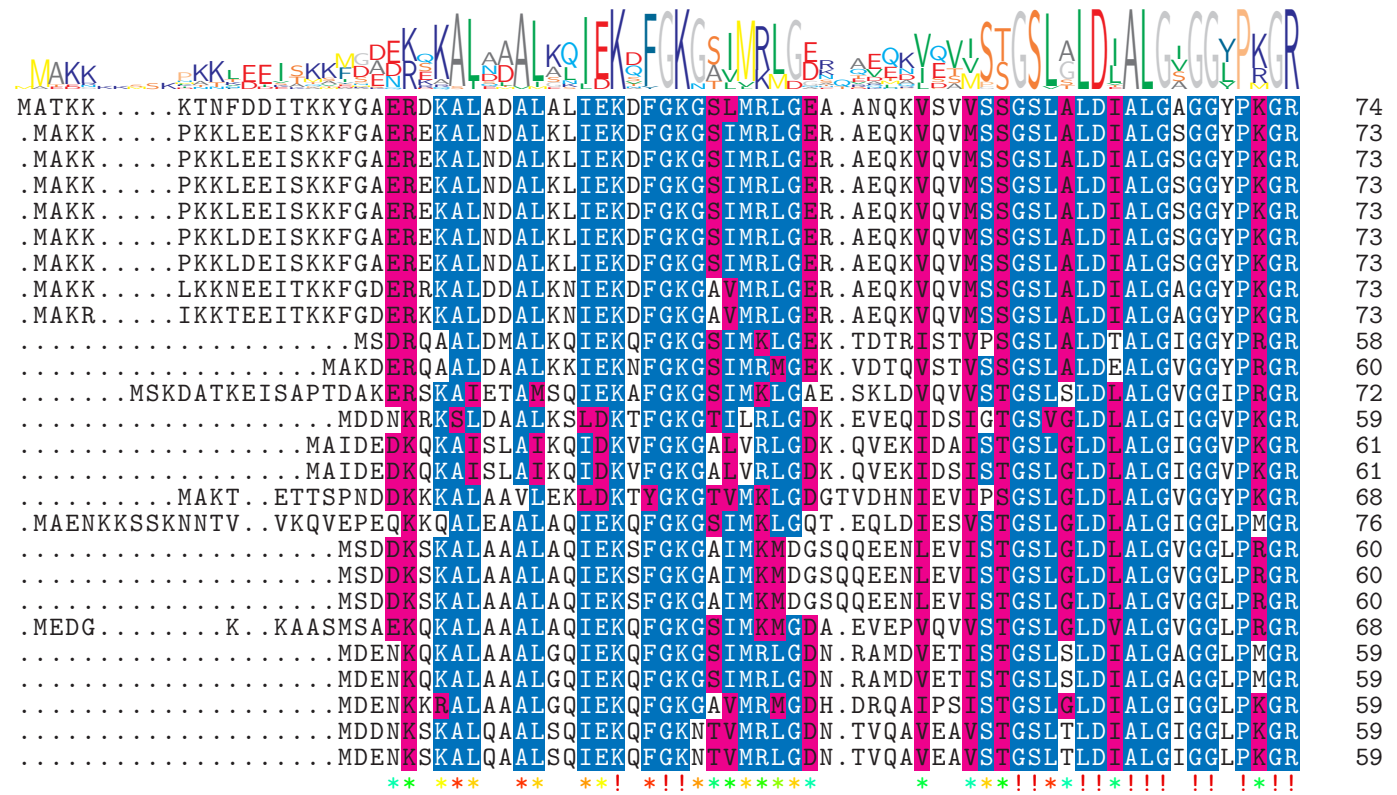


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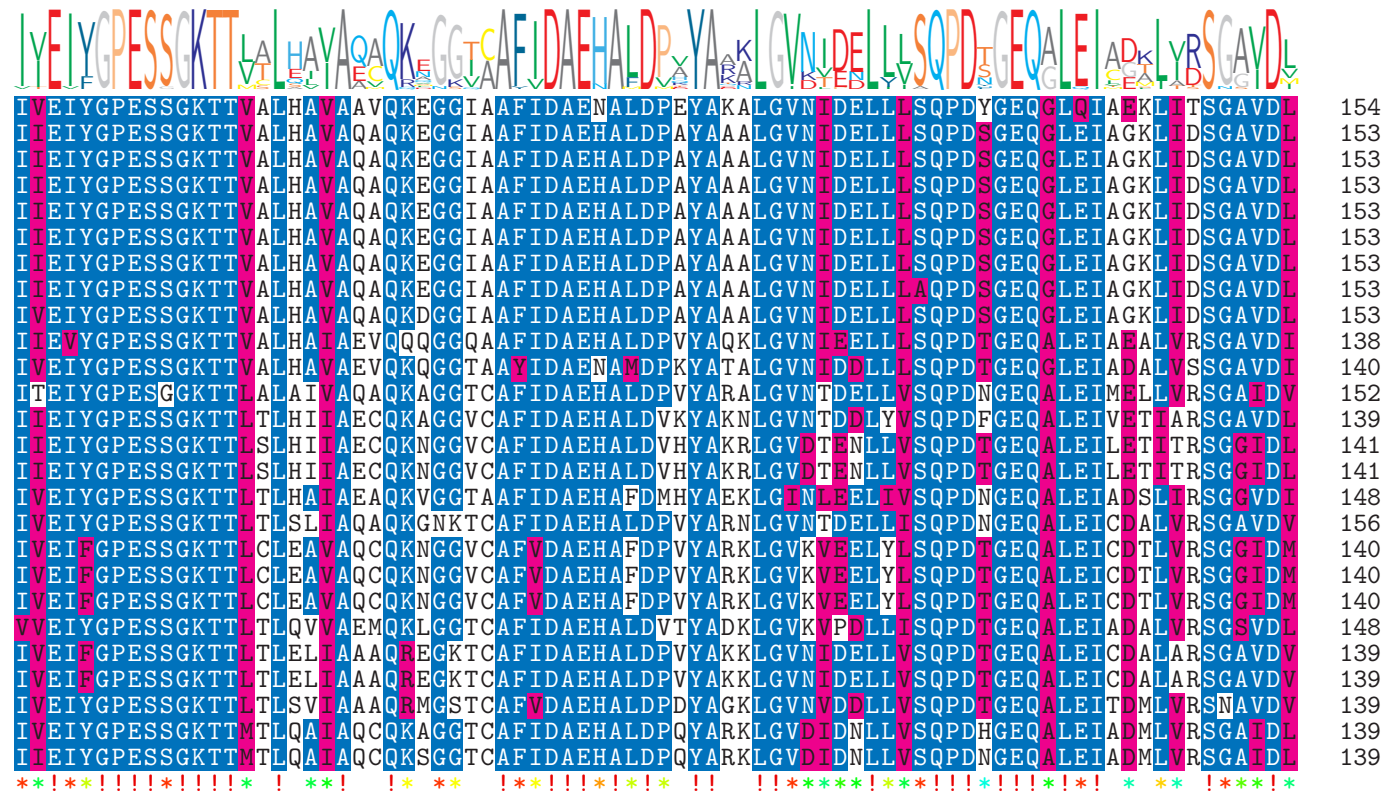
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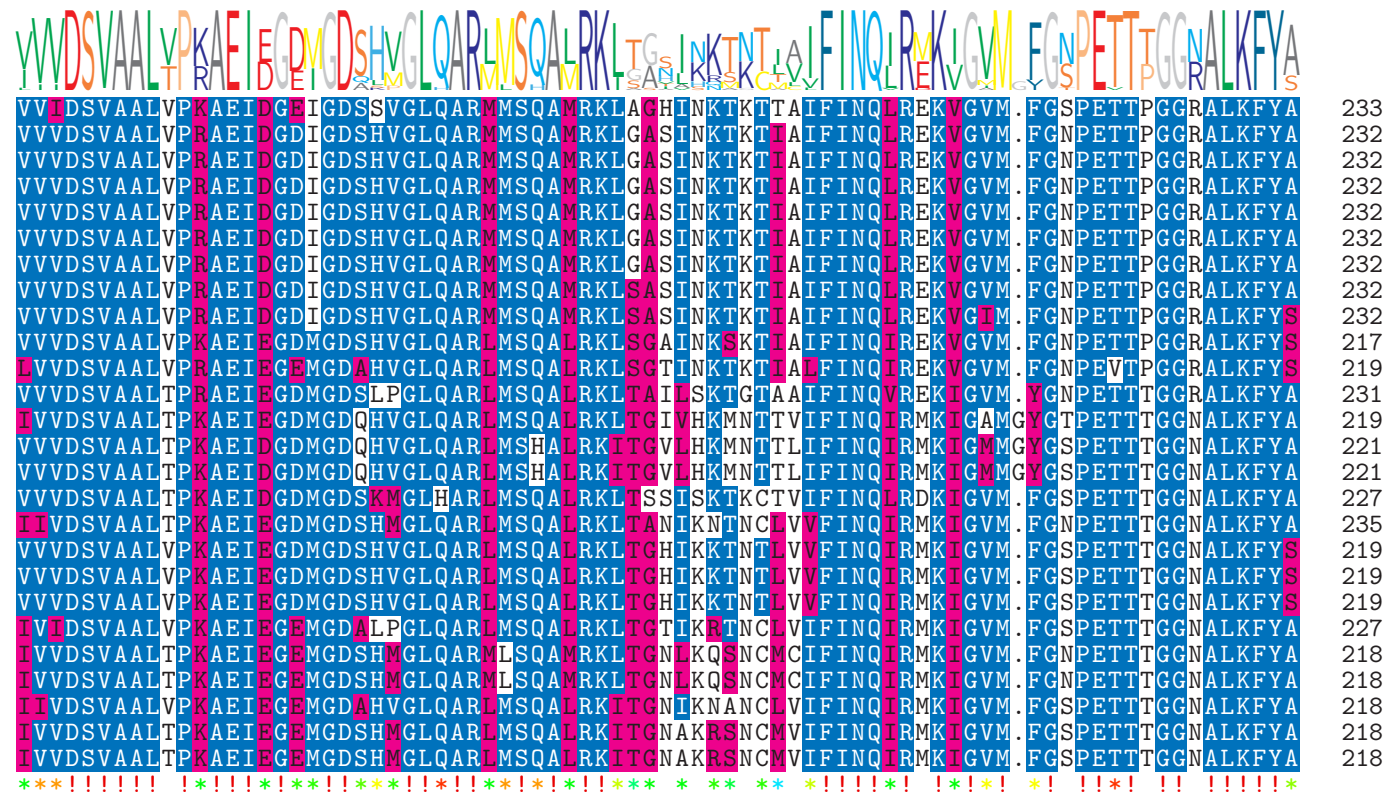
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consensus

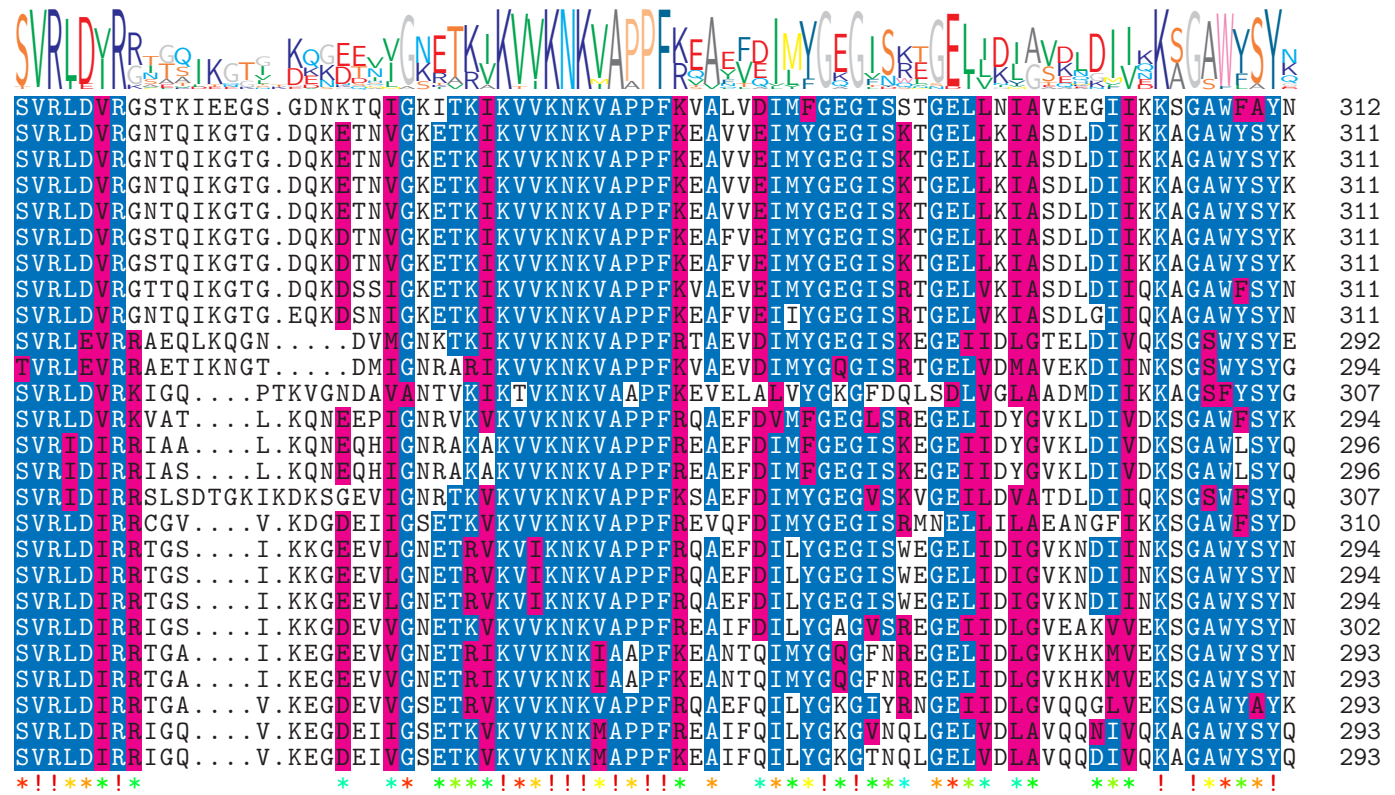
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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Sequence	Score
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DEKIGQGS ⁺ ENAK ⁺ KYLA ⁺ EHPE ⁺ IF ⁺ DE ⁺ ID ⁺ KQ ⁺ VR ⁺ SKF ⁺ GLIDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ ENAK ⁺ KYLA ⁺ EHPE ⁺ IF ⁺ DE ⁺ ID ⁺ KQ ⁺ VR ⁺ SKF ⁺ GLIDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ ENAK ⁺ KYLA ⁺ EHPE ⁺ IF ⁺ DE ⁺ ID ⁺ KQ ⁺ VR ⁺ SKF ⁺ GLIDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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GEKIGQGS ⁺ ENAK ⁺ RYLAD ⁺ HP ⁺ QL ⁺ FD ⁺ DE ⁺ ID ⁺ RV ⁺ VR ⁺ VKF ⁺ GLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGS ⁺ ENAK ⁺ KFLAD ⁺ NP ⁺ EI ⁺ FD ⁺ DE ⁺ ID ⁺ HK ⁺ VR ⁺ VQY ⁺ CLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGGGREN ⁺ AK ⁺ FLKEN ⁺ KD ⁺ IMLM ⁺ IQ ⁺ IREHY ⁺ CLDNNGVVQQQAETQEEL.EFE.E.....	348
SERIGGGREN ⁺ AK ⁺ NYLAD ⁺ HED ⁺ VE ⁺ DEV ⁺ LR ⁺ KV ⁺ RAAY ⁺ GISDVPEEDLPTTEDE.QI.NILP ⁺ DST.....EE.....	355
DERIGGGK ⁺ EKTI ⁺ AIY ⁺ A ⁺ ERP ⁺ EME ⁺ Q ⁺ EIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE..A.....	363
DKKLGGGREN ⁺ SKA ⁺ FLKEN ⁺ PEI ⁺ AD ⁺ EIT ⁺ KAI ⁺ QNSM ⁺ GI ⁺ TEGMI.....SGSEDDE..GE..E.....	343
DKKLGGGREN ⁺ AK ⁺ ALLKED ⁺ KALADE ⁺ ITL ⁺ KIK ⁺ ESIG ⁺ SNEEI.....MPLPDEP..LE..E.....	347
DKKLGGGREN ⁺ AK ⁺ ALLKED ⁺ KALADE ⁺ ITL ⁺ KIK ⁺ ESIG ⁺ SNEEI.....MPLPDEP..LE..E.....	347
DTKLGGGRD ⁺ TV ⁺ KE ⁺ LID ⁺ KN ⁺ PEI ⁺ AD ⁺ EI ⁺ AK ⁺ IREKLQAK.....	343
GTKIGGGK ⁺ NNAV ⁺ KLKEN ⁺ PEI ⁺ AD ⁺ KE ⁺ IQ ⁺ EIRNTLASNPNS.....VLKATASDSND..ESDVVEEFSEEEEL.....	372
GAKIGGGK ⁺ DN ⁺ VR ⁺ VWLKEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AK ⁺ IRALN ⁺ GVEMHI.....TEGT.....Q..DET ⁺ DGERPEE.....	348
GAKIGGGK ⁺ DN ⁺ VR ⁺ VWLKEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AK ⁺ IRALN ⁺ GVEMHI.....TEGT.....Q..DET ⁺ DGERPEE.....	348
GAKIGGGK ⁺ DN ⁺ VR ⁺ VWLKEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AK ⁺ IRALN ⁺ GVEMHI.....TEGT.....Q..DET ⁺ DGERPEE.....	348
GERIGGGRD ⁺ NC ⁺ REFL ⁺ RNAEL ⁺ LARE ⁺ EN ⁺ KV ⁺ REHL ⁺ GVTPMG....AV.TLAEVEVE..D.....	352
GDKIGGGK ⁺ ANACK ⁺ YLKEN ⁺ PEI ⁺ AK ⁺ TL ⁺ DKK ⁺ IREMLLN ⁺ PNEM....QLIAETSSAAD..DVEFGAVPEEF.....	354
GDKIGGGK ⁺ ANACK ⁺ YLKEN ⁺ PEI ⁺ AK ⁺ TL ⁺ DKK ⁺ IREMLLN ⁺ PNEM....QLIAETSSAAD..DVEFGAVPEEF.....	354
GSKIGGGK ⁺ ANAAK ⁺ YLEEN ⁺ PEI ⁺ GQ ⁺ EIE ⁺ QQI ⁺ RAKL ⁺ YTSAN....TKATPVAEDLA..DADI.....	347
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GNKIGGGK ⁺ NNVIR ⁺ YFEEN ⁺ PNQI ⁺ AEI ⁺ ERN ⁺ IREQLLT ⁺ TGTN....GAVQ.I.EDEE..EPDLL..LES.....	349

logo

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consensus

 non conserved
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 ≥ 50% conserved