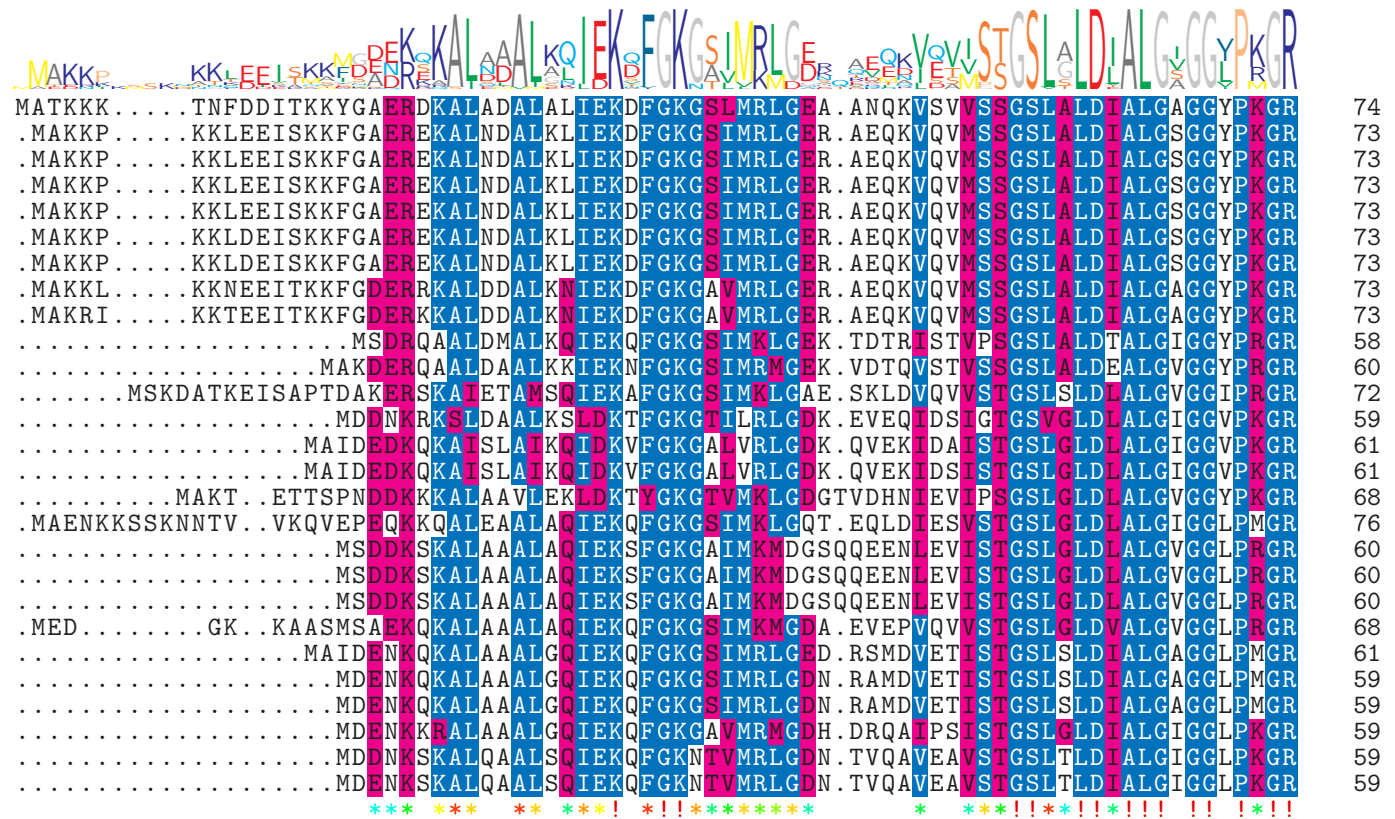


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consensus

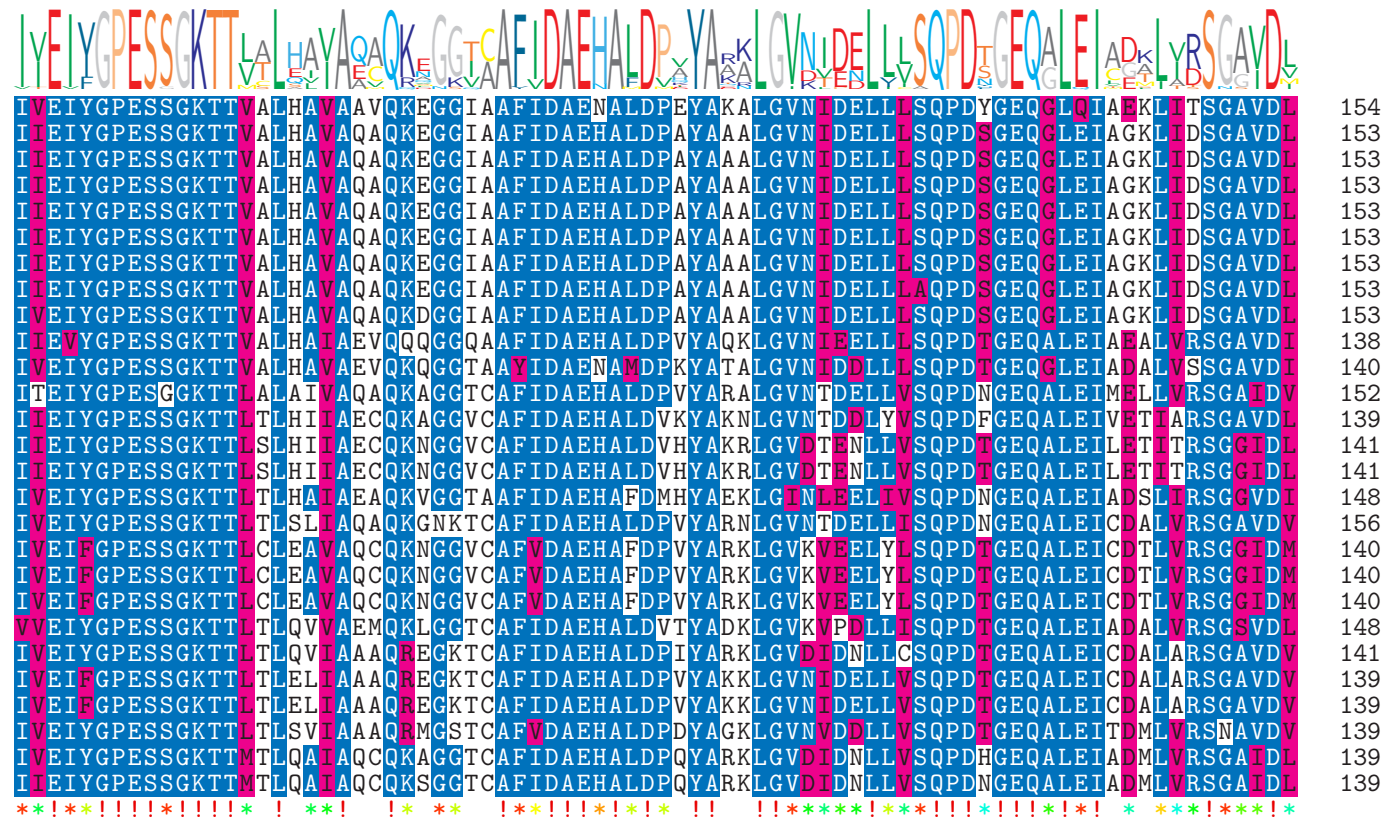
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

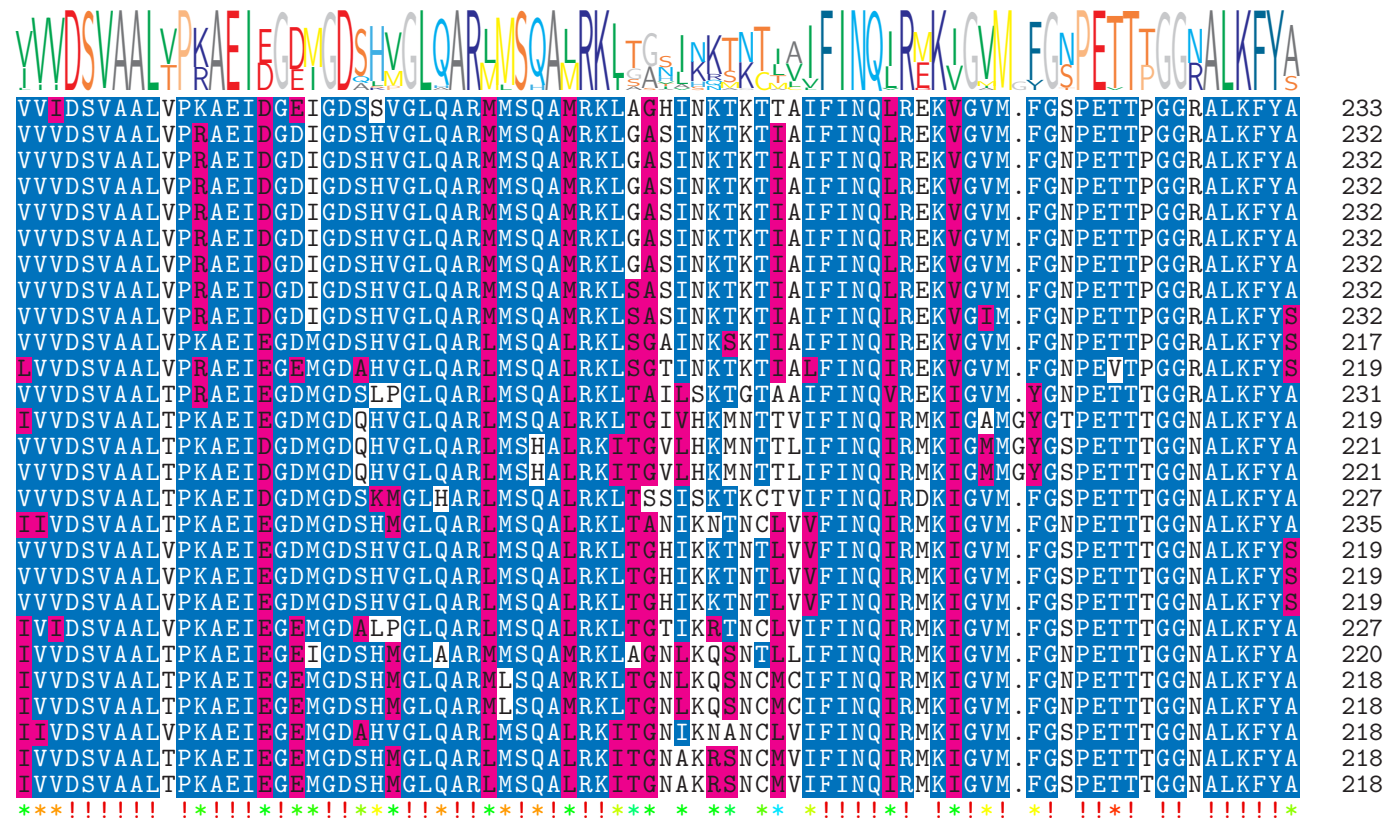
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consensus

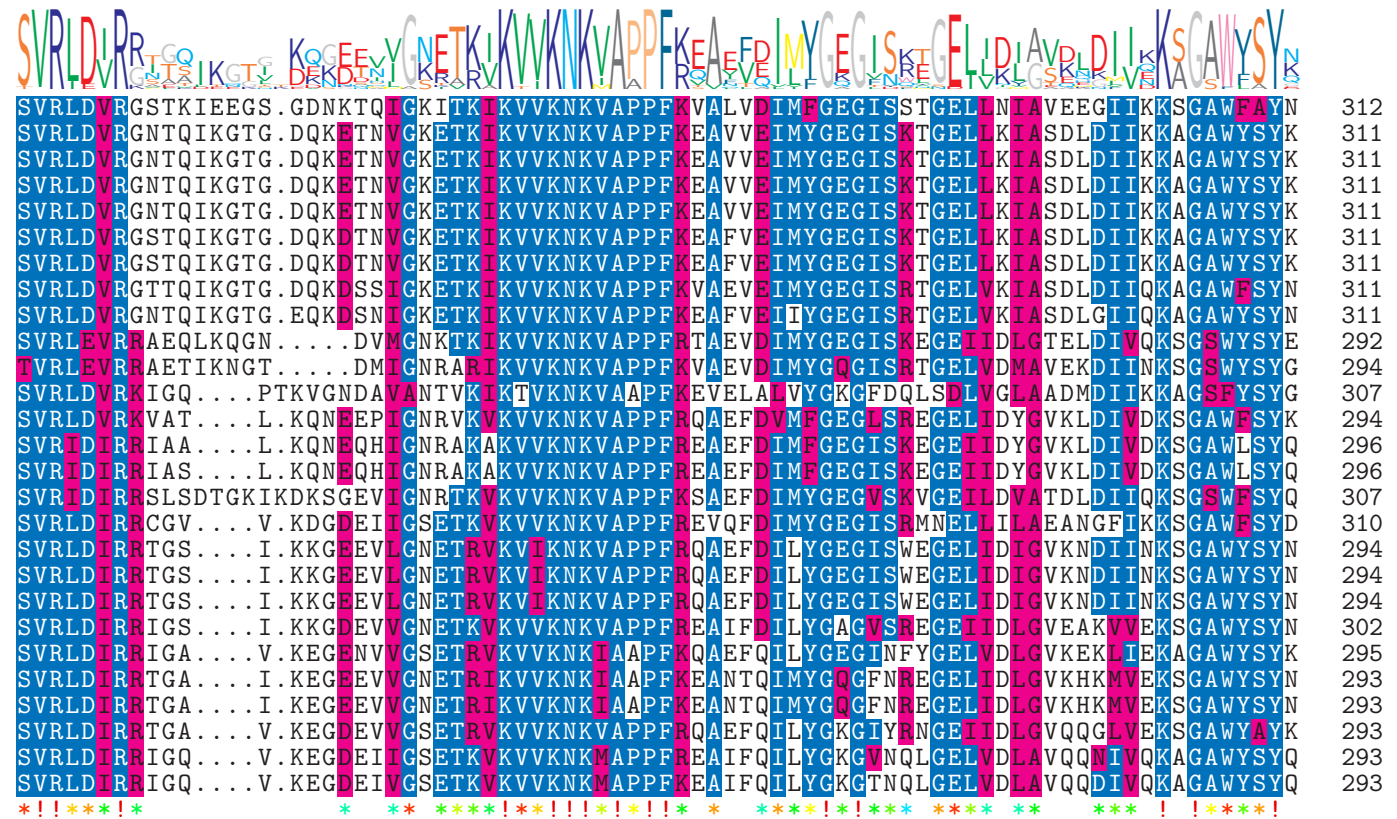
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consensus

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consensus

Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGQGSENAKKYLAADNPVFDEIDHQVRVQFGLIDGEEIAES...KKDE...VAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPFIIDIDHKVRVQFGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGGGRENAAKQFLKENKDIMLMIQEQIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGGGRENAAKRYLADHEDVEDEVRLKVRVQFGLSDVPEEDLPTTEDE.QI.NILPDDST....EE.....	355
DERIGGKEKTIATIAERPMEQETIDRVRMAAIRAGNAGEAPALAPAPAAPEAA...EA.....	363
DKKLGGGRENKAFLENPEIADITKATINSMGTIEGM.....ISGSEDDGE.....EE.....	343
DKKLGGGRENAAKALLKEDKALADEITLKIKESIGSNEE.....IMPLPDEPL.....EEME.....	347
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DTKLGGGRDITVKELIKDNPELAEELEAKIREKLQAK.....	343
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GERIGGGRDNCREFLRENAELAREIENKVRHLGVTPM.....GAVTLAEE.VE...ED.....	352
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GSKIGGQKANAAKYLEENPEIGQIEEQIRAKLLVTSA....NTKATPVAEDL..ADADI.....	347
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GNKIGGQKNNVIRIFYEENQIAEIERNIEREQLLTGT....NGAVQ.I.EDE...EEDPLLLES	349

logo

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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349

consensus

 non conserved
 similar
 ≥ 50% conserved