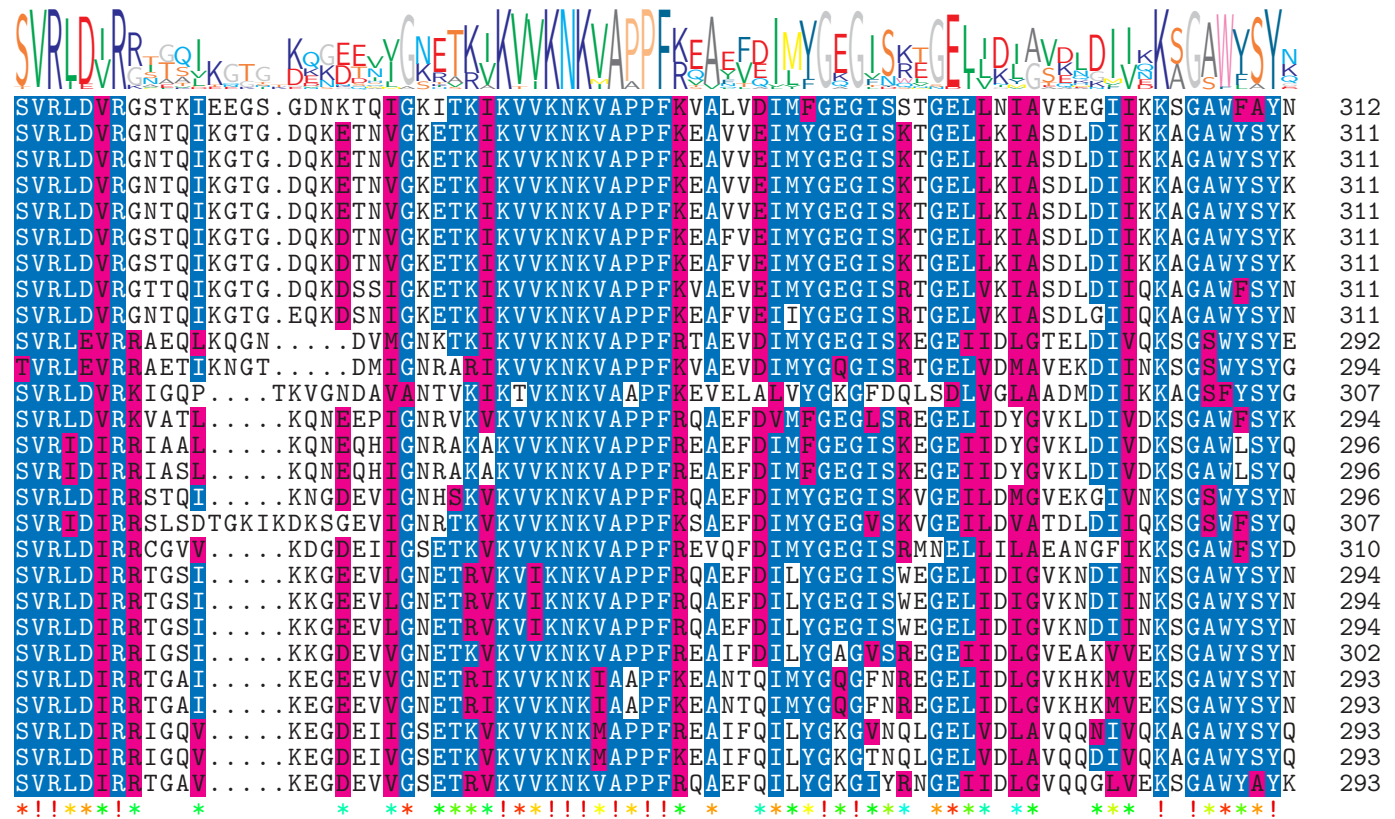
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consensus

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consensus

Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK...KDE...PKKEEA..VNEEVPLDLGDE	381
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DEKIGQGSENAKKYLAHNPEVFAEIDHQRVVFGLIDGEEAAES...K...KDE...VAQVDS..VNEEVPLDLGDE	377
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVAS...E.....E.....TDDLALDLNG	373
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKE...K.....ETDEK..ADKNLVLELDDT	378
EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQ...EEL.EFE.E.....	348
SERIGQGRENAKNYLADHEDVEDEVRLKVRAAAYGISDVPEEDLPTTEDE...QI.NILPDDST...EE...	355
DERIGQGEKTIATYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAEA.....	363
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DKKLGGGRENKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP..LEEME	347
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GNKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTKASD...QTAA.HDETEEEPDLL..ES	349
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logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved