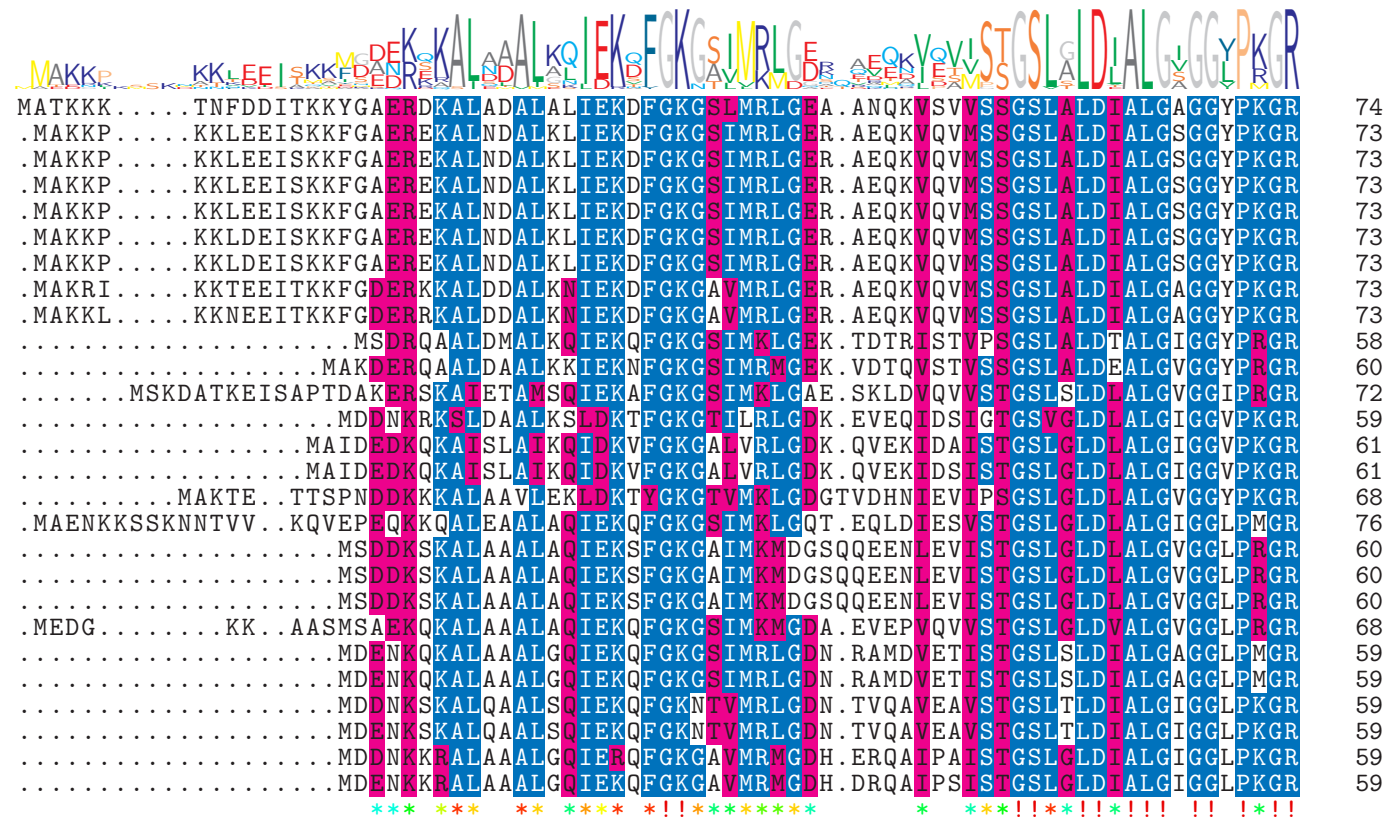


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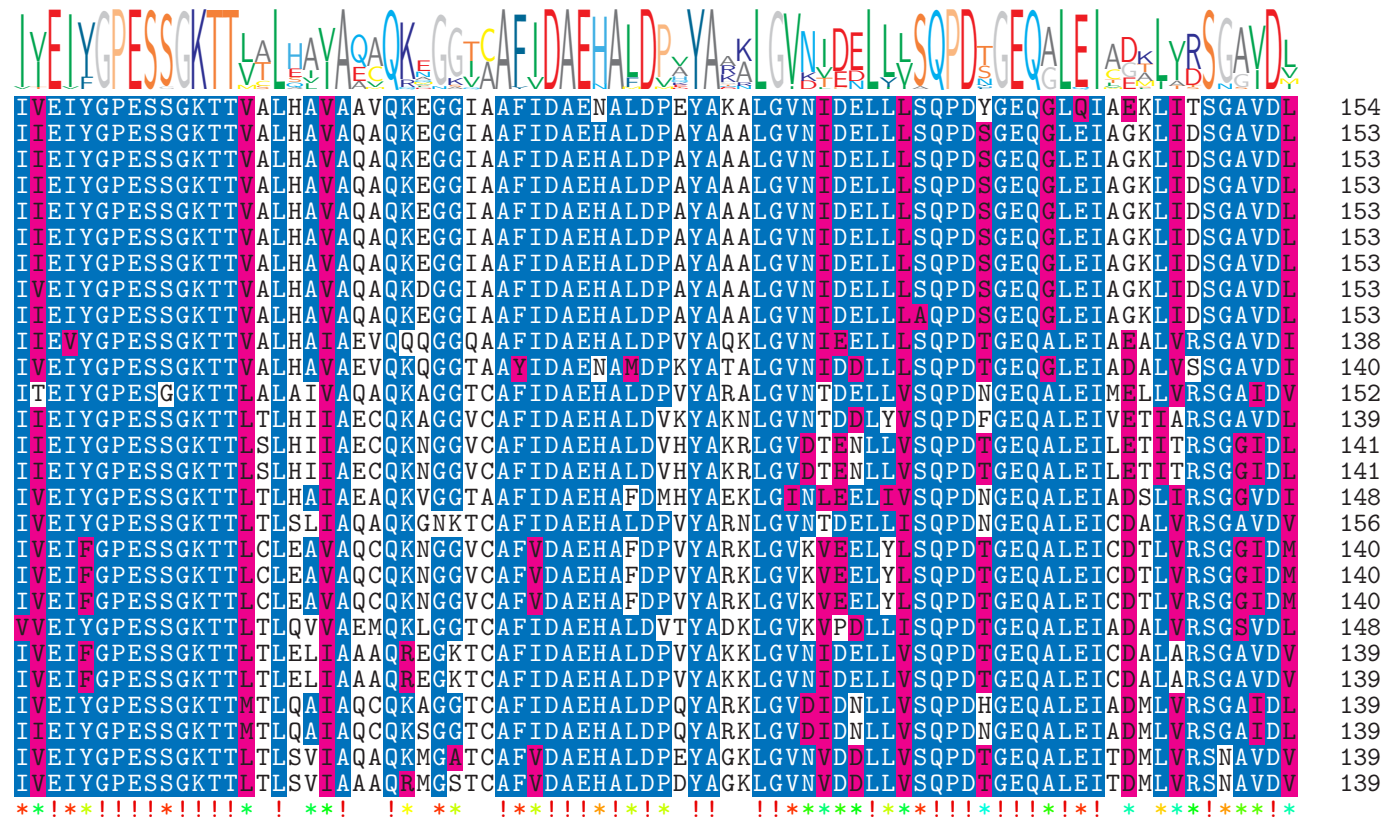
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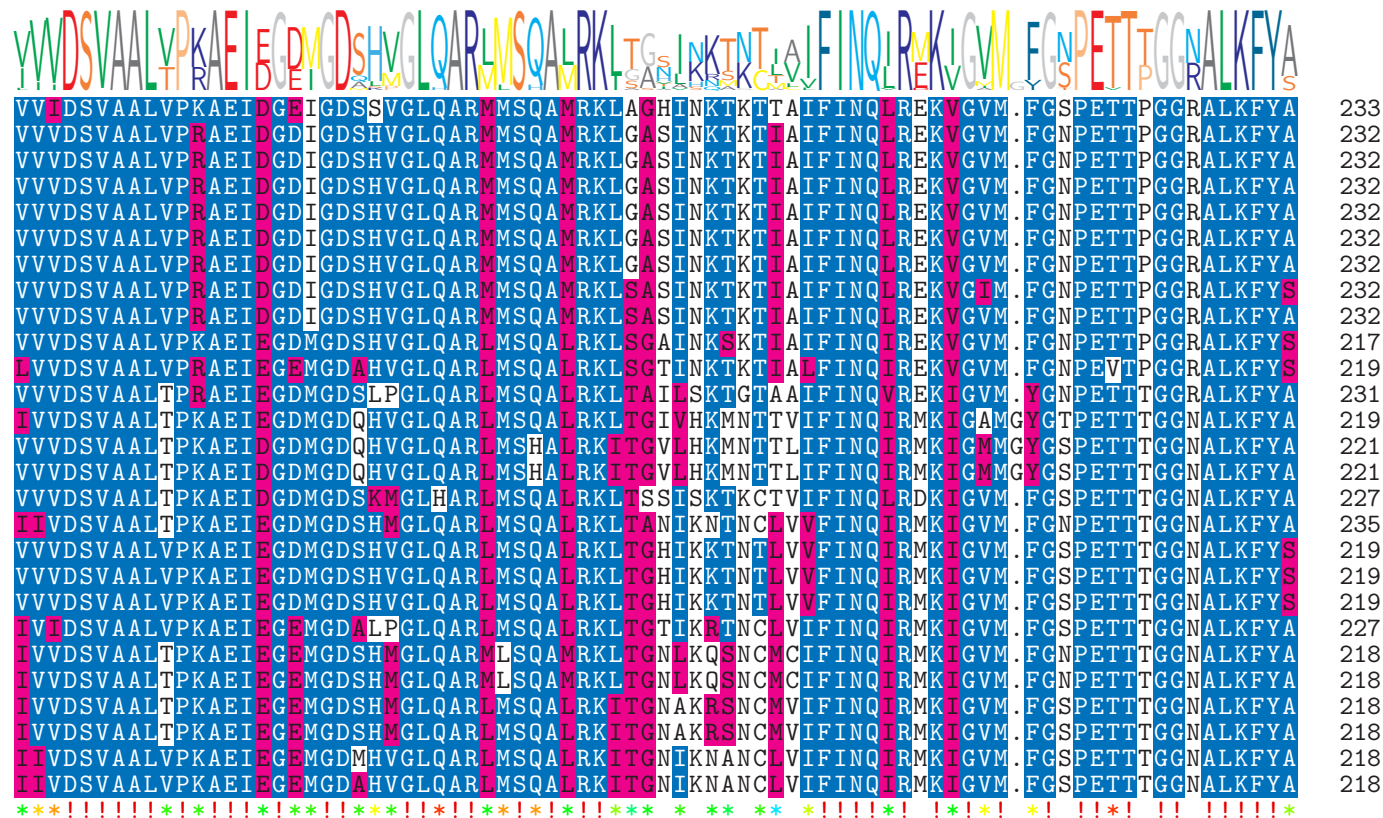
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consensus

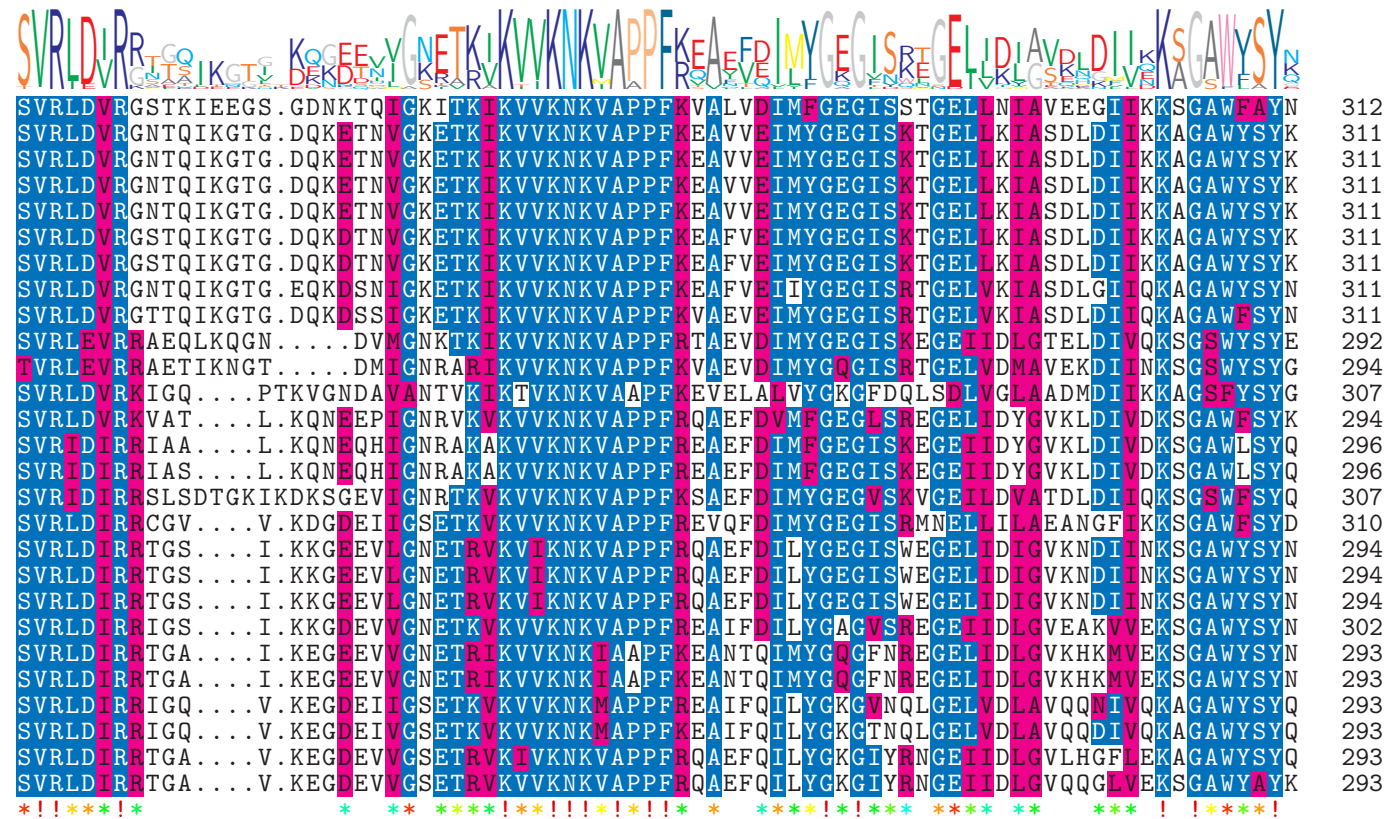
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NTDB id 314948 D3Z90 RS20775 WP 136477796.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKD.E...PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSENAKKYLAHNPEVFAEIDHQVRVQFGLIDGEEAAE...SKKD.E...VAQVDS..VNEEVTLDLGDELEI	380
DEKIGQGSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAE...SKKD.E...AAQVDS..VNEEVTLDLGDELEI	380
GEKIGQGSENAKFLADNPEIFDDIDHKVRVQFGLIEEDNTEEEKQSSKEK.....ETDEK..ADKNLVLELDDTIEL	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEEESAMAVASE.....E.....TDDLALDLNGIEI	376
EERLGQGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEE.L.EFE.E.....	348
SERIGQGGRENAKNYLADHEDVEDEVRLKVRVQFGLSDVPEEDLPTTEDE.Q.I.NILPDDST.....EE.....	355
DERIGQGKEKTIAYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAP.....AAPEAAEA.....	363
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GTKIGQGKNNNAVKWLKENPEIADKLEQIRNTLASNPNSVLKATASDSNDESDV...EEFSE...EEL.....	372
GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVMHITEGTQD.....ETDGERPEE.....	348
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GNKIGGQKNNVIRYFEENNQIAEIEIERNIREQLLTGTNGAVQI.....EDEEEPDLLE.....S.....	349
GSKIGGQKANSAKFLQDNPEIAATLEKQIRDKLLNPGAVEAAGKAA.....AVNAEPEDEAE.....TEV.....G.....	354
GSKIGGQKANAAYLEENPEIQEIEQQIRAKLLVTSANTKATPVA.....E.....DLAD.....ADI.....	347

logo

NTDB id 600 KW2 RS01765 WP 011675413.1		387
NTDB id 269 KZH43 RS08665 WP 001085462.1	EIEE	388
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NTDB id 1205 PSJM300 05545 AFN77183.1		347

consensus

 non conserved
 similar
 ≥ 50% conserved