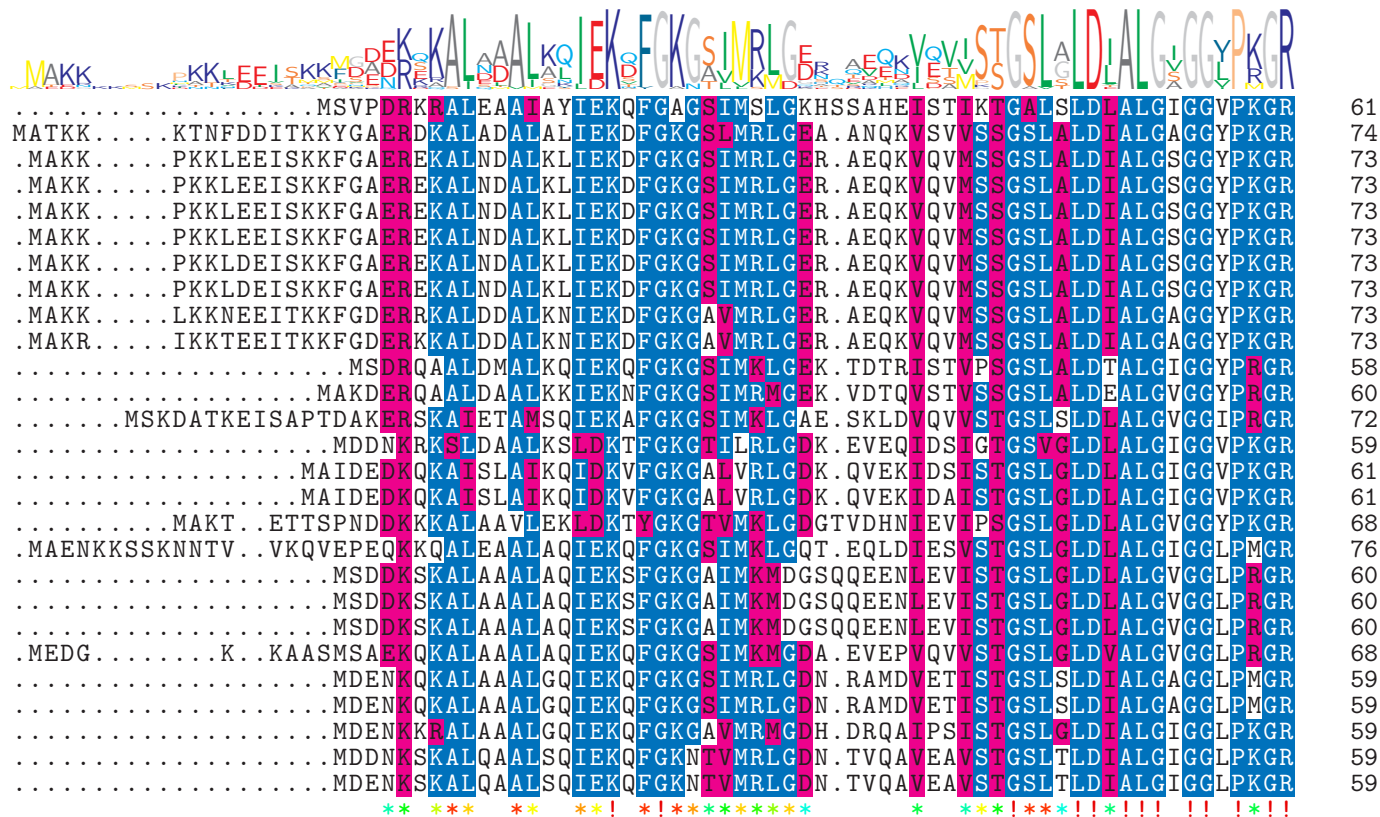


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

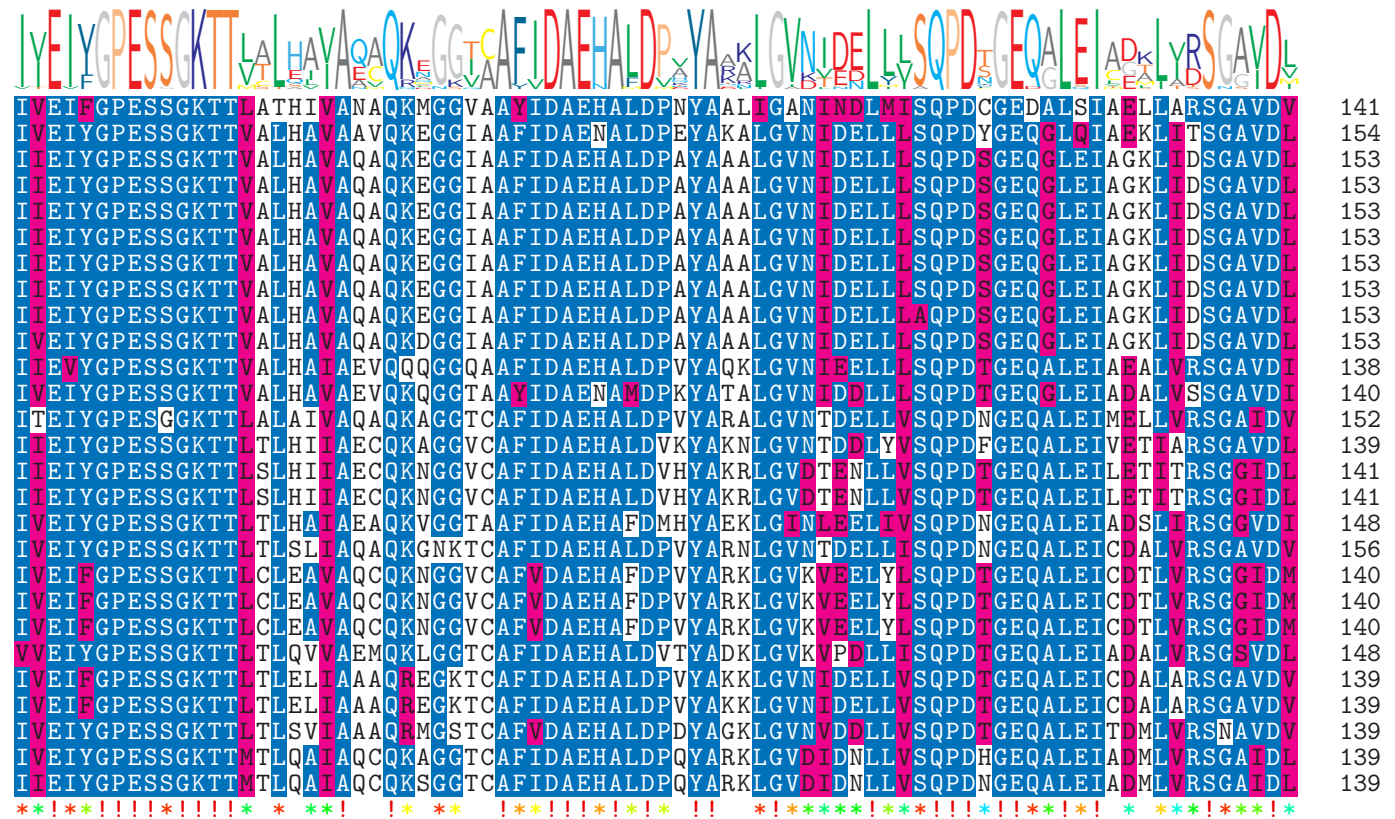
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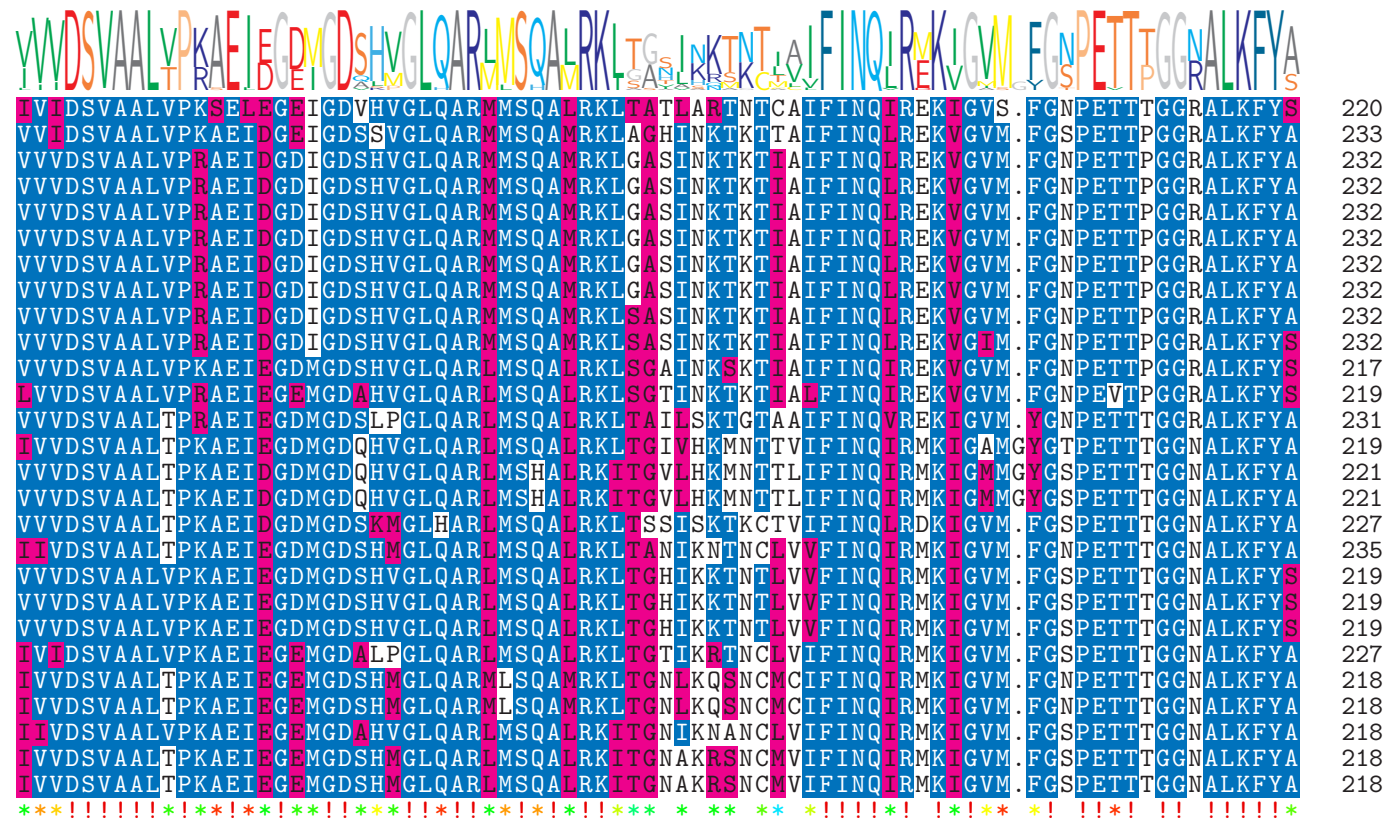
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consensus

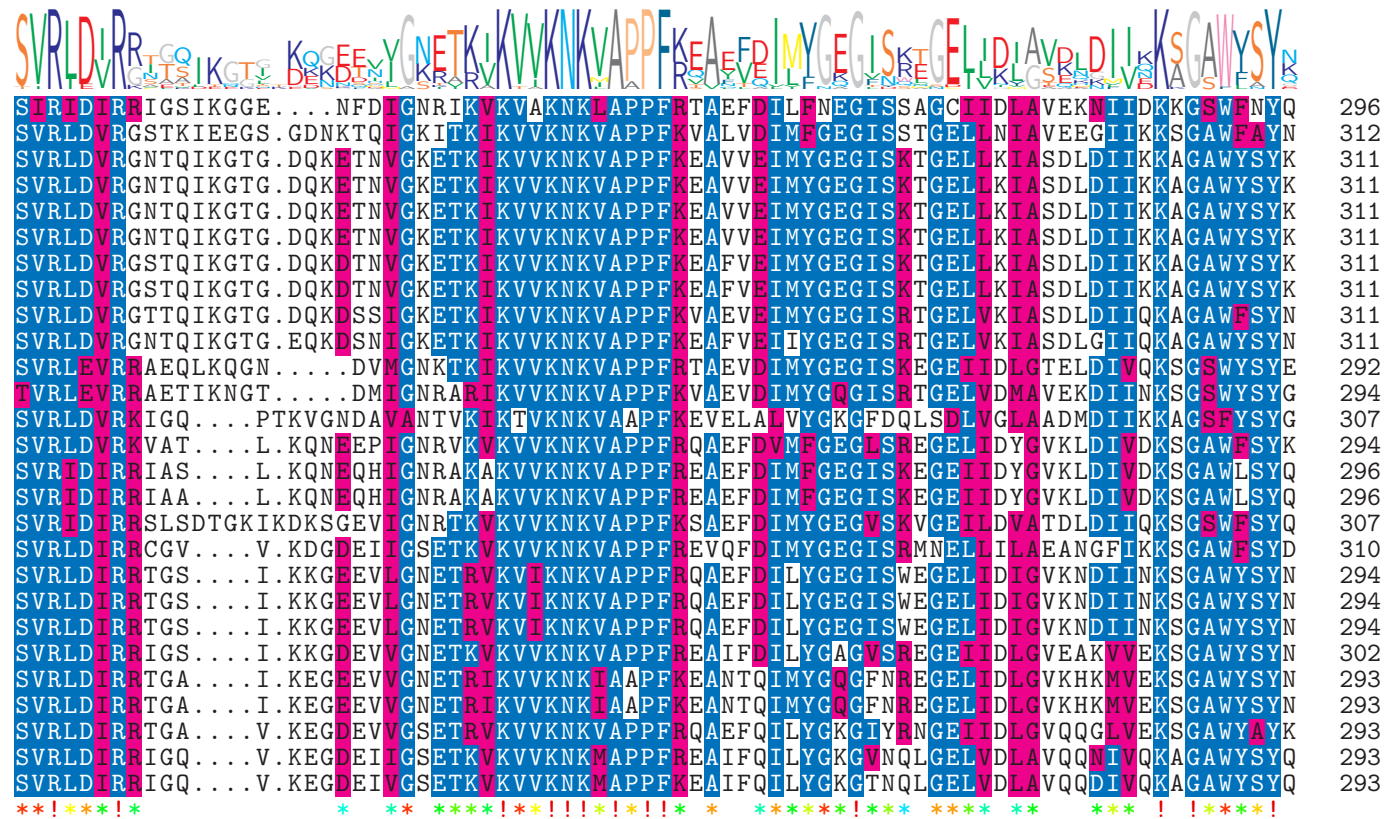
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consensus

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NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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Sequence	Score
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DEKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGQGSSENAKKYLAADNPEIFDEIDHQVRVQFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDLEIE	381
DEKIGQGSSENAKKYLAADNPEVFAEIDHQVRVQFGLIDGEEAAES...KKDE...VAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....ETDEK..ADKNLVLELDDTIELE	377
GEKIGQGSSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGRENAKQFLKENKDIMLMIQEIGIREHYGLDNGGVVQQAAETQEEL.EFE.E.....	348
SERIGQGRENAKNYLADHEDVEURLKVRVRAAYGISDVPEEDLPTTEDE.QI.NILPDST.....EE.....	355
DIRIGQGEKTIAYIAIERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE..A.....	363
DKKLGQGRENSKAFLENPEIADAITKAIQNSMGTIEGMI...SGSEDDE..GE..E.....	343
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP..LE..EME.....	347
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP..LE..EME.....	347
DTKLGGGRDTPVKELIKDNPELAEELEAKIREKLQAK.....	343
GTKIGQGGKNNAVKWLKENPEIADKLEQEIRNTLASNPNS...VLKATASDSND..ESDVVEEFSEEL..	372
GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....Q..DETDERPEE.....	348
GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....Q..DETDERPEE.....	348
GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....Q..DETDERPEE.....	348
GERIGQGRDNCREFLRENAELARENENKVRHLGVTPMG...AV.TLAEVEVE..D.....	352
GDKIGQGGKANACKYLKENPEIAKTLDKKLREMLLNPNEM...QLIAETSSAAD..DVEFGAVPEEF..	354
GDKIGQGGKANACKYLKENPEIAKTLDKKLREMLLNPNEM...QLIAETSSAAD..DVEFGAVPEEF..	354
GSKIGQGGKANAAKYLEENPEIGQEIEQQIRAKLLVTSAN...TKATPVAEDLA..DADI.....	347
GNKIGQGGKNNVIRYLEENPQISTEIEAVIREQLLTKASD...QTAA.HDETEE..EPDLL..ES.....	349
GNKIGQGGKNNVIRIFYENNQIAEIERNIREQLLTGTN...GAVQ.I.EDEE..EPDLL..LES.....	349

logo

		IEE	
NTDB id 29858 CTL RS00095 WP 009872817.1	...	352	
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	...	349	
NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349	
consensus			

- X non conserved
- X similar
- X ≥ 50% conserved