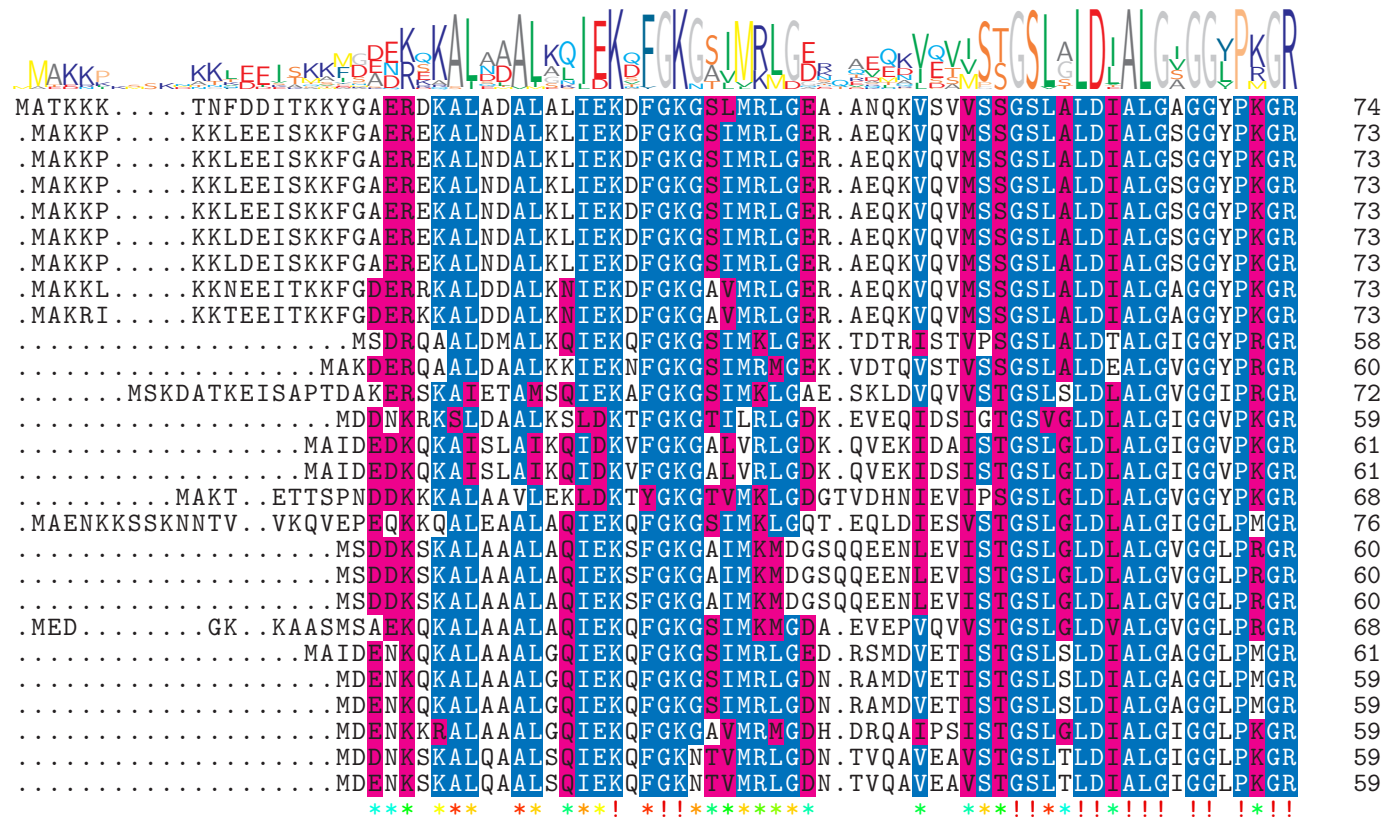


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consensus

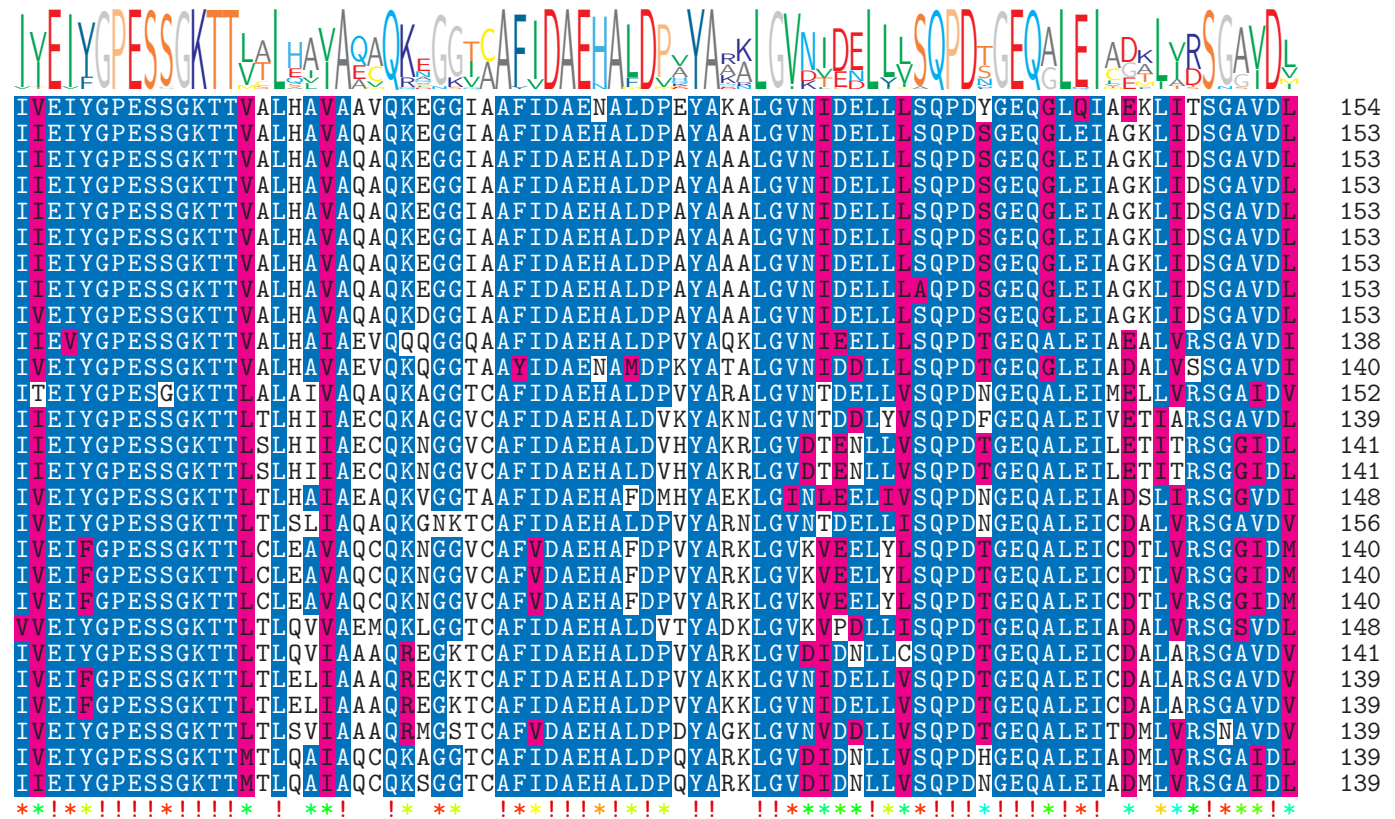
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consensus

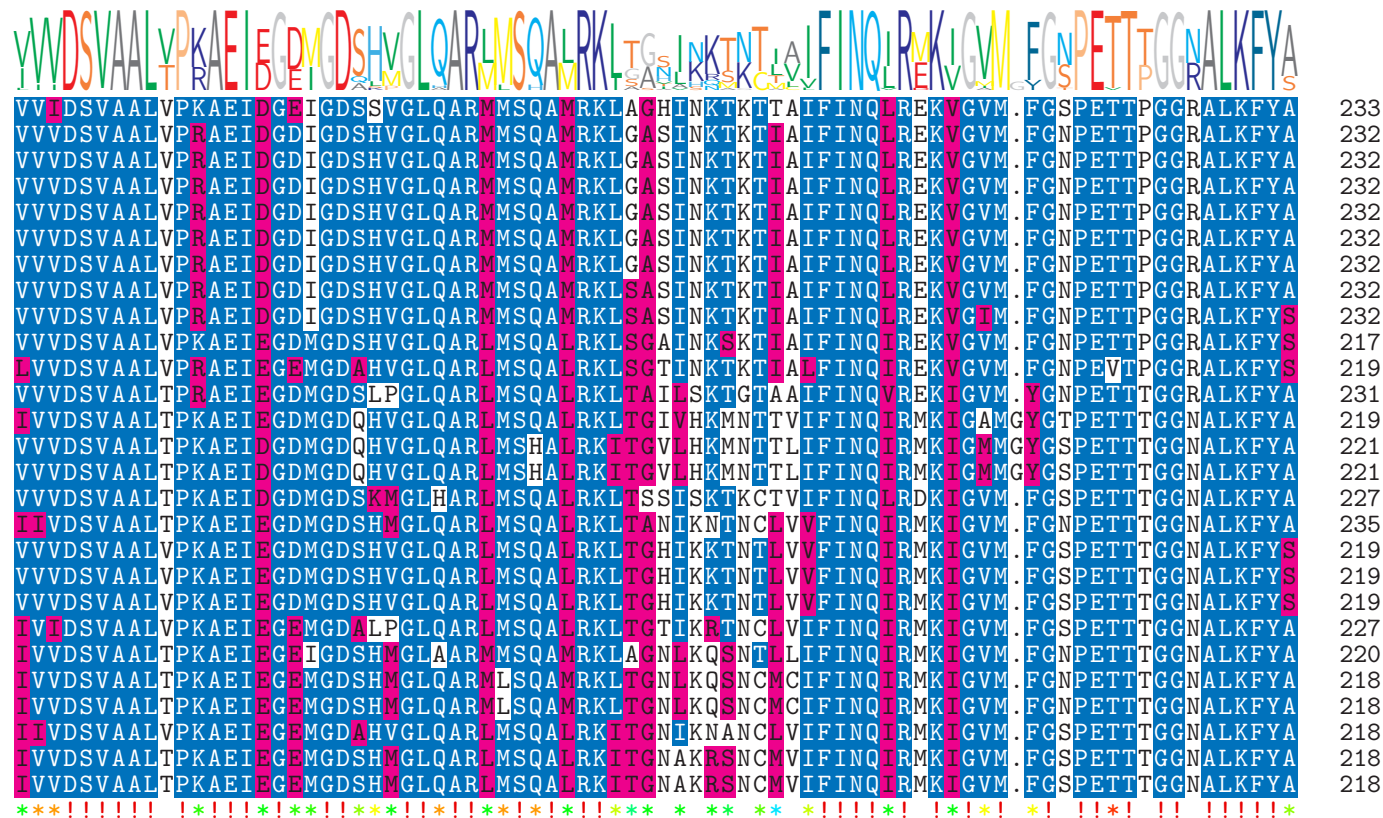
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consensus

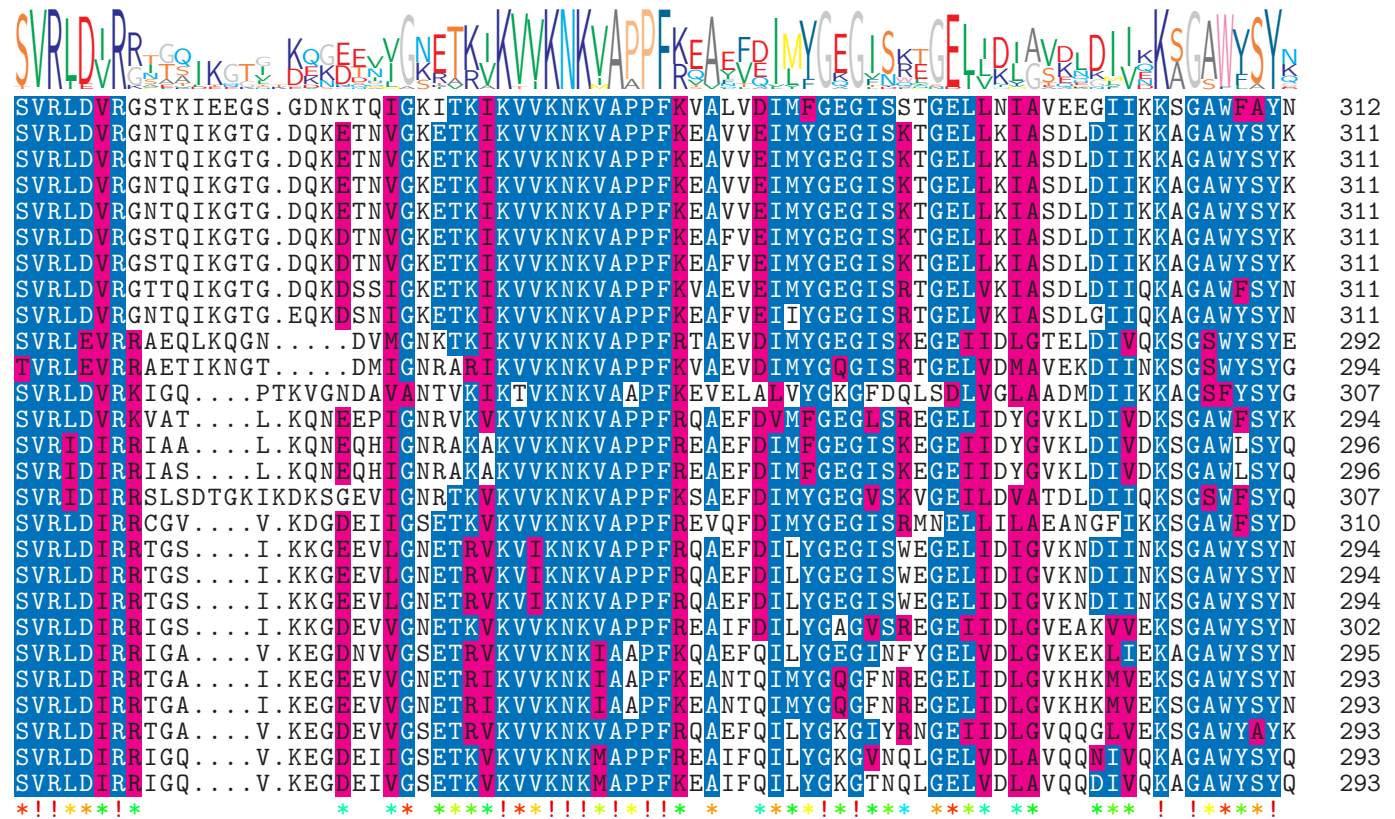
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consensus

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Sequence	Score
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DEKIGQGSENACKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDNTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGQGSENACKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDNTENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGSENACKKYLAENPEIFDEIDHQVRVQFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDLEIE	381
DEKIGQGSENACKKYLAENPEVFAEIDHQVRVQFGLIDGEEAAES...KKDE...VAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGQGRENAKNYLADHEDVEDEVRLKVRAAAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
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GERIGQGRDNCREFLRENAELAREIENKVRHLGVTPM....GAVTLAE.EVE...ED.....	352
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GSKIGQKANAAKYLEENPEIGQIEEQQIRAKLLVTSA....NTKATPVAEDL..ADADI.....	347
GNIKIGQKNNVIRYLEENPQISTEIEAVIREQLLTKAS....DQTAA.HDETE...EEPDLLES.....	349
GNKIGQKNNVIRIFYENNQIAEIEIRNIREQLLTGT....NGAVQ.I.EDE...EEPDLLES.....	349

logo

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NTDB id 1022 ACIAD RS06390 WP 004925639.1	...	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349

consensus

 non conserved
 similar
 ≥ 50% conserved