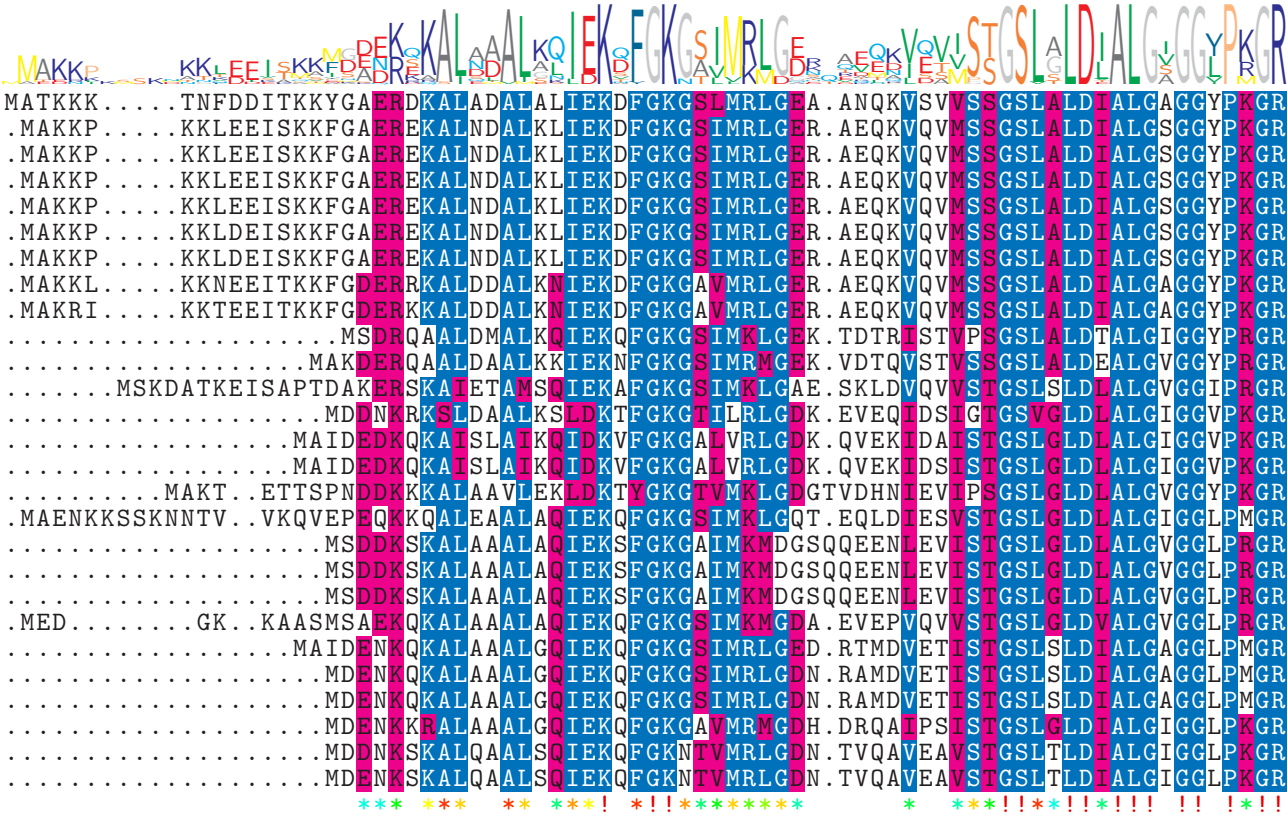


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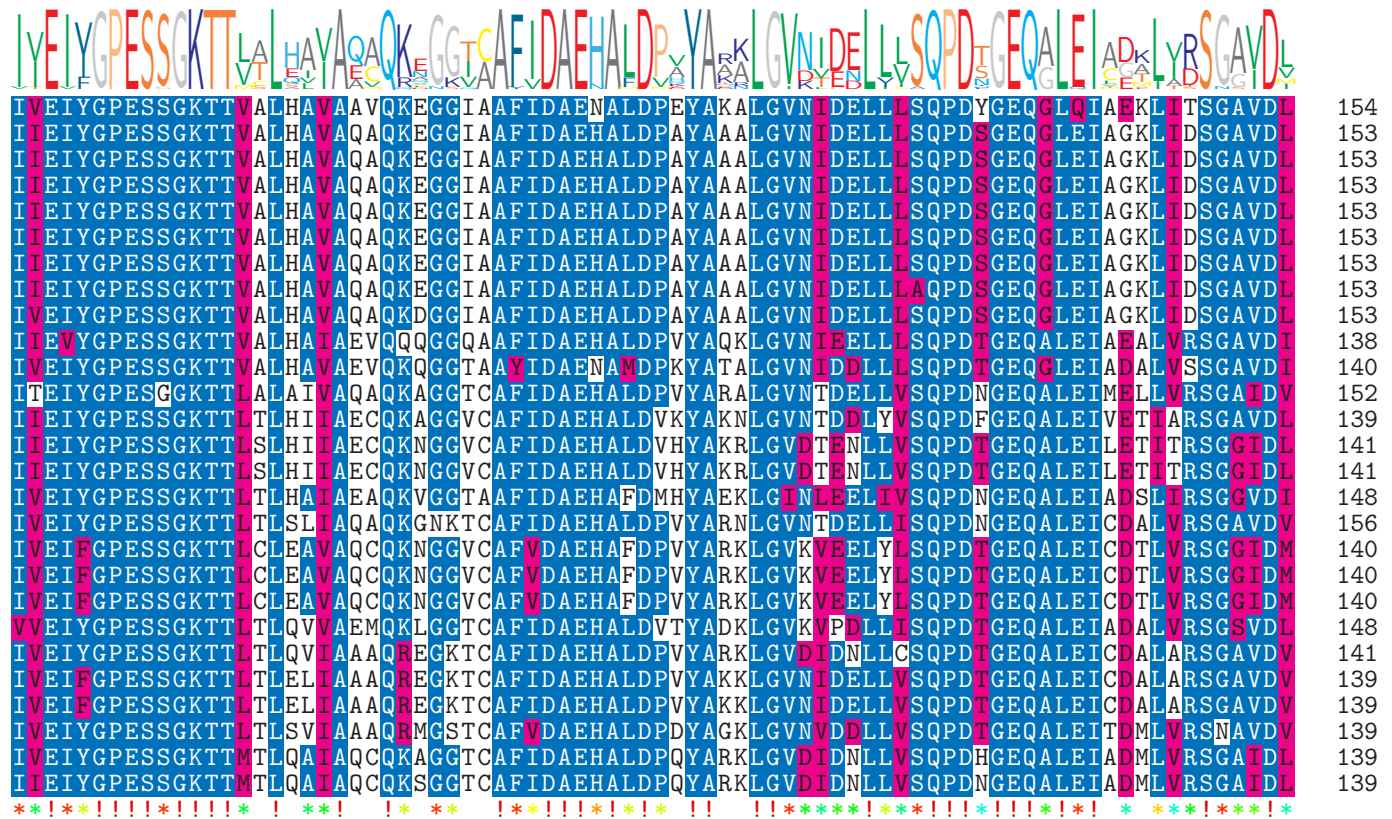


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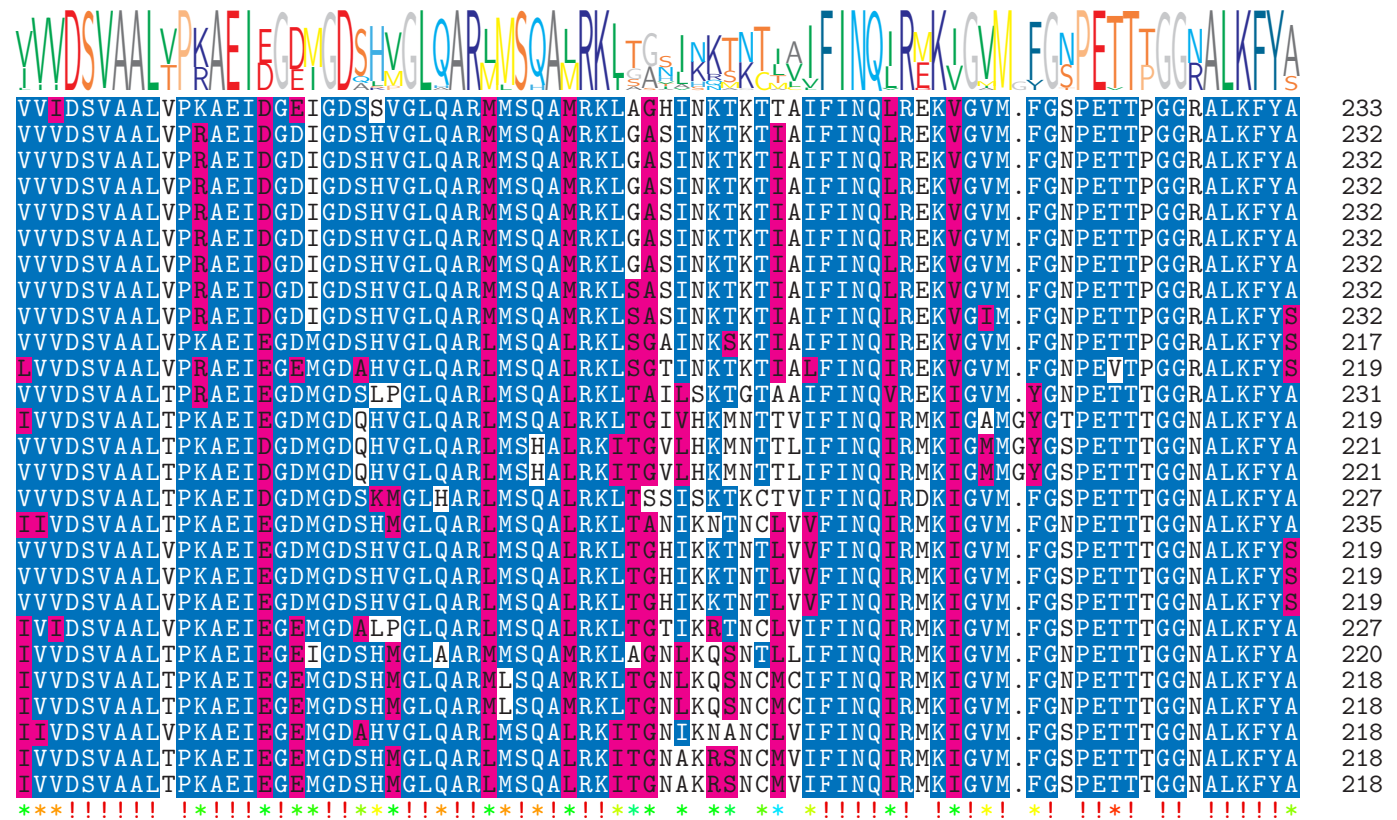
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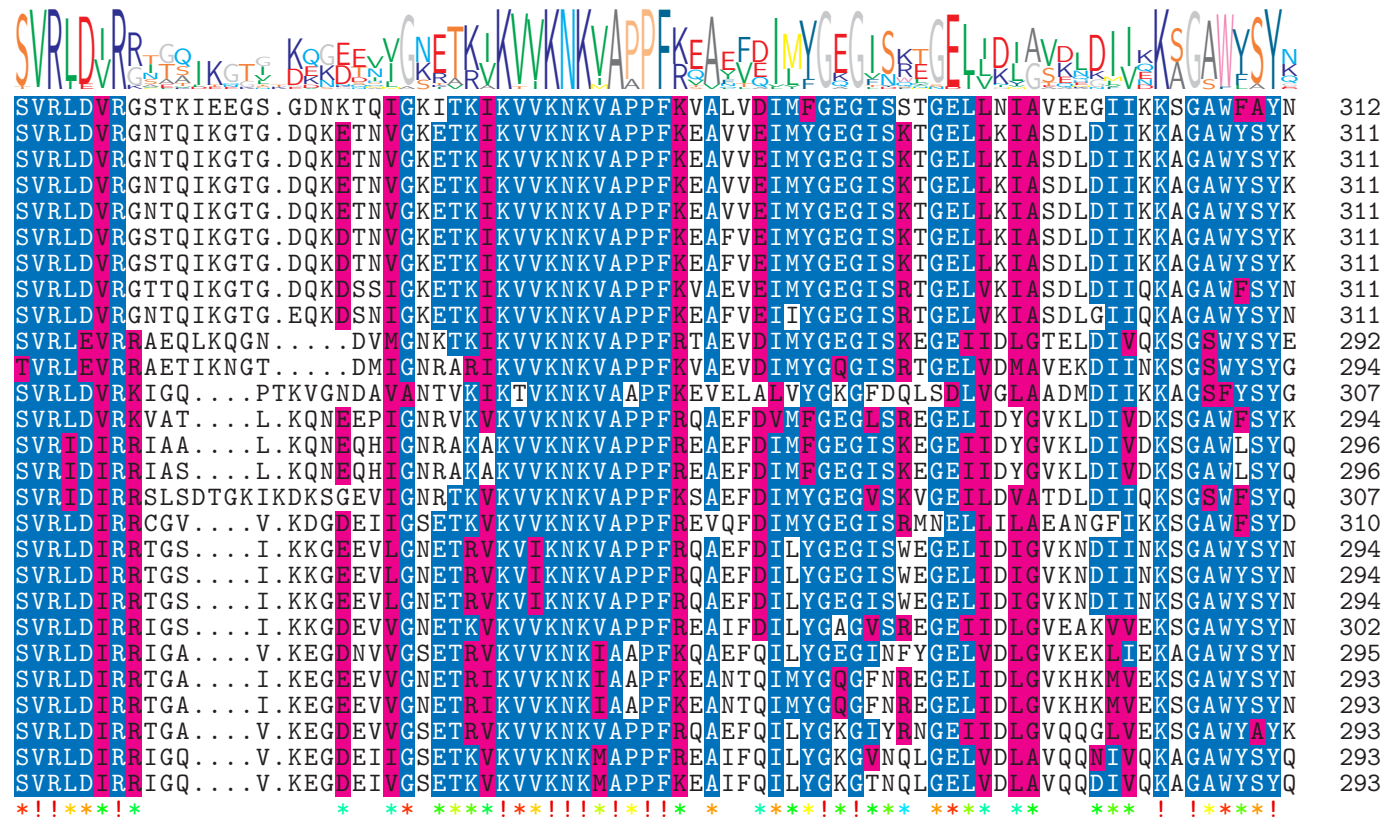
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consensus

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Sequence	Score
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DEKIGGGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EIF ⁺ DEID ⁺ DKQ ⁺ VR ⁺ SKF ⁺ CL ⁺ IDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGGGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AD ⁺ NPE ⁺ VF ⁺ AEID ⁺ DHQ ⁺ VR ⁺ VQF ⁺ CL ⁺ IDGEEAAE....SKKDE....VAQVDS..VNEEVTLDLGDELEIE	381
DEKIGGGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AD ⁺ NPE ⁺ VF ⁺ DEID ⁺ DHQ ⁺ VR ⁺ VQF ⁺ CL ⁺ IDGEEIAE....SKKDE....AAQVDS..VNEEVTLDLGDELEIE	381
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GEKIGGGS ⁺ EN ⁺ AK ⁺ KFL ⁺ AD ⁺ NPE ⁺ IF ⁺ DDI ⁺ DHK ⁺ VR ⁺ VQY ⁺ CL ⁺ IEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EE ⁺ RL ⁺ LG ⁺ GG ⁺ RE ⁺ NA ⁺ KQ ⁺ FL ⁺ KE ⁺ NKD ⁺ IMLM ⁺ IQE ⁺ QIREHY ⁺ CL ⁺ DMNGVVQQQAEEETQEEL.EFE.E.....	348
SERIGGG ⁺ RE ⁺ NA ⁺ KNYL ⁺ AD ⁺ HED ⁺ VE ⁺ DEV ⁺ RLK ⁺ V ⁺ RAAY ⁺ CI ⁺ SDVPEEDLPTTEDE.QI.NILPDDST....EE.....	355
DERIGGG ⁺ KE ⁺ KTI ⁺ AYI ⁺ AE ⁺ RP ⁺ EME ⁺ QE ⁺ IRDR ⁺ VMAAIRAGNAGEAPALAPAPAA.PEAAEA.....	363
DKKL ⁺ GG ⁺ GRE ⁺ NSK ⁺ AF ⁺ LKEN ⁺ PEI ⁺ AD ⁺ EITKAT ⁺ IQNSM ⁺ CI ⁺ EGMISGSEDDE..GE.E.....	343
DKKL ⁺ GG ⁺ GRE ⁺ NA ⁺ KALL ⁺ KED ⁺ KAL ⁺ ADEI ⁺ TLK ⁺ IKESIG ⁺ SNEEIMPLPDEP..LE.EME.....	347
DKKL ⁺ GG ⁺ GRE ⁺ NA ⁺ KALL ⁺ KED ⁺ KAL ⁺ ADEI ⁺ TLK ⁺ IKESIG ⁺ SNEEIMPLPDEP..LE.EME.....	347
DTKL ⁺ GG ⁺ GRD ⁺ TV ⁺ KEL ⁺ IKD ⁺ NPE ⁺ LAE ⁺ ELE ⁺ AKI ⁺ REKLQAK.....	343
GTKIGGG ⁺ KNN ⁺ AVK ⁺ WL ⁺ KEN ⁺ PEI ⁺ AD ⁺ KL ⁺ EQE ⁺ IRNTL ⁺ ASNPNSVLKATAS..D...SNDES DVEEFSEEE..L.....	372
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GAKIGGG ⁺ KDN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AKI ⁺ RALN ⁺ GVEMHITEGTQDE..TD.....GERPEE.....	348
GAKIGGG ⁺ KDN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AKI ⁺ RALN ⁺ GVEMHITEGTQDE..TD.....GERPEE.....	348
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GNKIGGG ⁺ KNN ⁺ VIRY ⁺ FEEN ⁺ NCI ⁺ AEI ⁺ ERN ⁺ IREQLLTGTNGAV..Q...I..EDEEPPDLLS.....	349

logo

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consensus

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