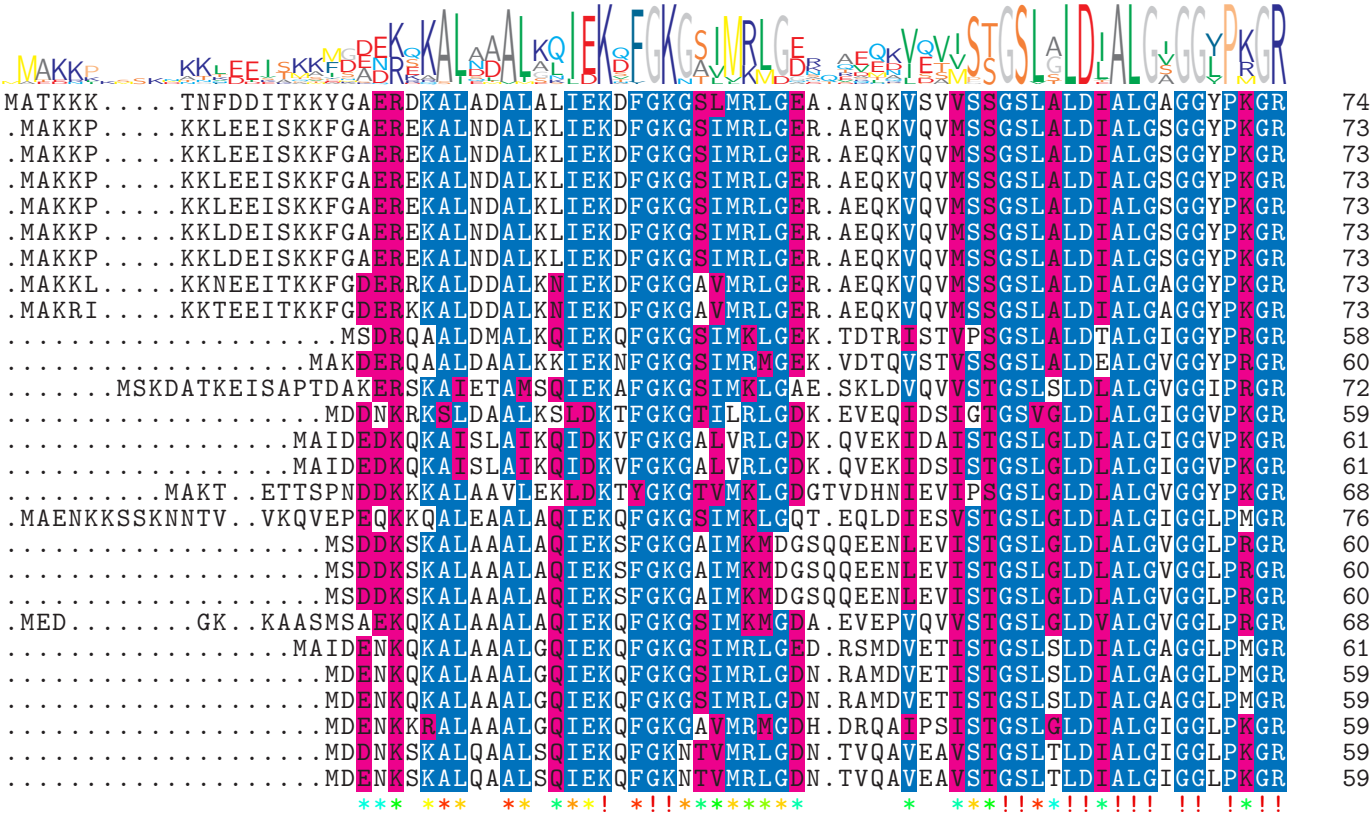


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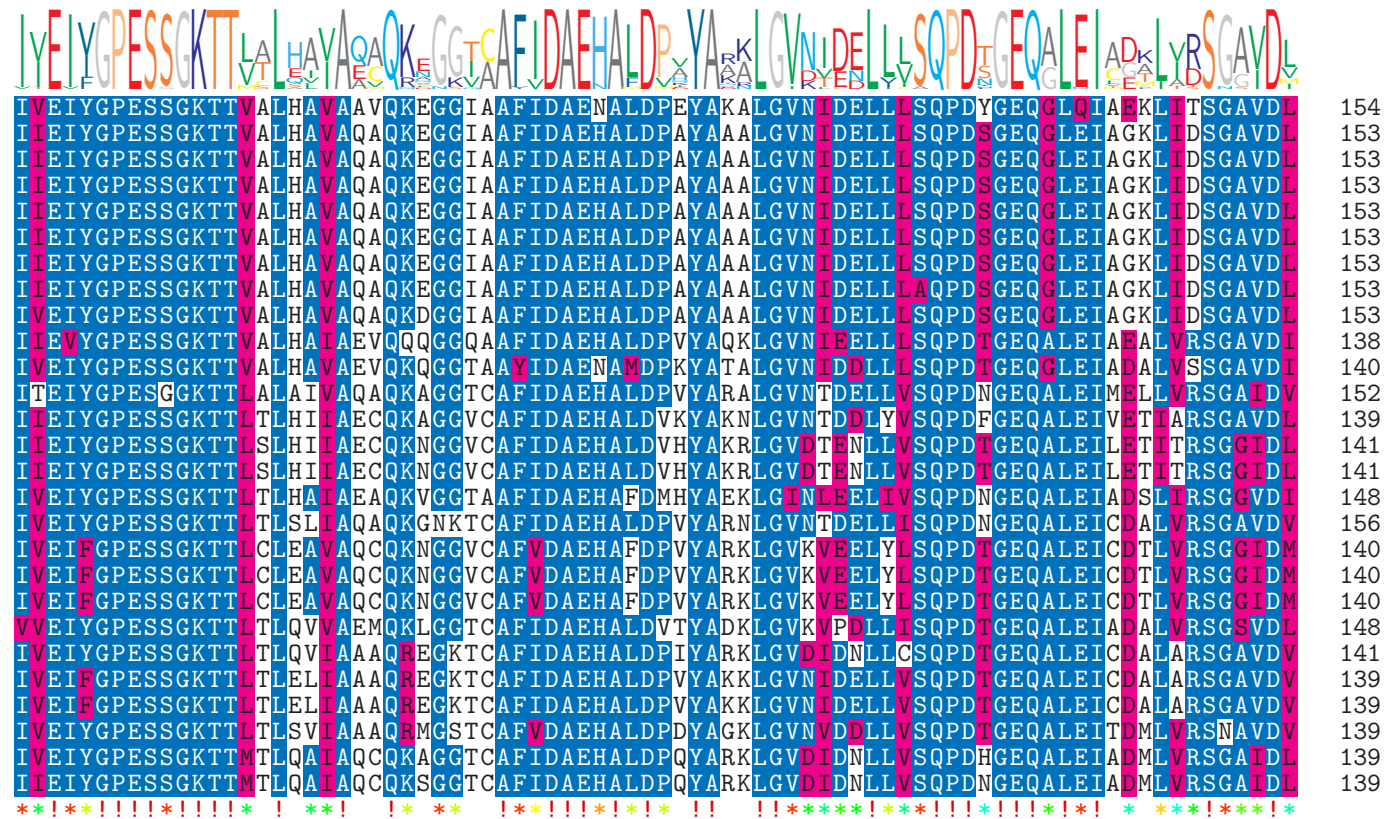
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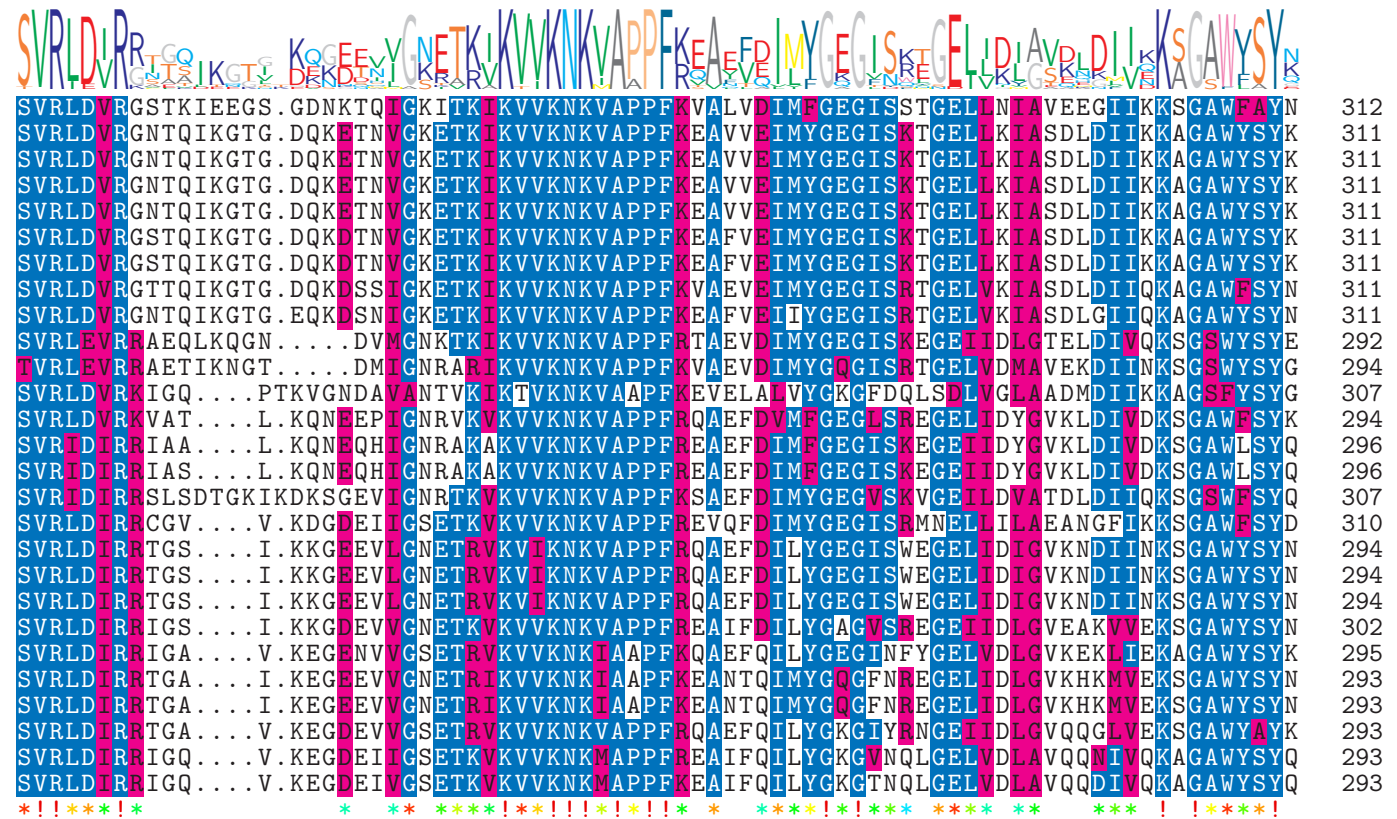
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consensus

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Sequence	Score
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DEKIGQGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PEIF ⁶ DEID ⁷ KQ ⁸ VR ⁹ SKF ¹⁰ GL ¹¹ IDGEEVSEQDTENKKDE ¹² ...PKKEEA ¹³ ..VNEEVPLDLGDELEIE	385
DEKIGQGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PEIF ⁶ DEID ⁷ KQ ⁸ VR ⁹ SKF ¹⁰ GL ¹¹ IDGEEVSEQDTENKKDE ¹² ...PKKEEA ¹³ ..VNEEVPLDLGDELEIE	385
DEKIGQGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PEIF ⁶ DEID ⁷ KQ ⁸ VR ⁹ SKF ¹⁰ GL ¹¹ IDGEEVSEQDTENKKDE ¹² ...PKKEEA ¹³ ..VNEEVPLDLGDELEIE	385
DEKIGQGS ¹ EN ² AK ³ KYLA ⁴ DN ⁵ PEIF ⁶ DEID ⁷ HQ ⁸ VR ⁹ VQF ¹⁰ GL ¹¹ IDGEEIAES ¹² ...KKDE ¹³ ...AAQV ¹⁴ DS ¹⁵ ..VNEEVTLDLGDLEIE	381
DEKIGQGS ¹ EN ² AK ³ KYLA ⁴ DN ⁵ PEVFA ⁶ EID ⁷ HQ ⁸ VR ⁹ VQF ¹⁰ GL ¹¹ IDGEEAAES ¹² ...KKDE ¹³ ...VAQV ¹⁴ DS ¹⁵ ..VNEEVTLDLGDLEIE	381
GEKIGQGS ¹ EN ² AK ³ RYLA ⁴ DHPQLF ⁵ DEID ⁶ RK ⁷ VR ⁸ VKF ⁹ GL ¹⁰ LEESEESAMAVASE ¹¹E ¹²TDDLALDLNGIEIE	377
GEKIGQGS ¹ EN ² AK ³ KFLA ⁴ DN ⁵ PEIF ⁶ DD ⁷ ID ⁸ HK ⁹ VR ¹⁰ VQY ¹¹ GL ¹² IEEDNTEEKQSSKEK ¹³ETDE ¹⁴ K ¹⁵ ..ADKNLVLELDDTIELE	382
EE ¹ RL ² GG ³ RENA ⁴ KQFL ⁵ KN ⁶ KDIMLMIQE ⁷ QIREHY ⁸ GL ⁹ DNNGVVQQAEETQEEL ¹⁰ .EFE ¹¹ .E ¹²	348
SERIGQGS ¹ RENA ² KNYLA ³ DH ⁴ ED ⁵ VE ⁶ DE ⁷ VR ⁸ LK ⁹ VRAAY ¹⁰ GIS ¹¹ SDVPEEDLPTTEDE ¹² .QI ¹³ .NILPDDST ¹⁴EE ¹⁵	355
DERIGQG ¹ KE ² KTIAY ³ IA ⁴ ERP ⁵ MEQE ⁶ FEIRDRVMAAIRAGNAGEAPALAPAPAAPEAA ⁷ ...EA ⁸	363
DKKLGGGREN ¹ SKA ² FL ³ IKEN ⁴ PEIA ⁵ DEIT ⁶ KA ⁷ IQNSM ⁸ GI ⁹ EGM ¹⁰ISGSEDDEG ¹¹EE ¹²	343
DKKLGGGREN ¹ AK ² ALL ³ KED ⁴ KALADEIT ⁵ TL ⁶ KIKESIG ⁷ SNEE ⁸IMPLPDEPL ⁹EEME ¹⁰	347
DKKLGGGREN ¹ AK ² ALL ³ KED ⁴ KALADEIT ⁵ TL ⁶ KIKESIG ⁷ SNEE ⁸IMPLPDEPL ⁹EEME ¹⁰	347
DTKLGGGRD ¹ TV ² KEL ³ IK ⁴ DN ⁵ PELA ⁶ E ⁷ LEAK ⁸ IREKLQAK ⁹	343
GTKIGQG ¹ KNN ² AVK ³ WL ⁴ KEN ⁵ PEIA ⁶ DK ⁷ LEQEIRNTLASNP ⁸ N.....SVLKATASDSN ⁹ ...DES ¹⁰ DVEEFSEEL ¹¹	372
GAKIGGQKDN ¹ VR ² VWL ³ KEN ⁴ PEIS ⁵ DEIDAK ⁶ IRALNG ⁷ VMH ⁸ITEG ⁹T ¹⁰ ...QDET ¹¹ DGERPEE ¹²	348
GAKIGGQKDN ¹ VR ² VWL ³ KEN ⁴ PEIS ⁵ DEIDAK ⁶ IRALNG ⁷ VMH ⁸ITEG ⁹T ¹⁰ ...QDET ¹¹ DGERPEE ¹²	348
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GERIGGGRD ¹ NC ² REFL ³ RENA ⁴ E ⁵ LAREIENK ⁶ VREHL ⁷ GVTPM ⁸GAVTLAEE ⁹ .VE ¹⁰ ...ED ¹¹	352
GEKIGGQKAN ¹ ATAWL ² KN ³ PETAK ⁴ EIEKKVRELLSNPN ⁵STPDFSV ⁶ ...D ⁷ ...DSEGVAETNEDF ⁸	353
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GSKIGGQKANA ¹ AKYLEN ² PEIG ³ QIEEQ ⁴ IRAKLL ⁵ VTSA ⁶NTKATPVAEDL ⁷ ...ADADI ⁸	347
GNKIGGQKNN ¹ VIRYLEN ² PEIQISTEIE ³ EAVIREQLLTKAS ⁴DQTA ⁵ A.HDETE ⁶ ...EED ⁷ LLLES ⁸	349
GNKIGGQKNN ¹ VIRYLEN ² PEIQIAEIE ³ ERNIREQLLTGT ⁴NGAVQ ⁵ .I.EDE ⁶ ...EED ⁷ LLLES ⁸	349

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- X non conserved
- X similar
- X ≥ 50% conserved