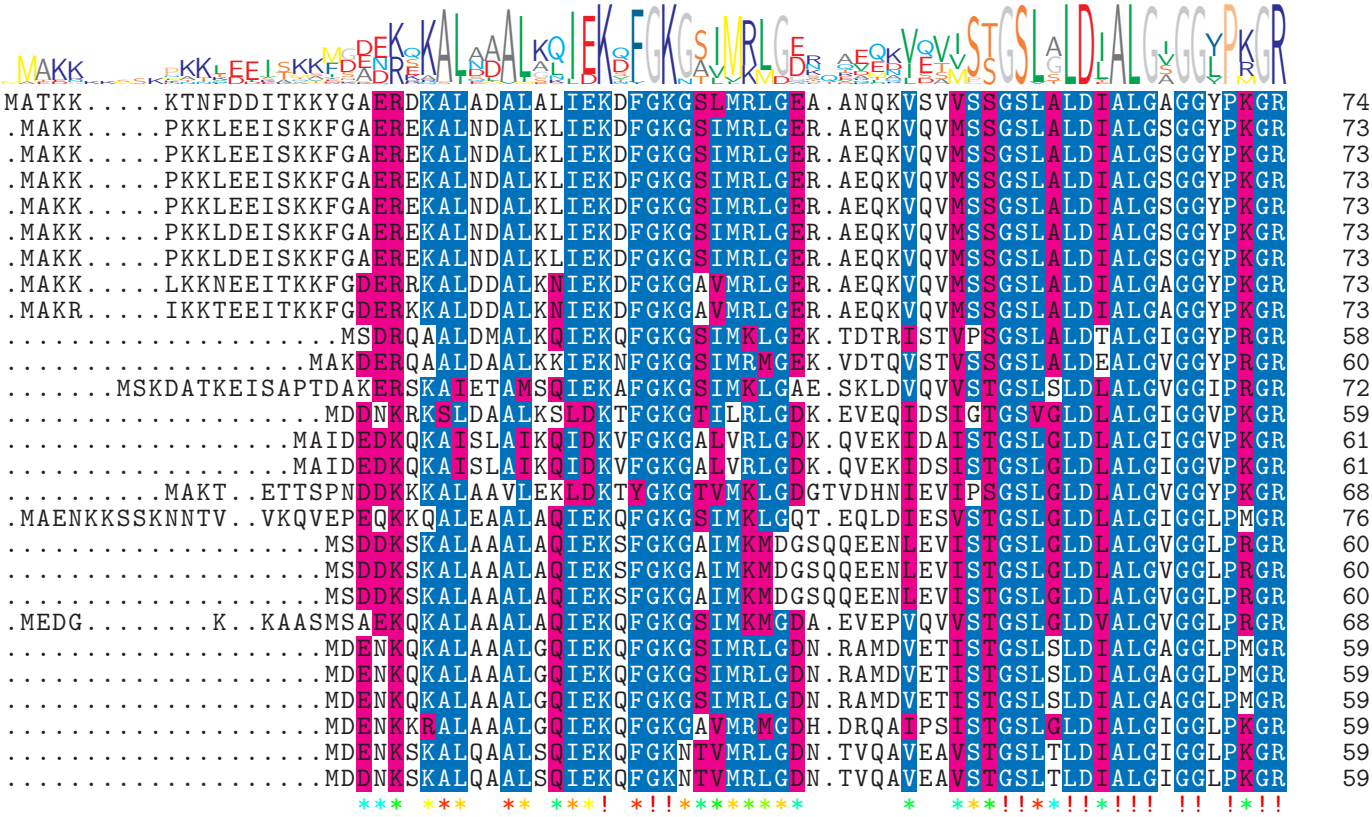


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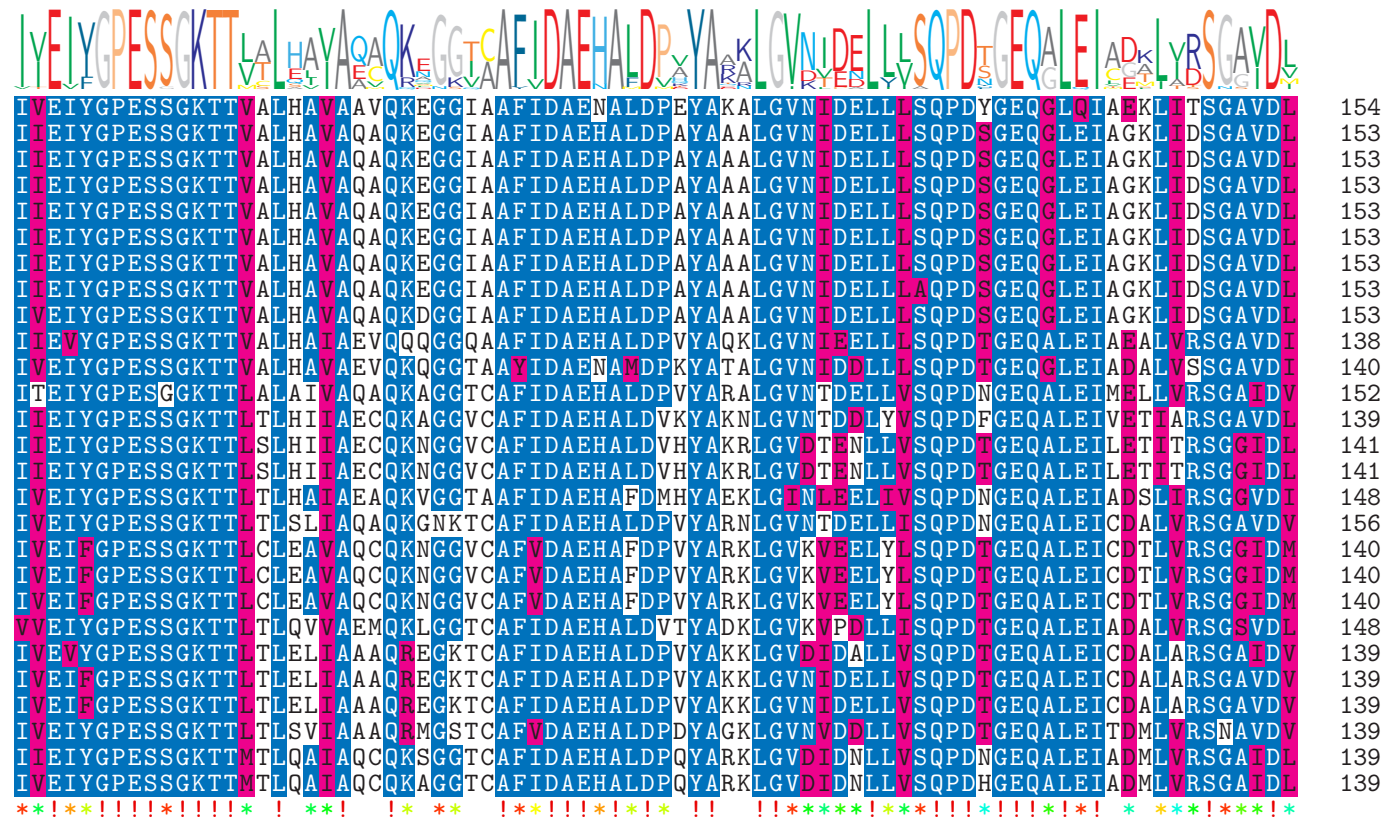
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
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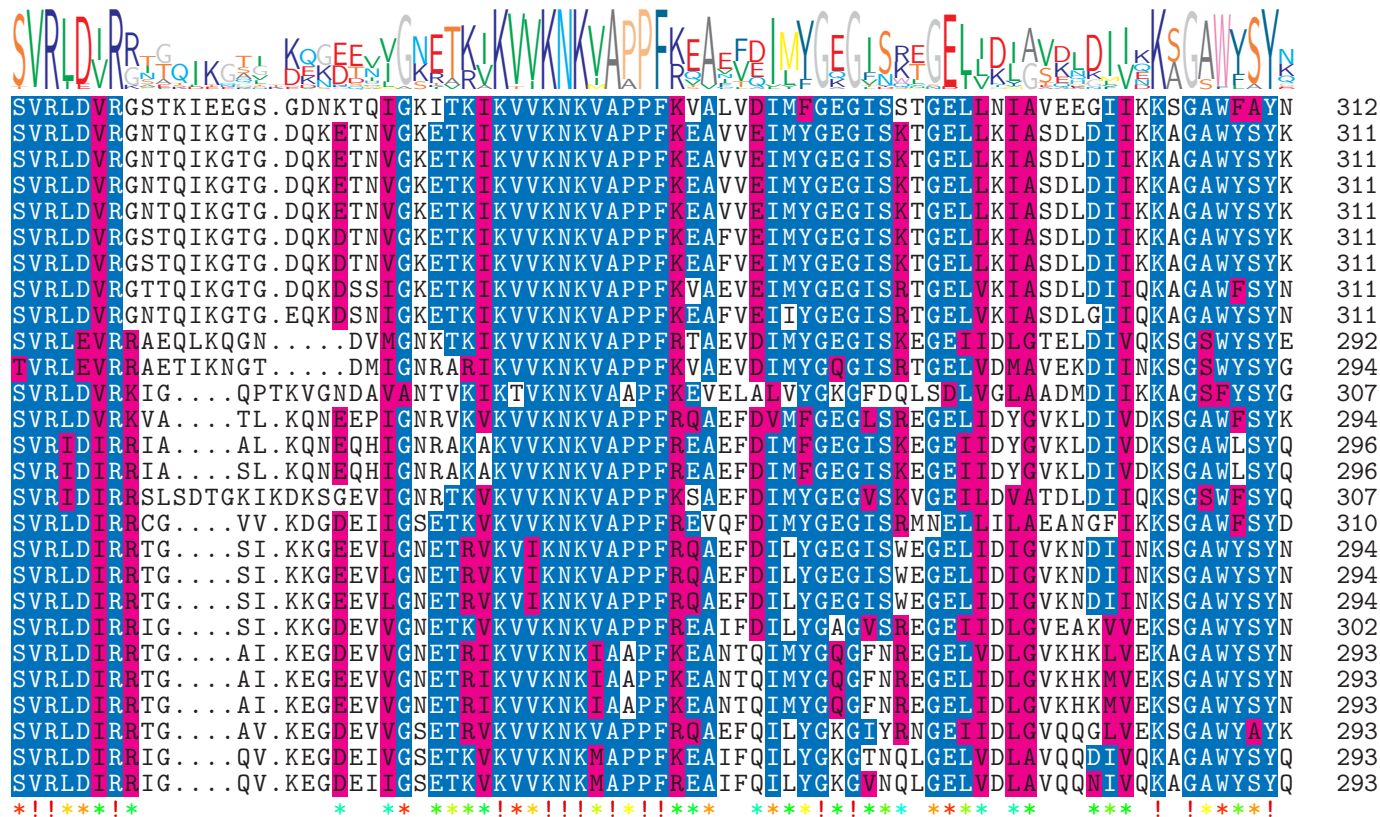
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Sequence	Score
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DEKIGQGSENAKKYLAHNPEVFAEIDHQRVVFGLIDGEEAAE...SKKDE...VAQVDS..VNEEVTLDLGDLEIE	381
DEKIGQGSENAKKYLAHNPEIFDEIDHQRVVFGLIDGEEAAE...SKKDE...AAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEEESAMAVASE.....E.....TDDLALDLNGIEIE	377
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EERLGGGRENAAKQFLKENKDIMLMIQEQUIREHYGLDNGVGVVQQAEETQEEL.EFE.E.....	348
SERIGGGRENAAKNYLADHEDVEDEVRLKVRAAAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE	355
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GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQD.....E.....TD.....GERPEE	348
GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQD.....E.....TD.....GERPEE	348
GERIGGGRDNCREFLRENAELAREIENKVRHGLVTPMGAVTLAE.EVEED.....	352
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GDKIGGQKANACKYLKENPEIAKTLDDKLREMLLNPNENMQLIAETSSAADVV.....EF.....GAVPEEF	354
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GNKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTASDQTAA.HDETEEEP.....DL.....LES	349

logo

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consensus

 non conserved
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 ≥ 50% conserved