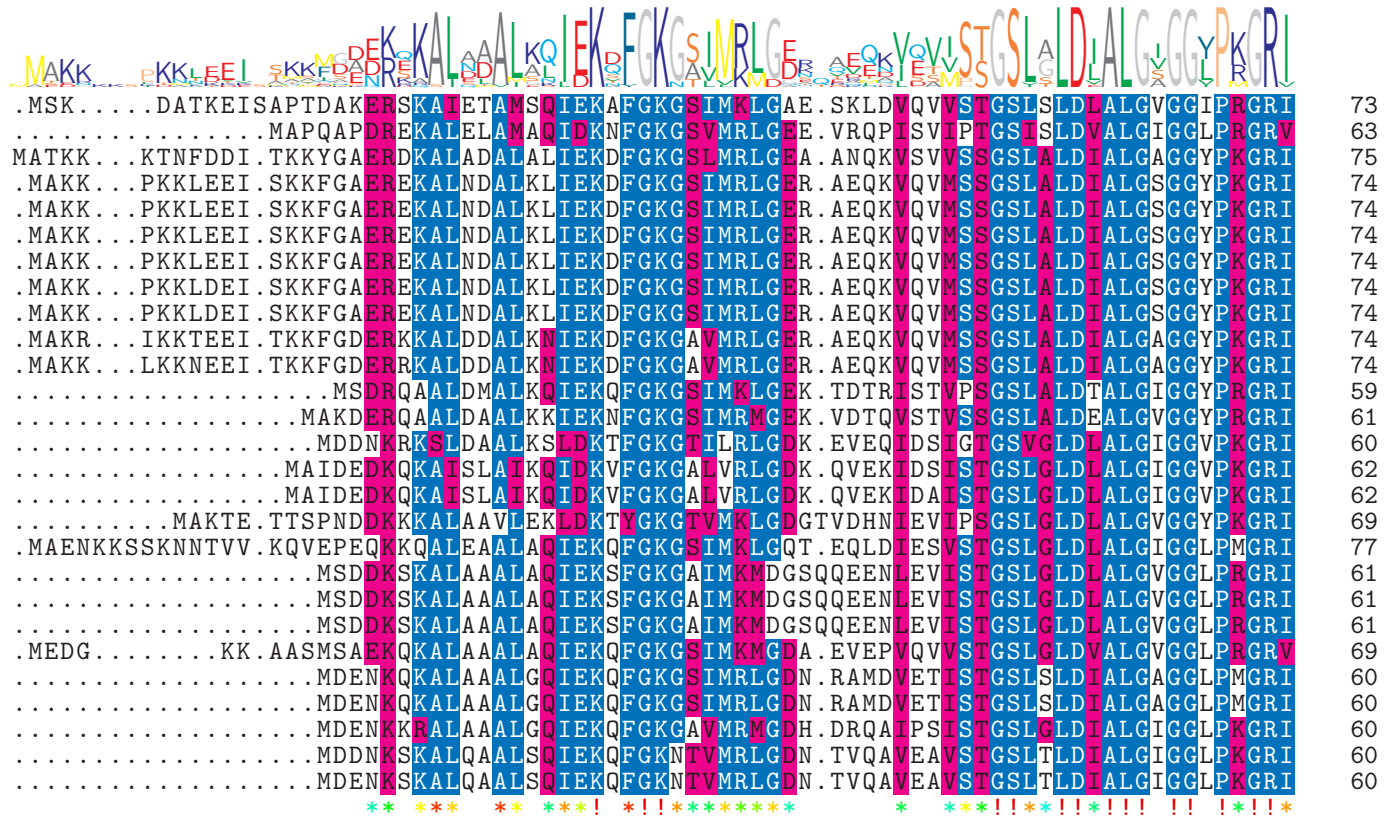


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consensus

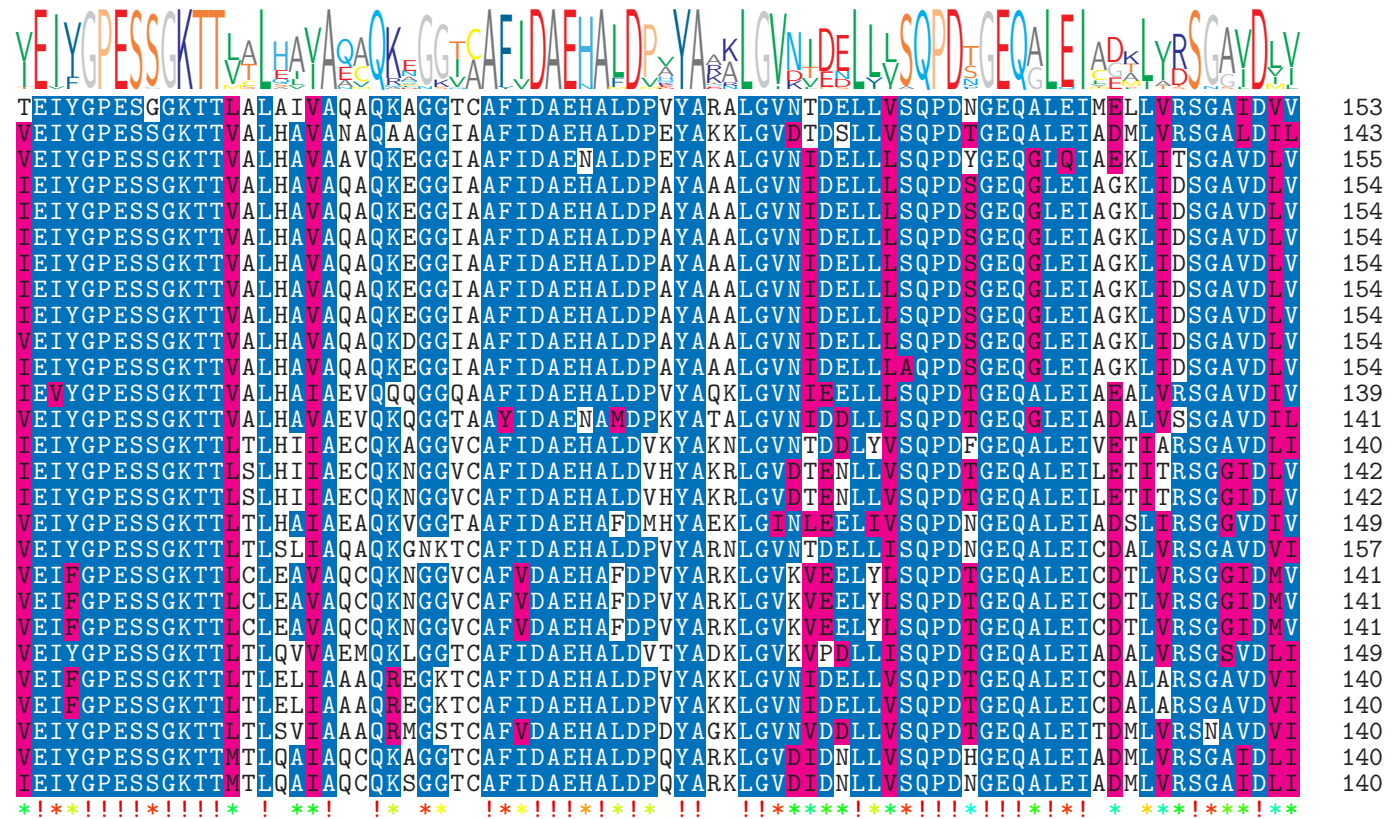
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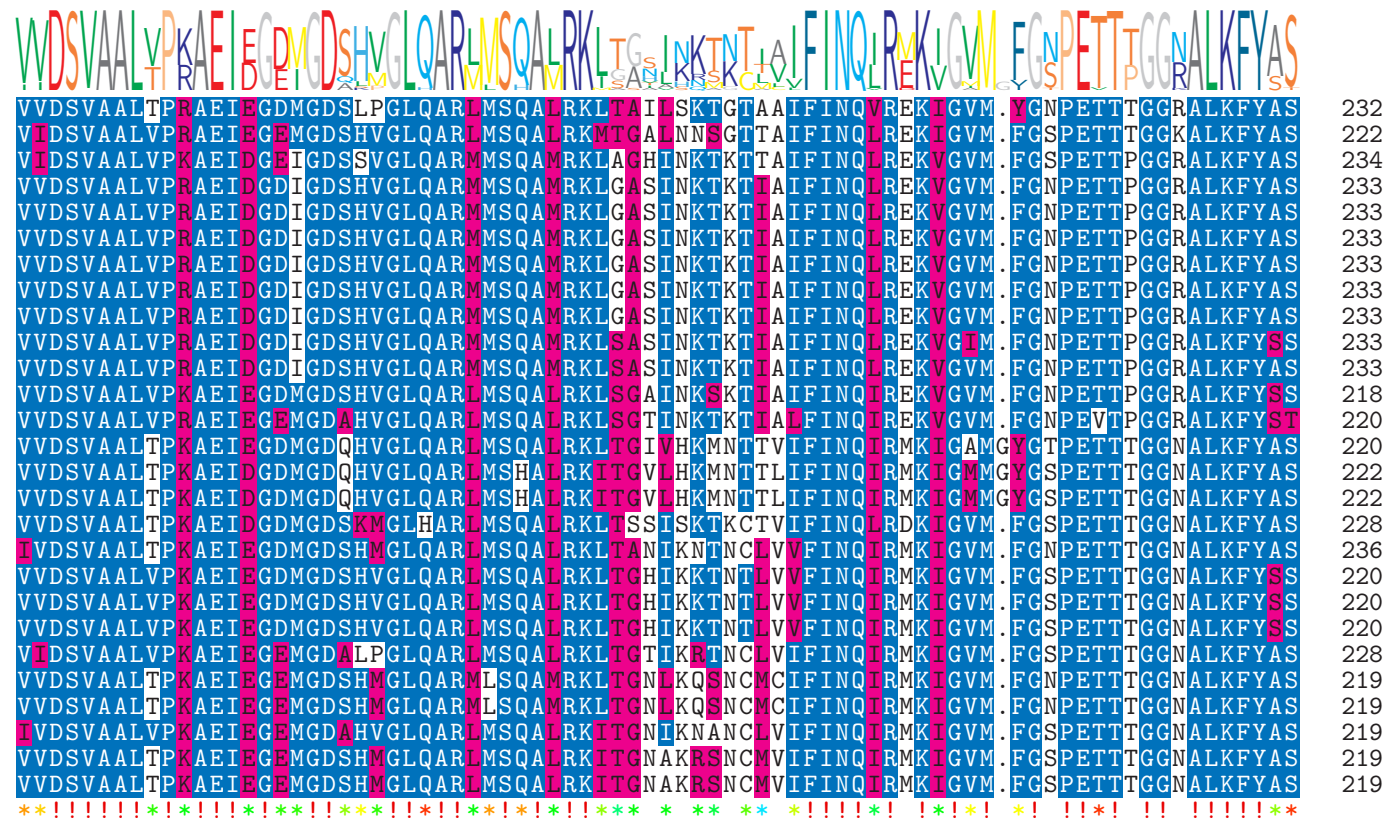
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consensus

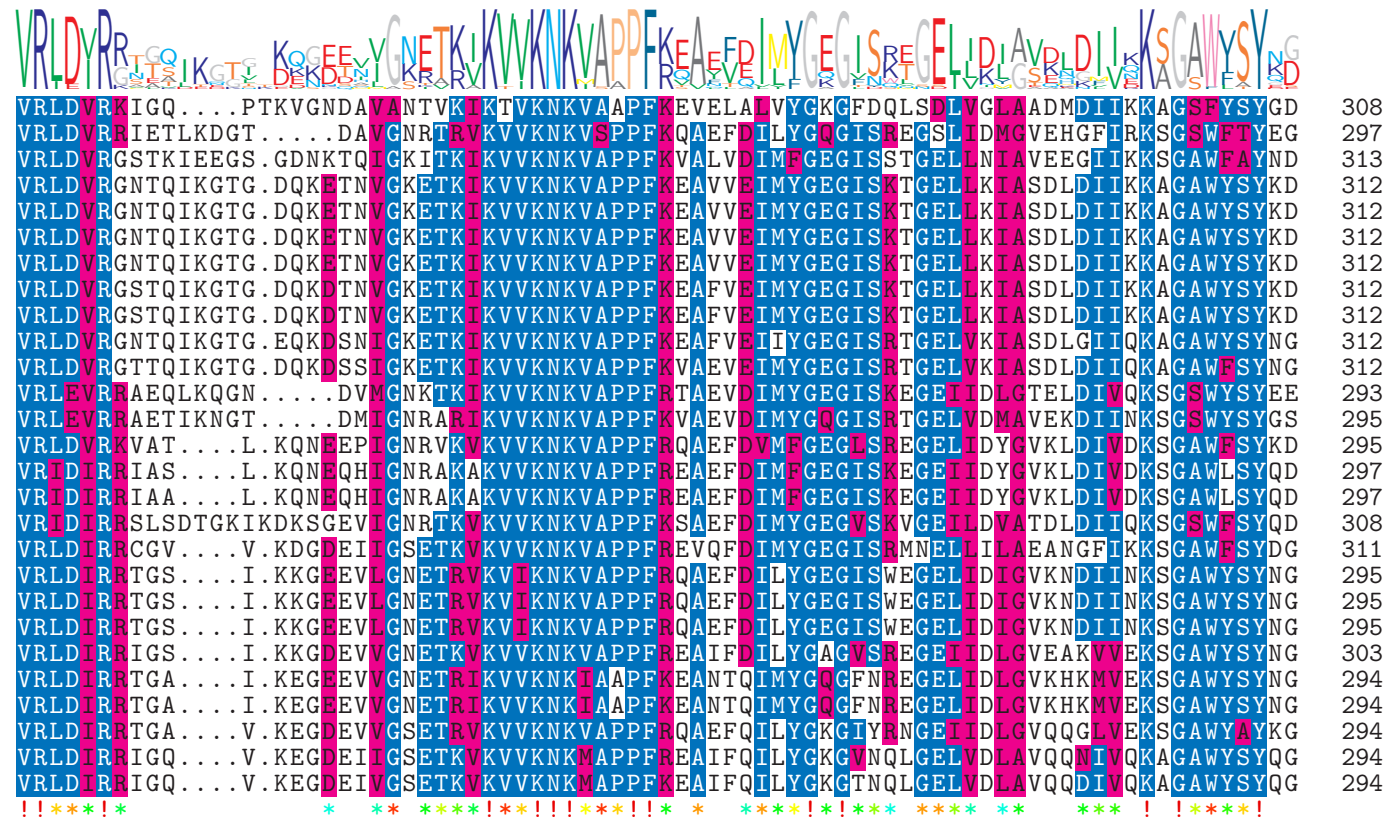
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

Sequence	Score
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EKIGQGAEKAKNYLKEHQDVFEIDHKVRAAHGILDDSEVATEEETTASKTAKAKKEEKAVETEEIELELED.....	387
EKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEI	386
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EKIGQGSSENAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDELEIEI	382
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KKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEIMPLPDEP..LEEM.E.....	347
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TKLGQGRDVTVKELIKDNPELAEELEAKIREKLQAK.....	343
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DKIGQGGKANACKYLKENPEIAKTLDDKKLREMLLNPNMQLIAETSSAADV.EFGAVPE.....EF.....	354
SKIGQGGKANAAKYLEENPEIGQETIEQQIRAKLLVTSANTKATPVAEDLADA.DI.....	347
NKIGQGGKNNVIRYLEENPQISTEIEAVIREQLLTKASDQTA.HDETEEEP.DLL..ES.....	349
NKIGQGGKNNVIRYFENNQIAEIERNIREQLLTGTNGAVQ.I.EDEEEP.DLL..LE.....S.....	349

logo

		EE	
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consensus

 non conserved
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 ≥ 50% conserved