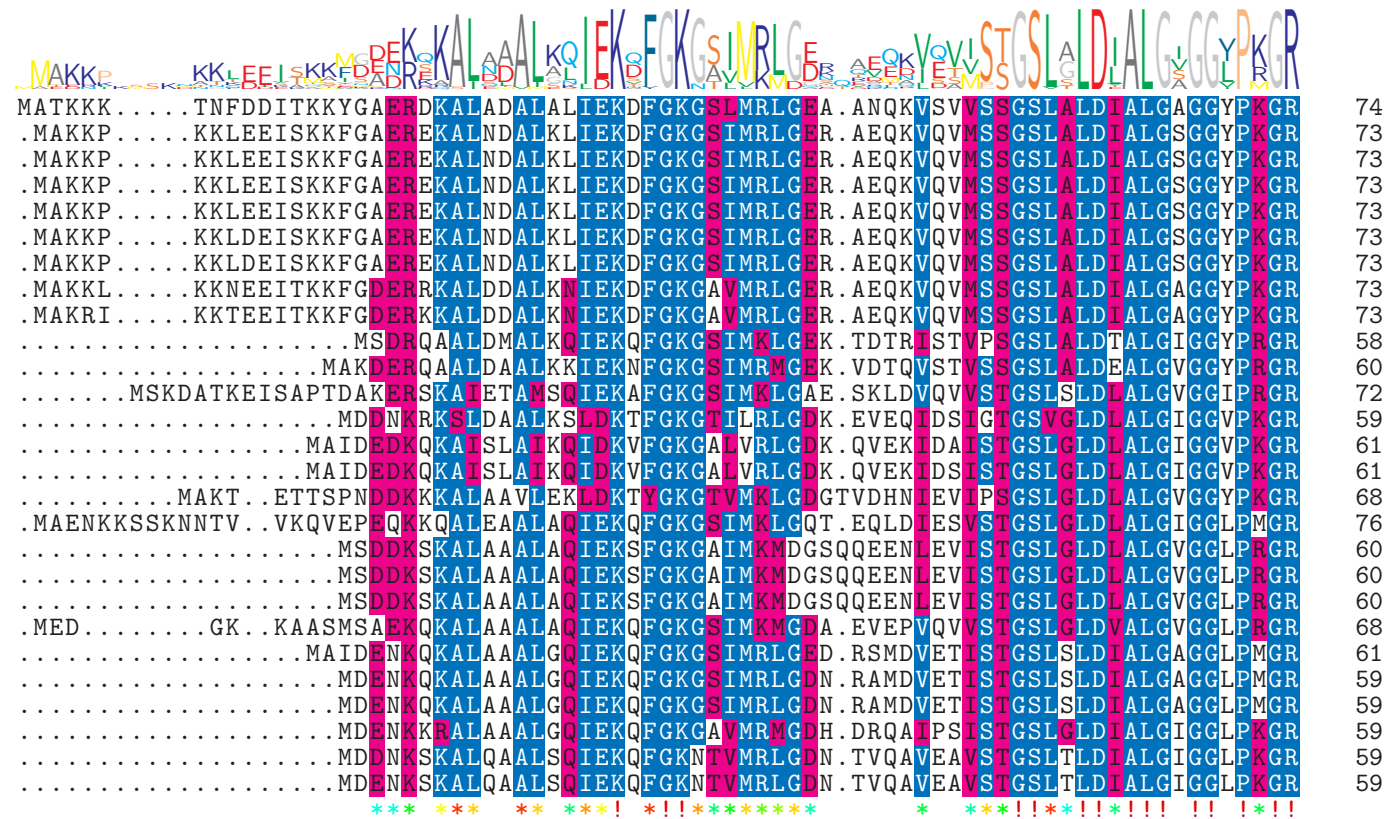


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consensus

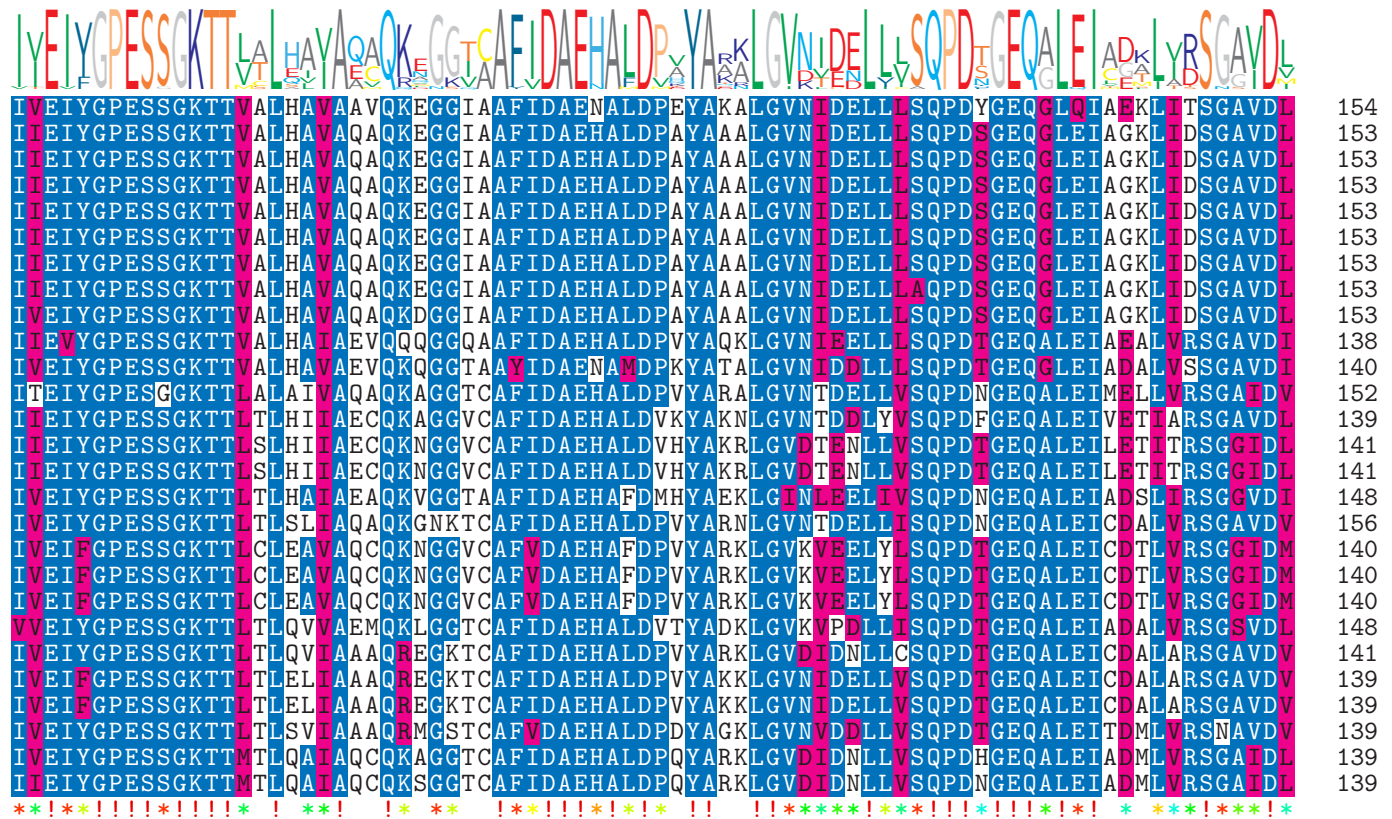
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

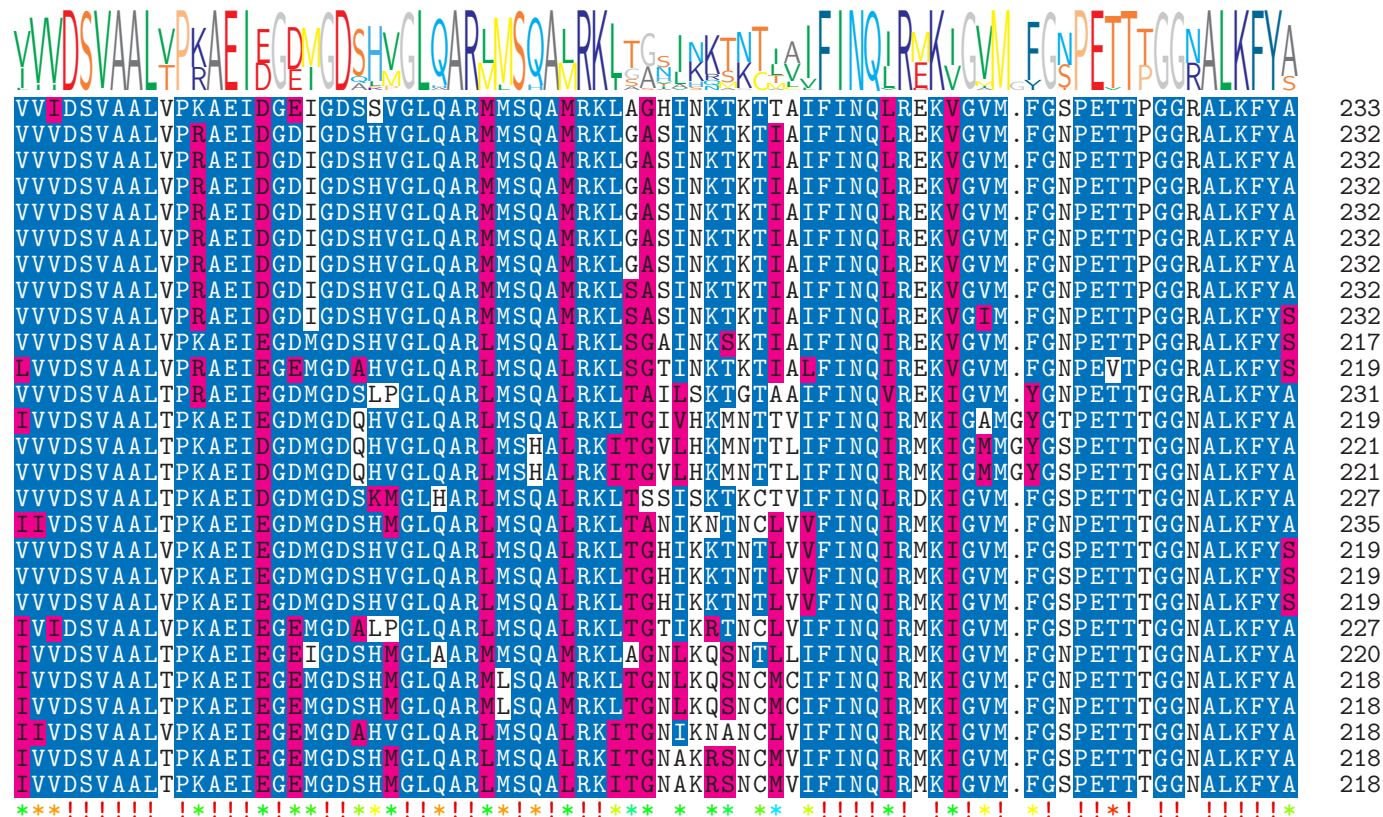
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

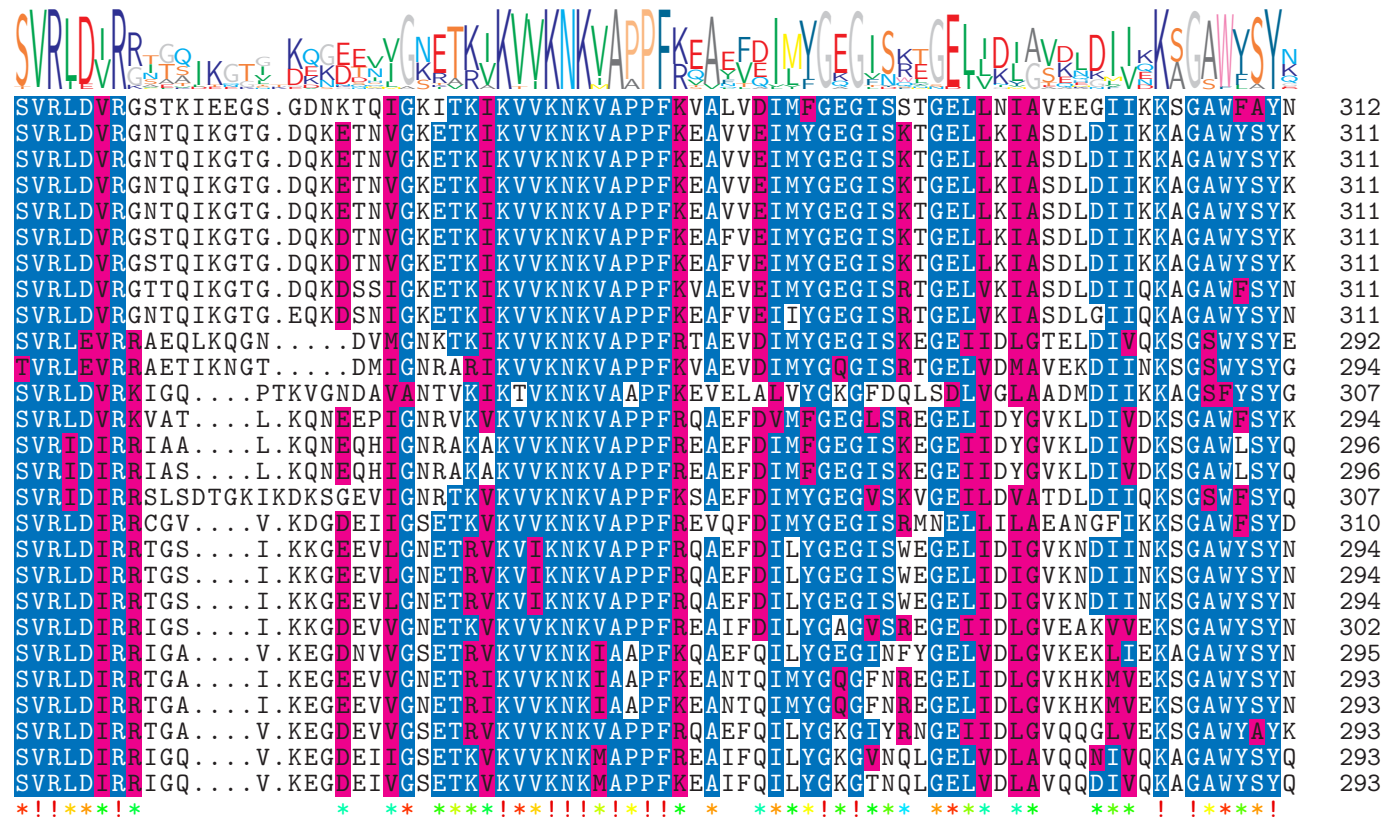
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consensus

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consensus

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Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGQGSENAKKYLAADNPVFDEIDHQVRVQFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDLEIE	381
DEKIGQGSENAKKYLAADNPVFDEIDHQVRVQFGLIDGEEIAES...KKDE...VAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPFIIDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGGGRENAAKQFLKENKDIMLMIQEIQIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGGGRENAAKNYLADHEDVEDEVRLKVRAAAYGISDVPPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGGQKEKTIATYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAA...EA.....	363
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GERIGGQRDNCREFLRENAELAREIENKVRHLGVTPM.....GAVTLAE.EVE...ED.....	352
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GDKIGGQKANACKYLKENPEIAKTLDDKKLREMLLNPN.....MQLIAETSSAA..DDVEFGAVPEEF.....	354
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GSKIGGQKANAACKYLEENPEIGQIEEQQIRAKLLVTSA.....NTKATPVAEDL...ADADI.....	347
GNIKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTKAS.....DQTAA.HDETE...EEPDLLES.....	349
GNKIGGQKNNVIRIFYENNQIAEIEIRNIREQLLTGT.....NGAVQ.I.EDE...EEPDLLES.....	349

logo

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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349

consensus

 non conserved
 similar
 ≥ 50% conserved