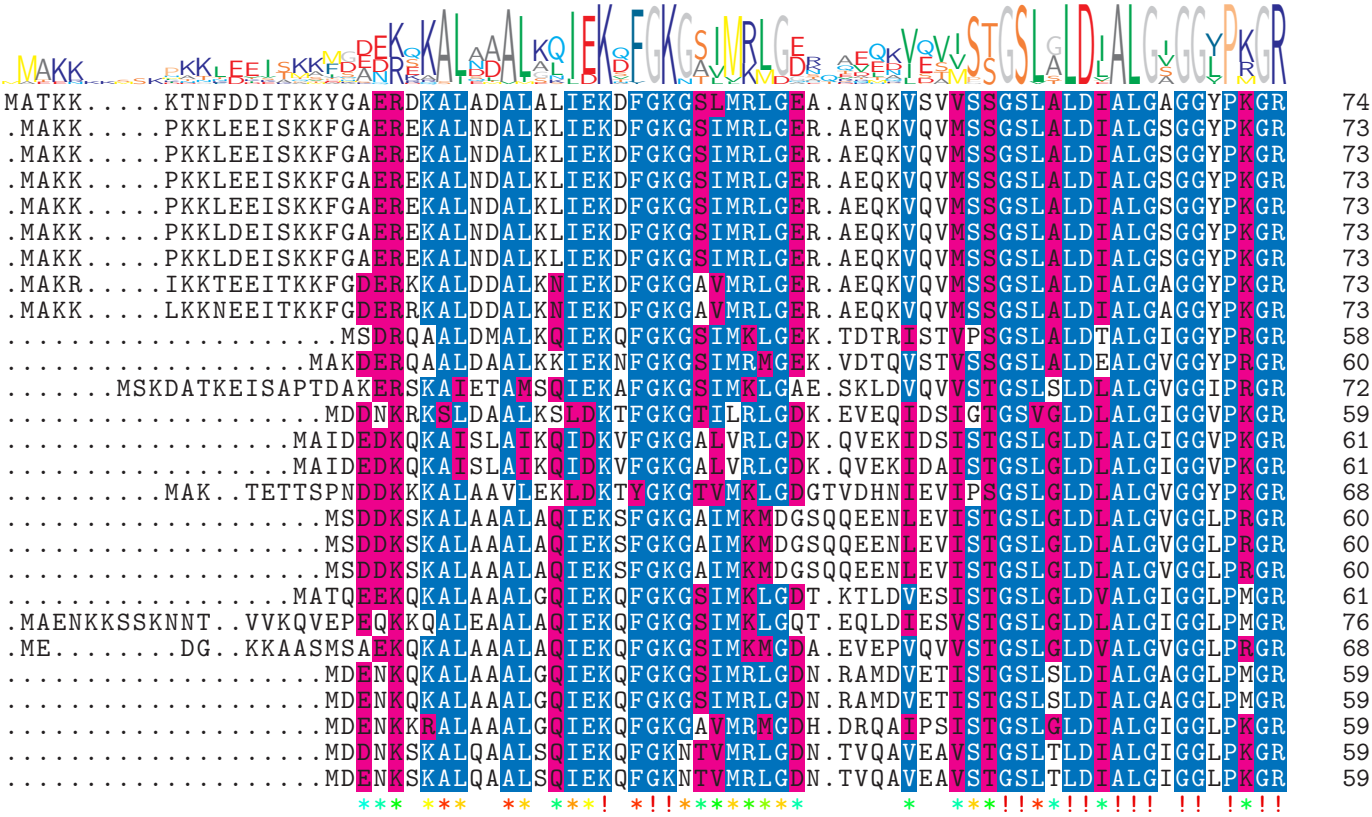


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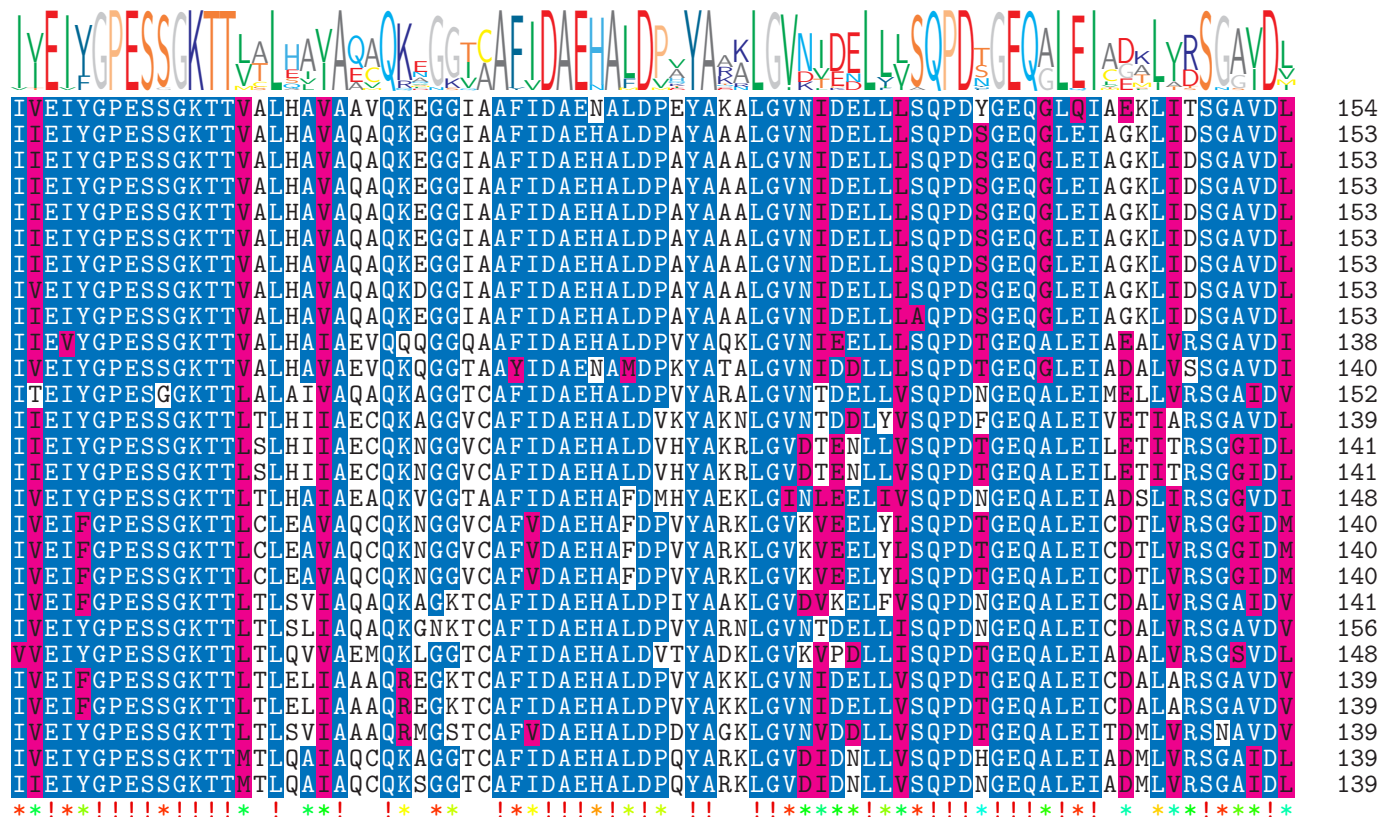
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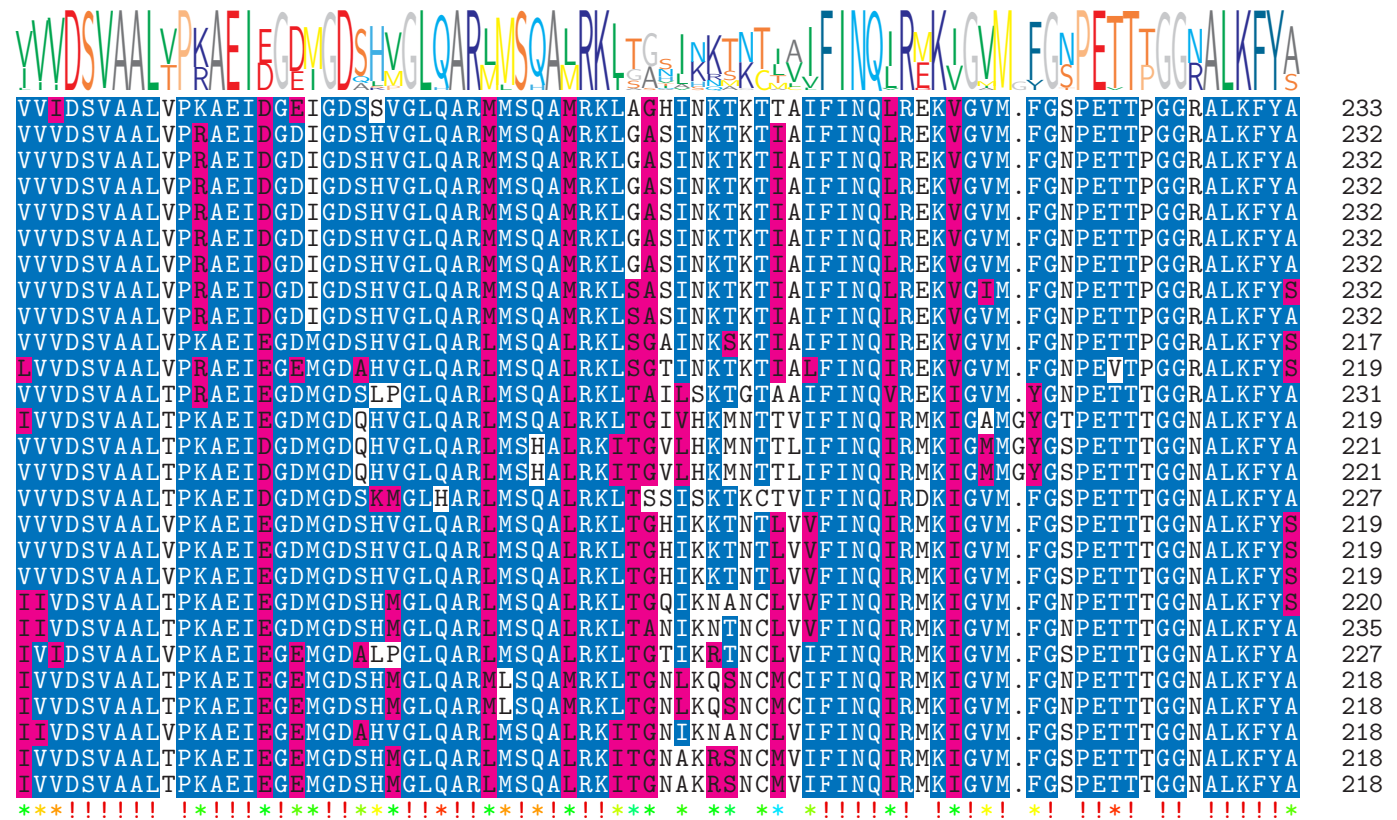
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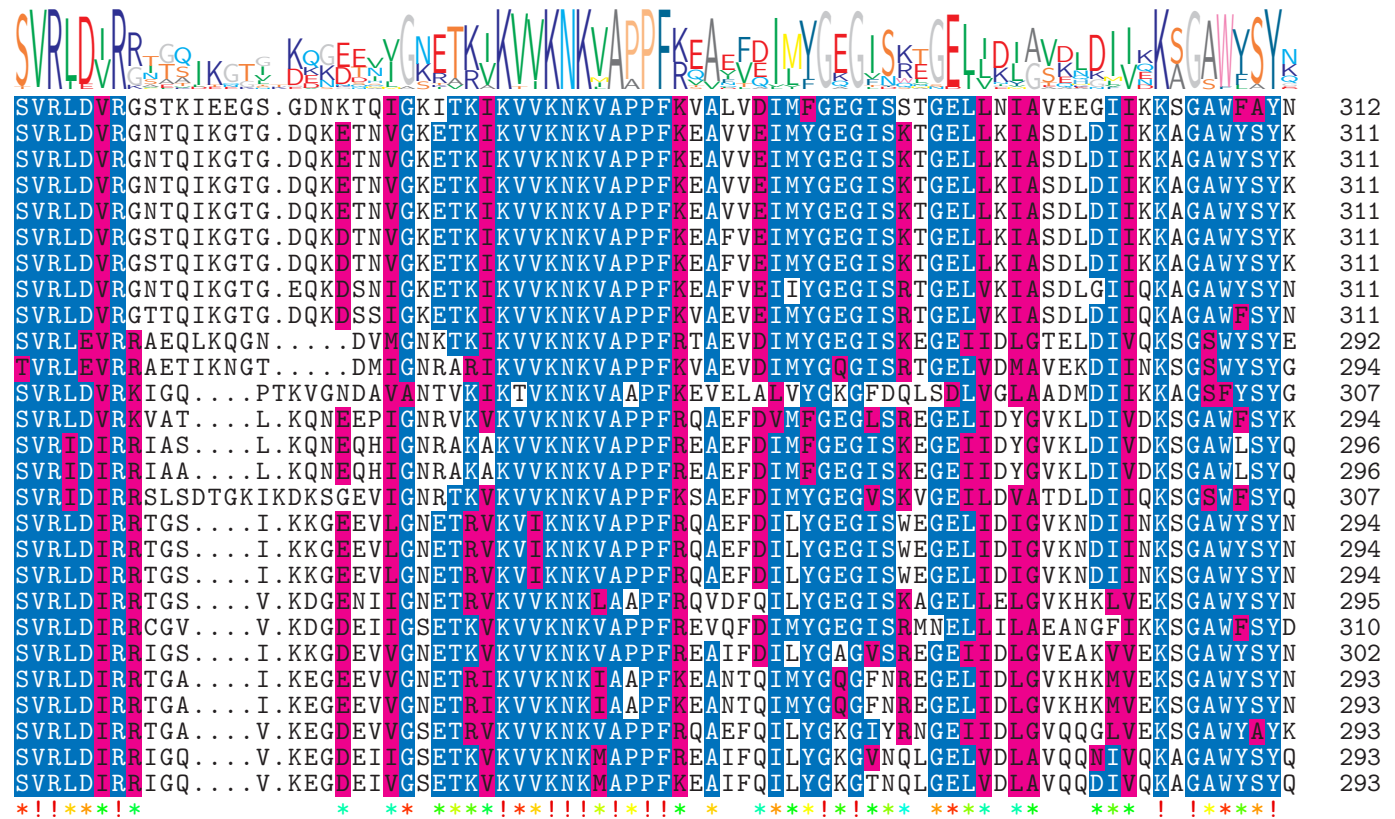
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Sequence	Score
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DEKIGQGSEN ⁺ AK ⁺ KYL ⁺ AEHP ⁺ EI ⁺ FDEI ⁺ DKQV ⁺ RSKF ⁺ CL ⁺ IDGEEVSEQD.TENKKDE....PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSEN ⁺ AK ⁺ KYL ⁺ AEHP ⁺ EI ⁺ FDEI ⁺ DKQV ⁺ RSKF ⁺ CL ⁺ IDGEEVSEQD.TENKKDE....PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSEN ⁺ AK ⁺ KYL ⁺ AEHP ⁺ EI ⁺ FDEI ⁺ DKQV ⁺ RSKF ⁺ CL ⁺ IDGEEVSEQD.TENKKDE....PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSEN ⁺ AK ⁺ KYL ⁺ ADNP ⁺ EI ⁺ FDEI ⁺ DHQR ⁺ VRVQ ⁺ FL ⁺ IDGEEIAES....KKDE....AAQVDS..VNEEVPLDLGDELEI	380
DEKIGQGSEN ⁺ AK ⁺ KYL ⁺ ADNP ⁺ EI ⁺ FDEI ⁺ DHQR ⁺ VRVQ ⁺ FL ⁺ IDGEEAAES....KKDE....VAQVDS..VNEEVPLDLGDELEI	380
GEKIGQGSEN ⁺ AK ⁺ KFL ⁺ ADNP ⁺ EI ⁺ FDDI ⁺ DHKV ⁺ RVQY ⁺ CL ⁺ IEEDNTEEKQ.SSKEK.....ETDEK..ADKNLVLELDDTIEL	381
GEKIGQGSEN ⁺ AK ⁺ KRYL ⁺ ADHP ⁺ QL ⁺ FDEI ⁺ DRKV ⁺ RVKF ⁺ CL ⁺ LEESEEEESAM.AVASE.....E.....TDDLALDLNGLIEI	376
EE ⁺ RL ⁺ GGGREN ⁺ AK ⁺ QFL ⁺ KEN ⁺ KDI ⁺ MLMI ⁺ QE ⁺ QIREHY ⁺ CL ⁺ DMNGVVQQQA.EETQEEL.EFE.E.....	348
SERIGGGREN ⁺ AK ⁺ NYL ⁺ ADHED ⁺ VE ⁺ DEV ⁺ RLKV ⁺ RAAY ⁺ CI ⁺ SDVPEEDLPT.TEDE.QI.NILPDDST....EE.....	355
DERIGGQKEK ⁺ TIAY ⁺ IA ⁺ ER ⁺ PEME ⁺ QE ⁺ IRDRV ⁺ MAAIRAGNAGEAPALAPAPAAPEAAEA.....	363
DKKLGGGREN ⁺ SKA ⁺ FL ⁺ KEN ⁺ PEI ⁺ ADEIT ⁺ TKAT ⁺ IQNSM ⁺ CI ⁺ EGMISGS..EDDEGE..E.....	343
DKKLGGGREN ⁺ AK ⁺ ALL ⁺ KED ⁺ KALADEIT ⁺ TLKIK ⁺ ESIG ⁺ SNEEIMPL..PDEPLE..E.....	347
DKKLGGGREN ⁺ AK ⁺ ALL ⁺ KED ⁺ KALADEIT ⁺ TLKIK ⁺ ESIG ⁺ SNEEIMPL..PDEPLE..E.....	347
DTKLGGGRD ⁺ TVKEL ⁺ IKD ⁺ NPEL ⁺ AE ⁺ ELE ⁺ AKI ⁺ REKLQAK.....	343
GAKIGQGKD ⁺ NVRV ⁺ WLKEN ⁺ PEIS ⁺ DEID ⁺ AKI ⁺ RALN ⁺ GVEMHITEG..T...QD...ETDGE.RPEE.....	348
GAKIGQGKD ⁺ NVRV ⁺ WLKEN ⁺ PEIS ⁺ DEID ⁺ AKI ⁺ RALN ⁺ GVEMHITEG..T...QD...ETDGE.RPEE.....	348
GAKIGQGKD ⁺ NVRV ⁺ WLKEN ⁺ PEIS ⁺ DEID ⁺ AKI ⁺ RALN ⁺ GVEMHITEG..T...QD...ETDGE.RPEE.....	348
GEKIGQGKAN ⁺ SMK ⁺ WL ⁺ NENI ⁺ EKS ⁺ DELE ⁺ ARL ⁺ RAELVANPEQALM..ADIEQSENNTESSEDFE.....	354
GTKIGQGKKN ⁺ NAV ⁺ KWLKEN ⁺ PEI ⁺ AD ⁺ KL ⁺ EQEIR ⁺ NTL ⁺ ASNPN ⁺ SVLK..ATASDSND ⁺ ESDVE.EFSE....EEL.....	372
GE ⁺ RIGGRDNC ⁺ REFL ⁺ REN ⁺ AE ⁺ LARE ⁺ ENK ⁺ VREHL ⁺ CV ⁺ TPMGAV..TLAEEVEED.....	352
GD ⁺ KIGQGKAN ⁺ ACKY ⁺ LKEN ⁺ PEI ⁺ AKT ⁺ LD ⁺ KKL ⁺ REMLLN ⁺ PNEMQLI..AETSSAAD ⁺ VEFGAVPE.....EF.....	354
GD ⁺ KIGQGKAN ⁺ ACKY ⁺ LKEN ⁺ PEI ⁺ AKT ⁺ LD ⁺ KKL ⁺ REMLLN ⁺ PNEMQLI..AETSSAAD ⁺ VEFGAVPE.....EF.....	354
GSKIGQGKAN ⁺ AAKY ⁺ LEEN ⁺ PEI ⁺ GQ ⁺ EIE ⁺ QQI ⁺ IRAKLL ⁺ Y ⁺ TSANTKA..TPVAEDLADADI.....	347
GNKIGQGKKN ⁺ NVIR ⁺ YLEEN ⁺ PEI ⁺ QI ⁺ STEI ⁺ EAVI ⁺ REQLLT ⁺ KASDQTA..A.HDETEE ⁺ PDLL..ES.....	349
GNKIGQGKKN ⁺ NVIR ⁺ YLEEN ⁺ PEI ⁺ QI ⁺ STEI ⁺ EARN ⁺ IREQLLT ⁺ TGTNGAV..Q.I.EDEE ⁺ PDLL..LE.....S.....	349

logo

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved