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NTDB id 1427 RS RS02760 WP 011000510.1
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NTDB id 1205 PSJM300 05545 AFN77183.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

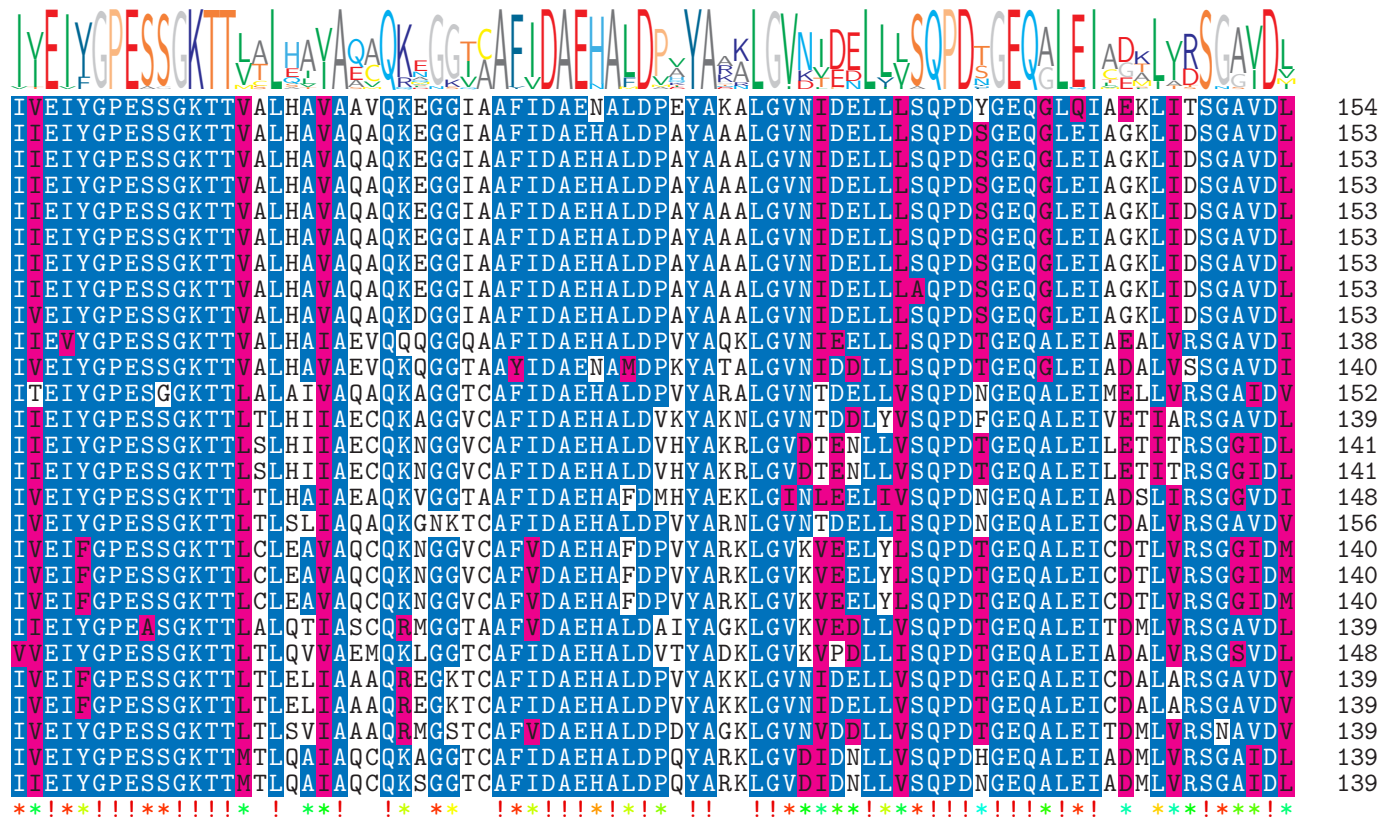
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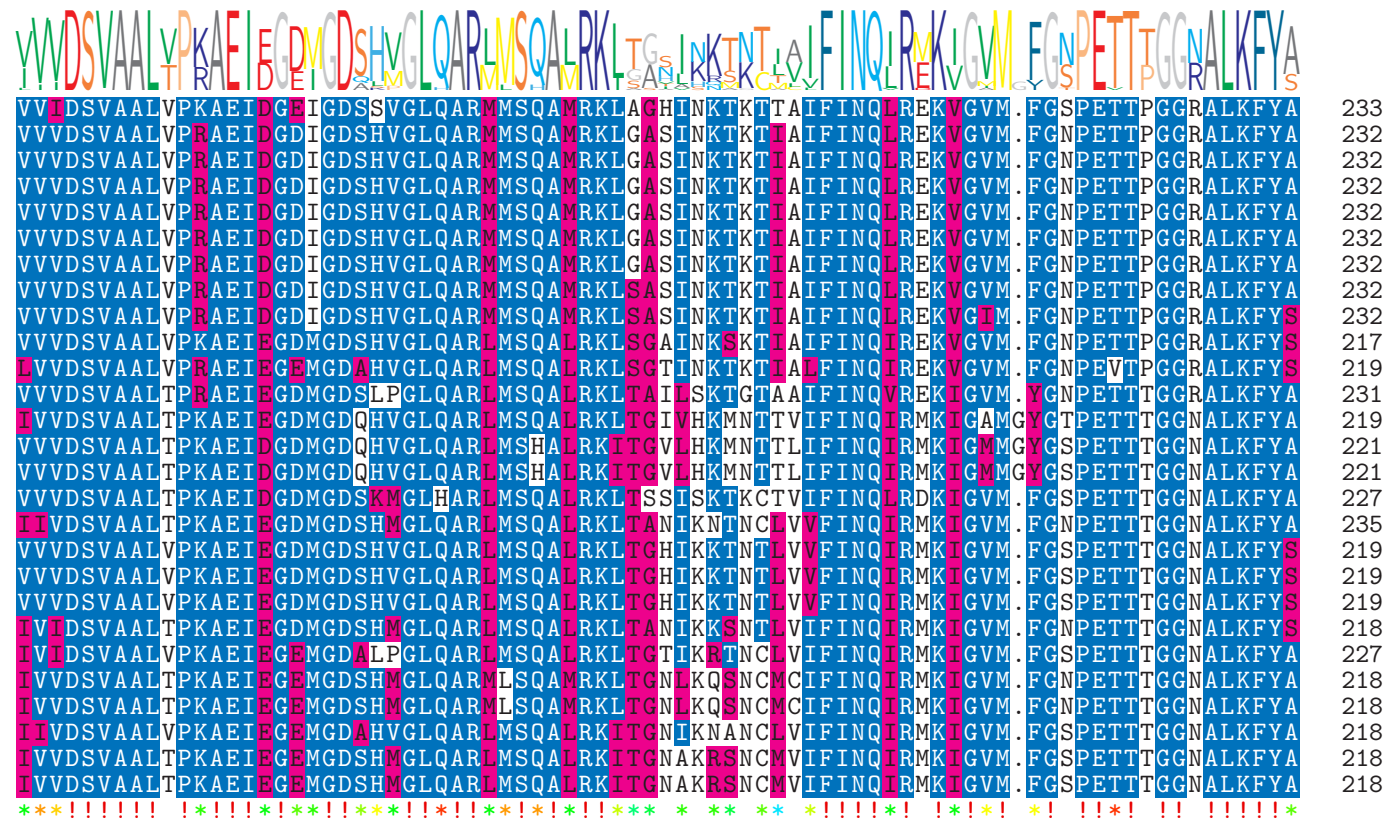
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consensus

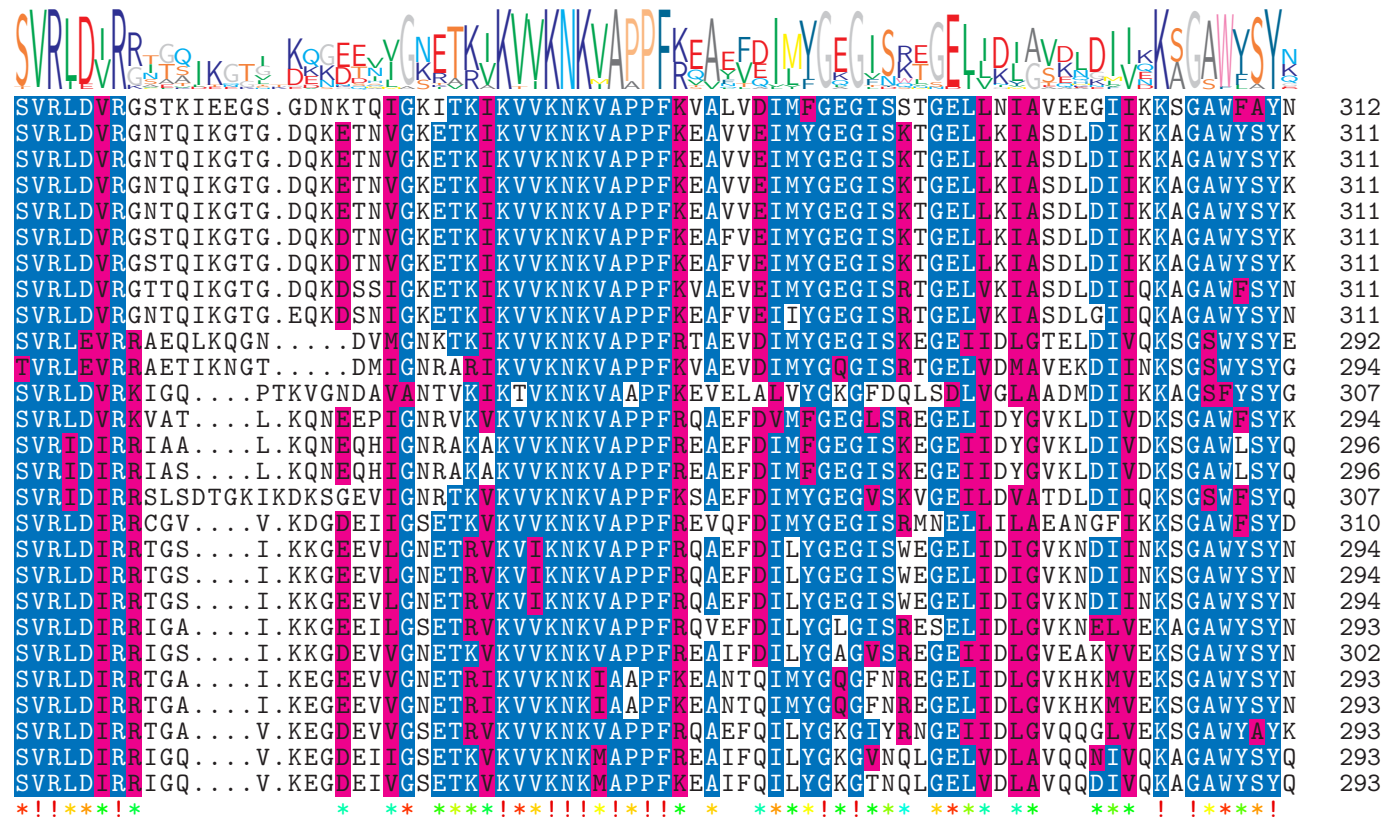
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK.KDE...PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK.KDE...PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK.KDE...PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSENAKKYLAADNPEIFDEIDHQRVQVFLIDGEEIAES...K.KDE...AAQVDS..VNEEVPLDLGDELEI	380
DEKIGQGSENAKKYLAADNPEVFAEIDHQRVQVFLIDGEEIAES...K.KDE...VAQVDS..VNEEVPLDLGDELEI	380
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVAS.E.....E.....TDDLALDLNGIEI	376
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQVGLIEEDNTEEKQSSKE.K.....ETDEK..ADKNLVLEDDTIEL	381
EERLGGGRENAAKQFLKENKDIMLMIQEQUIREHYGLDNGVGVQQAETQ.EEL.EFE.E.....	348
SERIGGGRENAAKNYLADHEDVEDEVRLKVRAAAYGISDVPEEDLPTTEDE..QI.NILPDDST.....EE.....	355
DERIGGGKEKTIAYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPAAEA.....	363
DKKLGGRGRENKAFLEKENPEIAEITTKAIQNSMGTIEGMISGSE..DDE..GEE.....	343
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DKKLGGRGRENAAKALLKEDKALADEITLKIKESIGSNEEIMPLP..DEP..LEEME.....	347
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GTKIGQGGKNNAVKWLKENPEIADKLEQAIRNTLASNPNSVLKA..TASDSNDES DVEEF.....SEEL.....	372
GAKIGQGGKNVRVWLKENPEISDEIDAKIRALNGVEMHITEGT.....QDET DGER.....PEE.....	348
GAKIGQGGKNVRVWLKENPEISDEIDAKIRALNGVEMHITEGT.....QDET DGER.....PEE.....	348
GAKIGQGGKNVRVWLKENPEISDEIDAKIRALNGVEMHITEGT.....QDET DGER.....PEE.....	348
GERIGQGGKENVRFLENPKIAGEIETRLREKLLPHRQGEKIA..EEVNAAE.....	343
GERIGQGGKGNREFLENAELARLEIENKVRHLGVTPMGAV.T.LAEVEED.....	352
GDKIGQGGKANACKYLKENPEIAKTLDKKLREMLLNPNMQLIA..ETSSAAD DVEFGAV.....PEEF.....	354
GDKIGQGGKANACKYLKENPEIAKTLDKKLREMLLNPNMQLIA..ETSSAAD DVEFGAV.....PEEF.....	354
GSKIGQGGKANAAKYLEENPEIGQIEEQQIRAKLLVTSANTKAT..PVAEDLADADI.....	347
GNKIGQGGKNNVIRYLEENPQISTEIEAVIREQLLTKASDQTAA...HDETEEEP DLL.....ES.....	349
GNKIGQGGKNNVIRYLEENPQIAEIEIERNIREQLLTGTNGAVQ...I.EDETEEEP DLL.....LES.....	349

logo

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NTDB id 228 SPD RS09265 WP 001085462.1	EIEE	388
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NTDB id 1074 ABD1 RS10195 WP 000344167.1		349

consensus

 non conserved
 similar
 ≥ 50% conserved