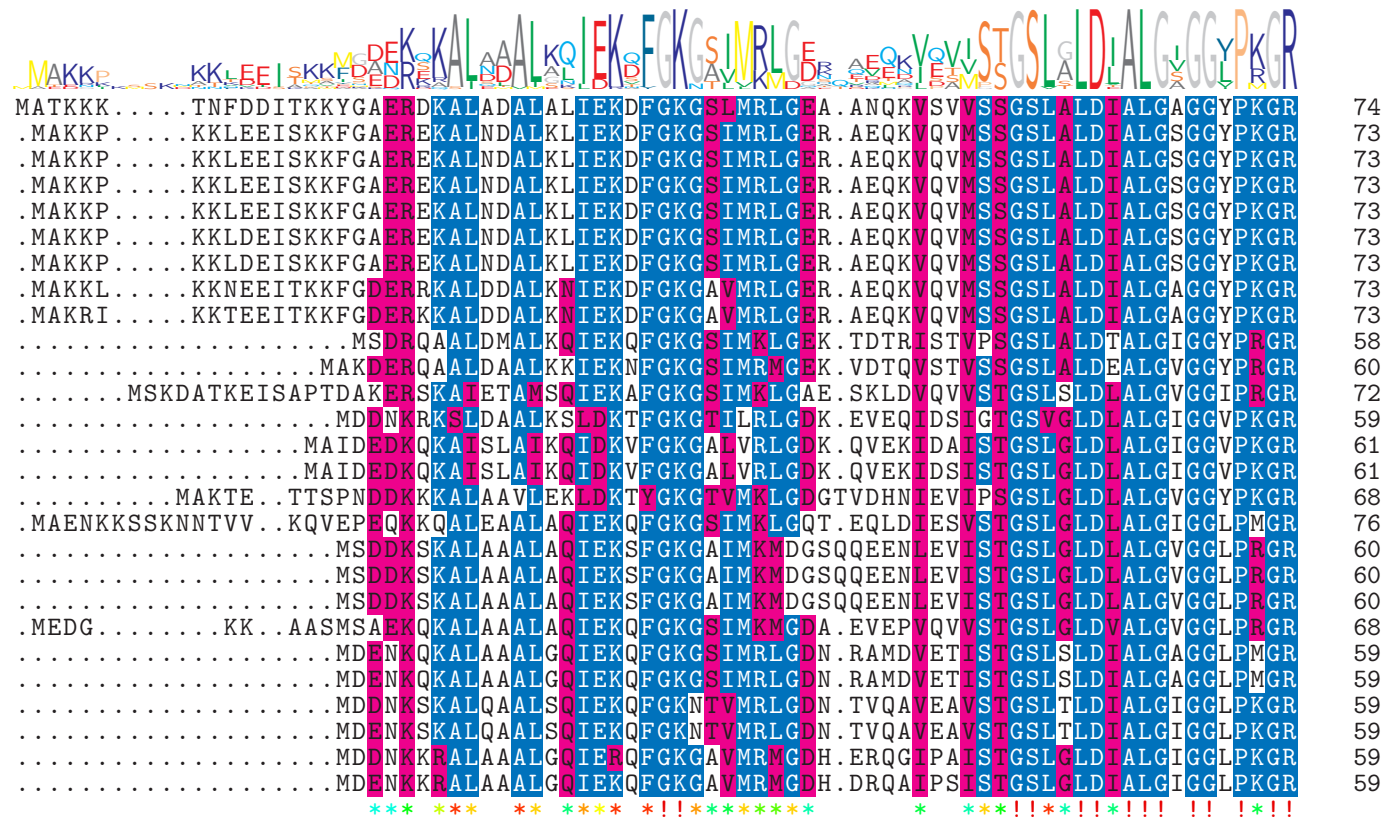


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consensus

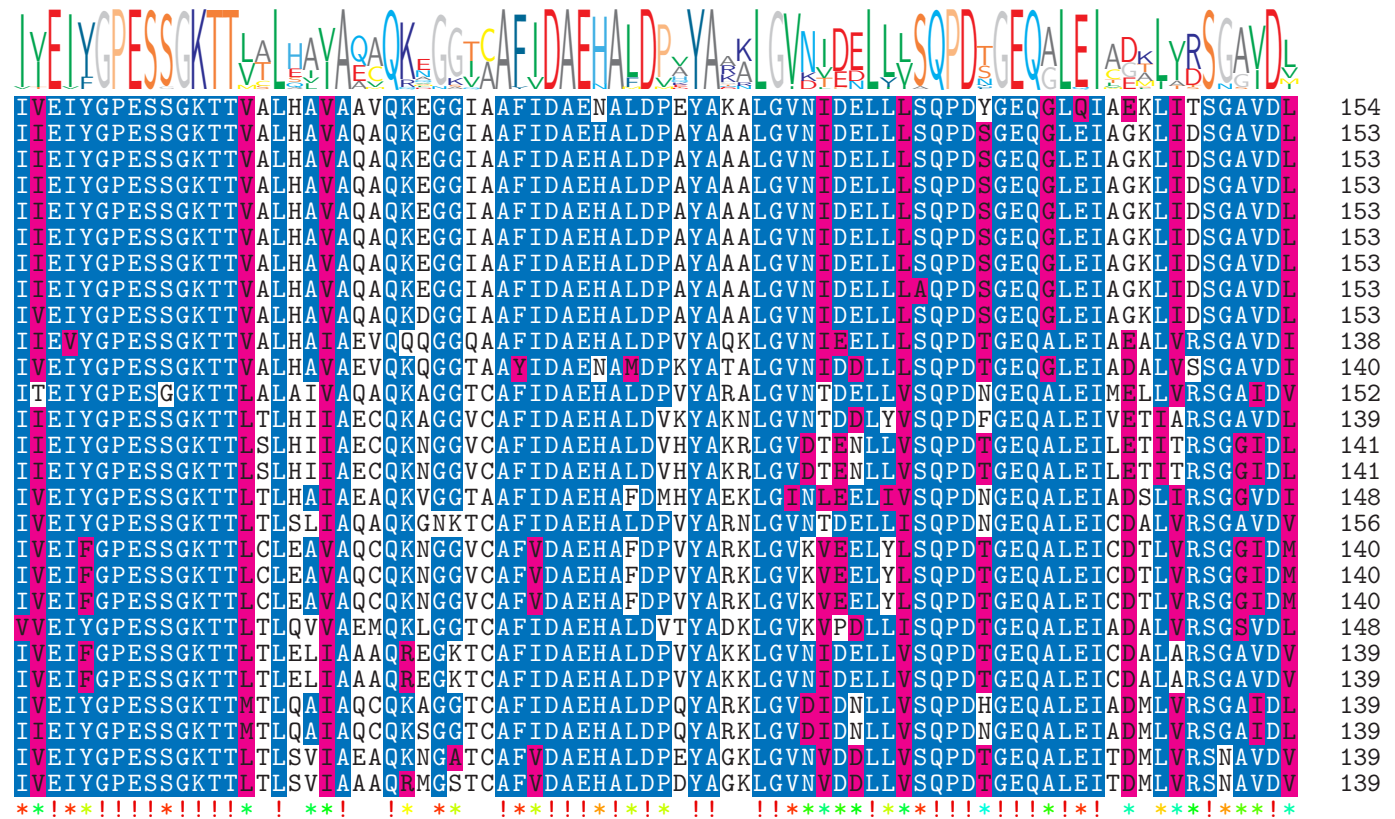
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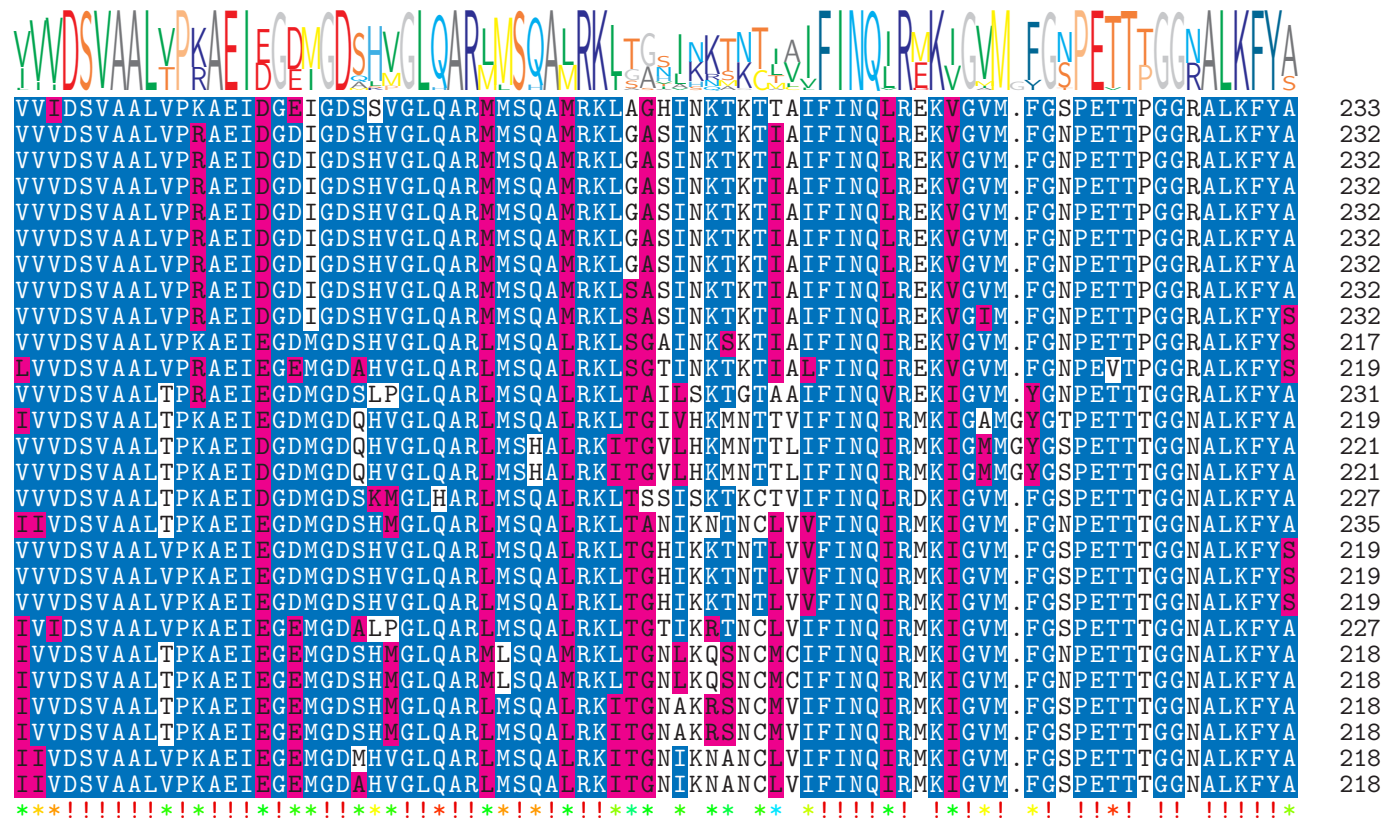
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

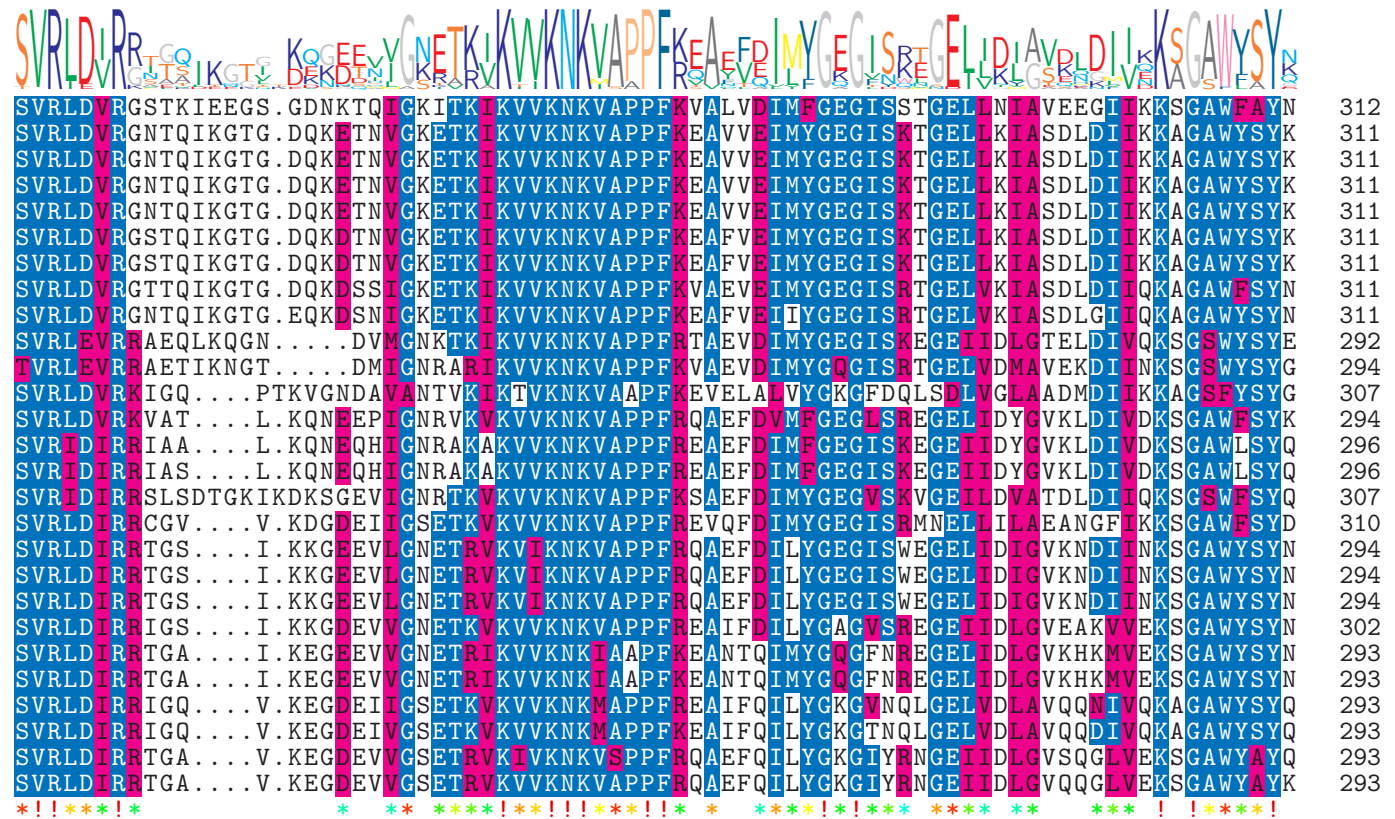
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consensus

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consensus

Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLAADNPEVFAEIDHQVRVQFGLIDGEEAAE...SKKDE...VAQVDS..VNEEVTLDLGDLEIE	381
DEKIGQGSENAKKYLAADNPEIFDEIDHQVRVQFGLIDGEEIAE...SKKDE...AAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGQGGRENAKKNYLADHEDVEDEVRLKVRAAYGISDVPPEEDLPTTEDE.QI.NILPDDST....EE.....	355
DERIGGQKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAP...A...APE..A...AEA.....	363
DKKILGQGGRENSKAFILKENPEIADEITKATQNSMGTIGMISGSEDDEGE.....E.....	343
DKKILGQGGRENAKALLKEDKALADEITLKIKESIGSNEEIMPLPDEPLE.....E.....ME.....	347
DKKILGQGGRENAKALLKEDKALADEITLKIKESIGSNEEIMPLPDEPLE.....E.....ME.....	347
DTKILGQGRDTPVKELIKDNPELAELEAKIREKLQAK.....	343
GTKIGQGKNNNAVKWLKENPEIADKLEQEIRNTLASNPNSVLKATASDSNDES...DVE..E...FSEEL.....	372
GAKIGQKDNVRVWLKENPEISDEIDAKIRALNGVMHITEGTQDET.D..GE...RPE..E.....	348
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GAKIGQKDNVRVWLKENPEISDEIDAKIRALNGVMHITEGTQDET.D..GE...RPE..E.....	348
GERIGQGRDNCREFLRENAELAREIENKVRHLGVTPM.GAVTLAEV..EE...D.....	352
GDKIGQGGKANACKYLKENPEIAKTLKKREMLLNPNMQLIAETSSA..AD..DVE.....FGAVPEEF.....	354
GDKIGQGGKANACKYLKENPEIAKTLKKREMLLNPNMQLIAETSSA..AD..DVE.....FGAVPEEF.....	354
GNKIGQGGKNNVIRYLEENPQISTEIEAVIREQLLTKASDQTAHDETE...E...EPD..L...LES.....	349
GNKIGQGGKNNVIRYFEENNQIAEIEIERNIREQLLTGTNGAVQI.EDE...E...EPD..L...LLES.....	349
GNKIGQGGKANAAKYLAENPAIGAIEEKQIREKLLKAGAAAEAGKAAAA..EA...DAD..D...MADADAGY.....	355
GSKIGQGGKANAAKYLEENPEIGQIEIEQRIAKLLVTSANTKATPVAE.....D...LADADI.....	347

logo

NTDB id 600 KW2 RS01765 WP 011675413.1	...	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	IEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	IEE	388
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NTDB id 1205 PSJM300 05545 AFN77183.1	...	347

consensus

 non conserved
 similar
 ≥ 50% conserved