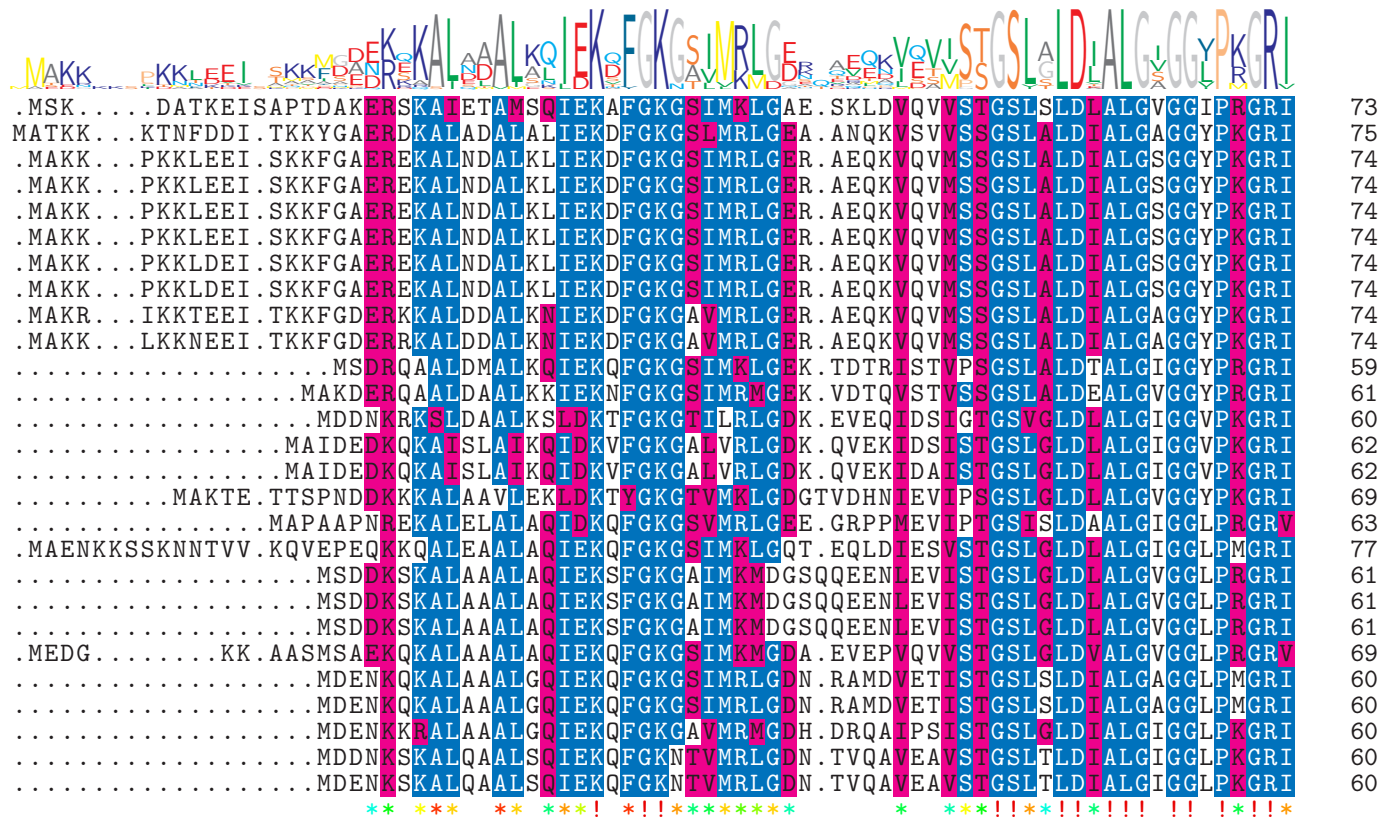


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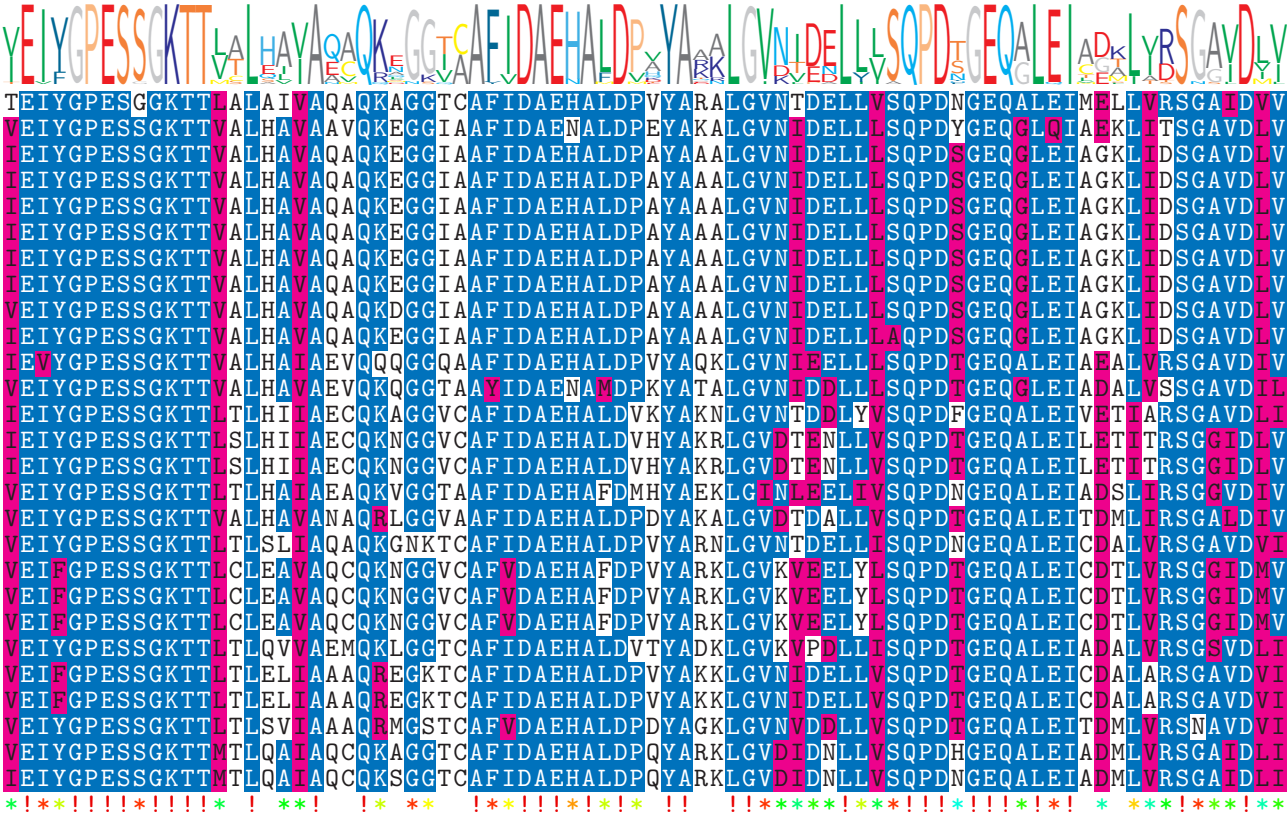
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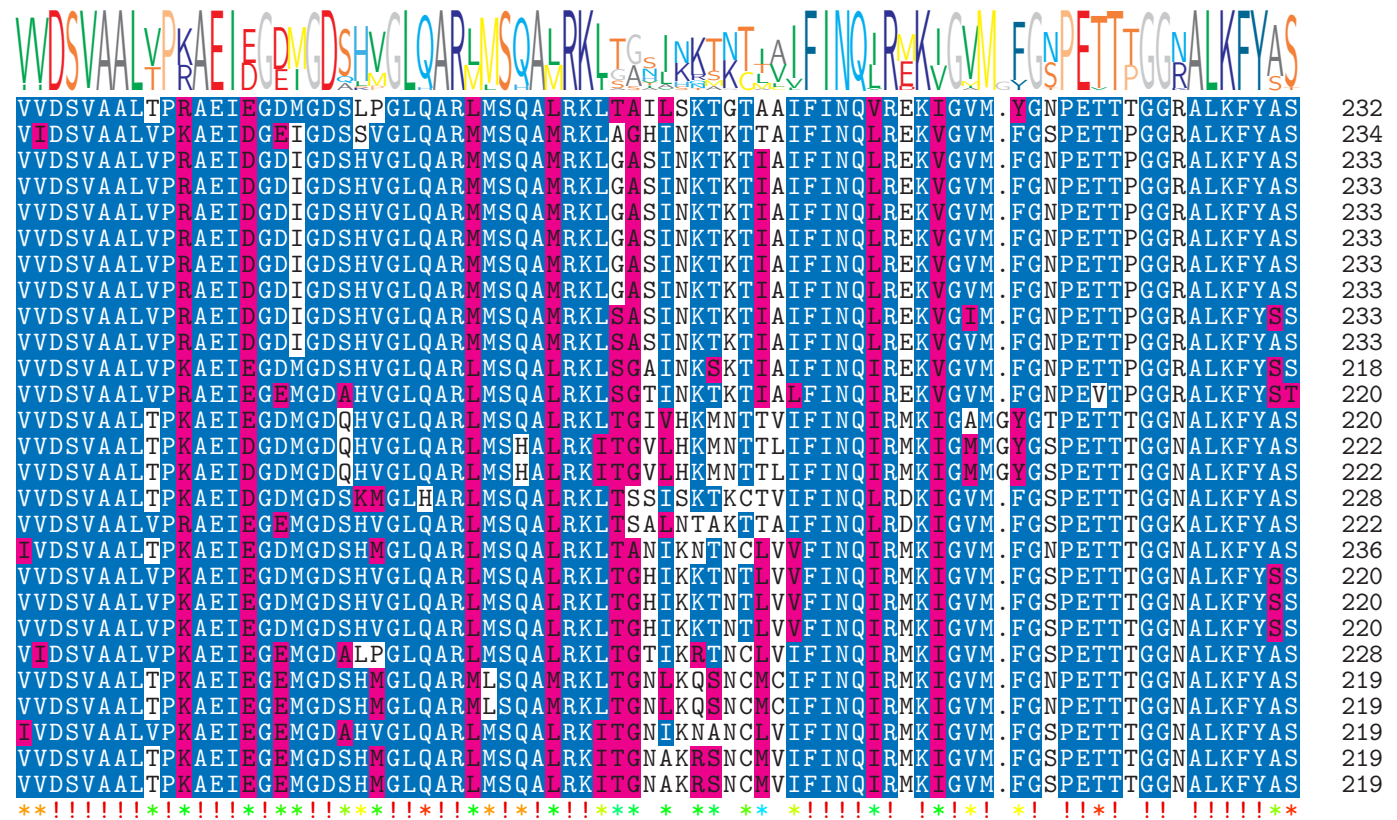
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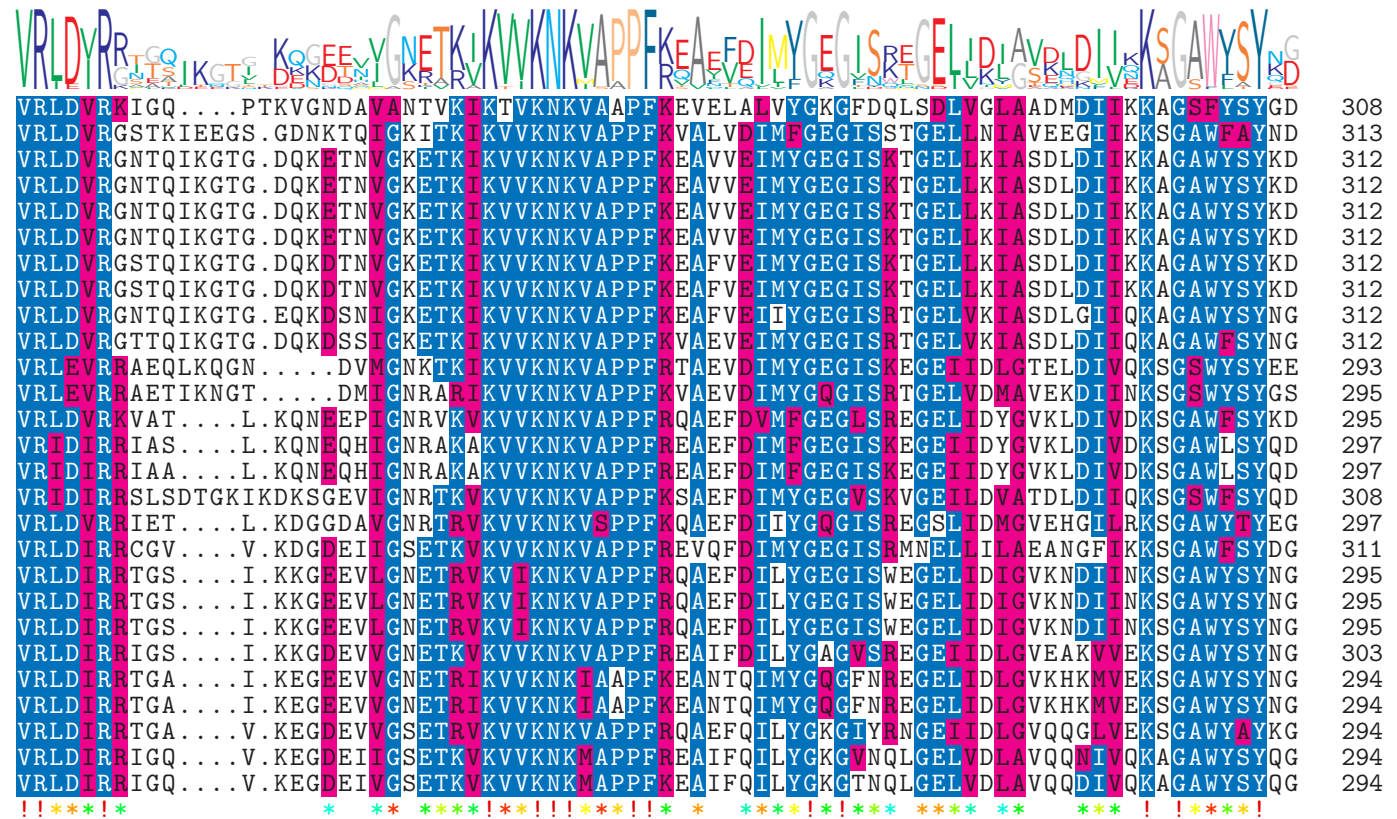


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

Sequence	Score
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EKIGQGAEAKAKNYLKEHQDVFEIDHKVRAAHGLDDSEVAETEETTASKTKAKAKKEEKAVETEEIELELED.....	387
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EKIGQGSSENAKKYLAHPFIDEIDKQVRSKFGLDIGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLAHPFIDEIDKQVRSKFGLDIGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLAHPFIDEIDKQVRSKFGLDIGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAES....KKDE....AAQVDS..VNEEVTLDLGDLEIEI	382
EKIGQGSSENAKKYLADNPEVFAEIDHQVRVQFGLIDGEEIAES....KKDE....VAQVDS..VNEEVTLDLGDLEIEI	382
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EKIGQGSSENAKKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIED	378
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KKLGGGRENSKAFLEKENPEIADEITTKAIQNSMGIEGMISGSEDDEG..EE.....	343
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NKIGQGKNNVIRYFENNQIAEIERNIREQLLTGTNGAVQ.I.EDEETEP.DLL..LE.....S.....	349

logo

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consensus

 non conserved
 similar
 ≥ 50% conserved