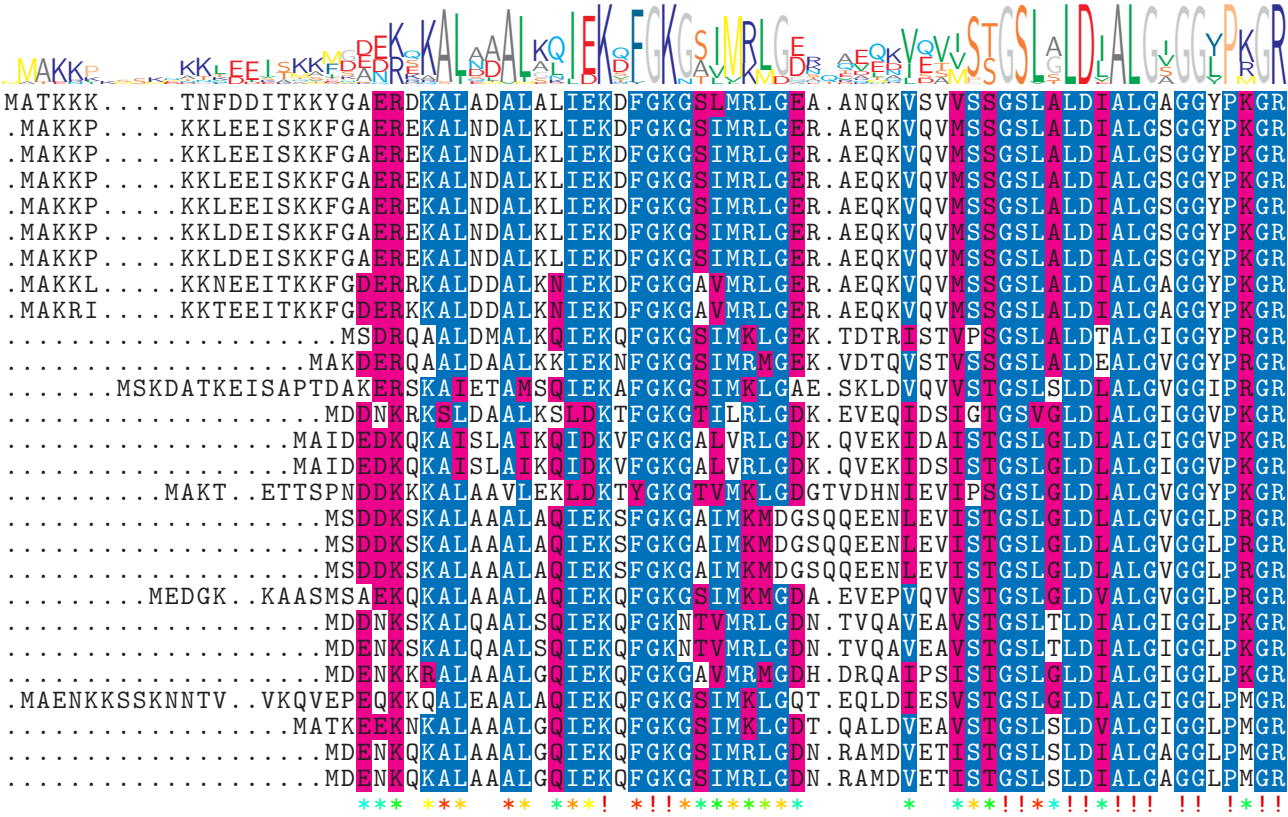


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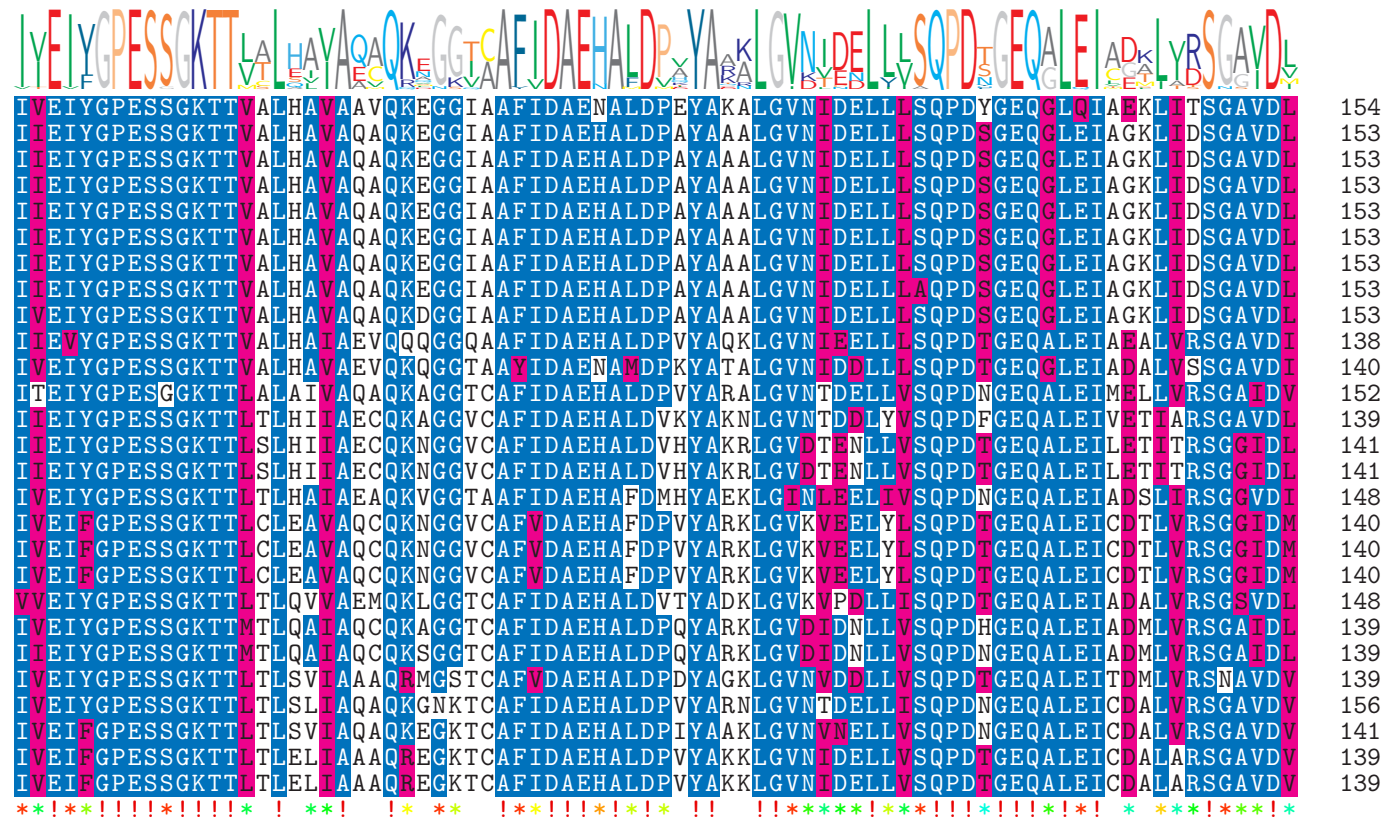


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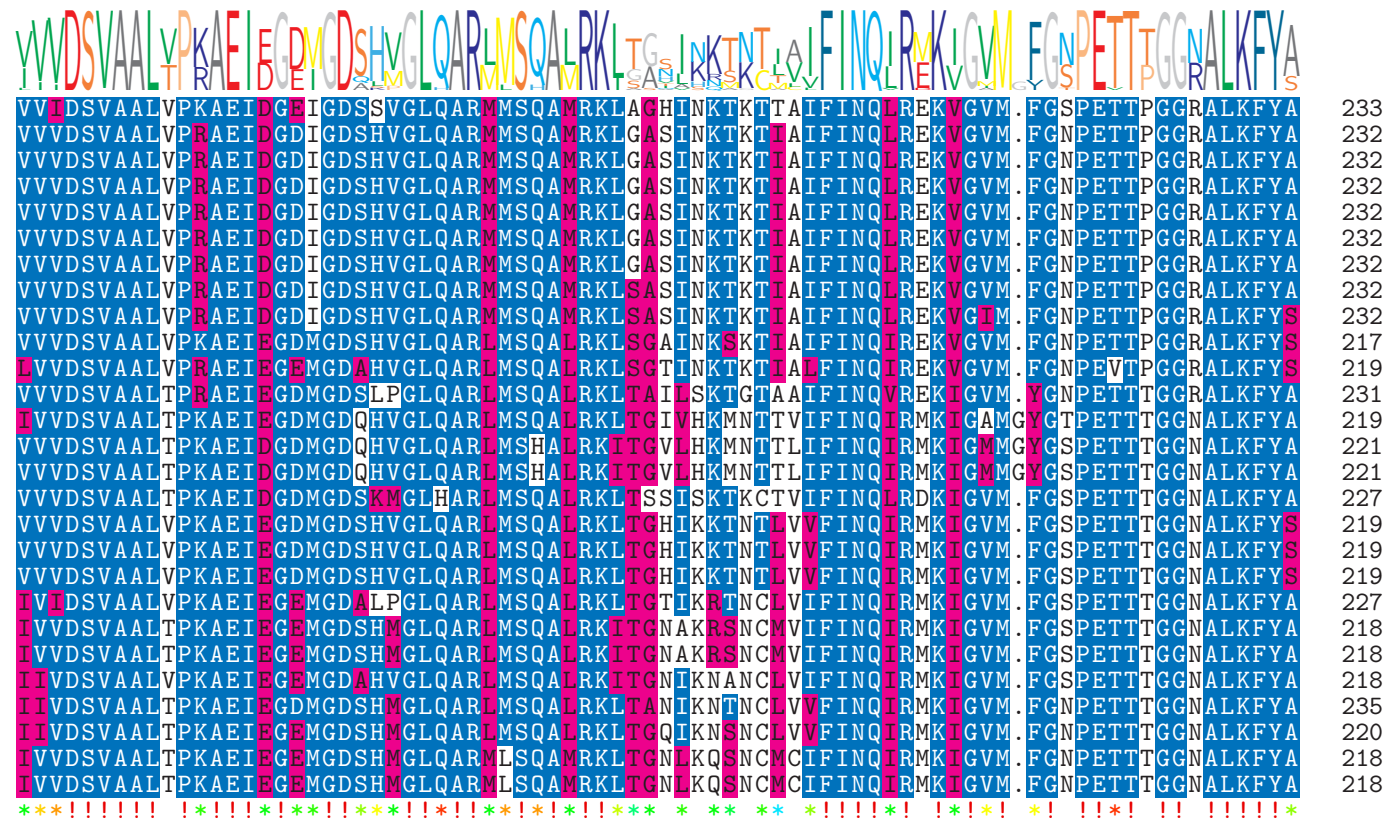
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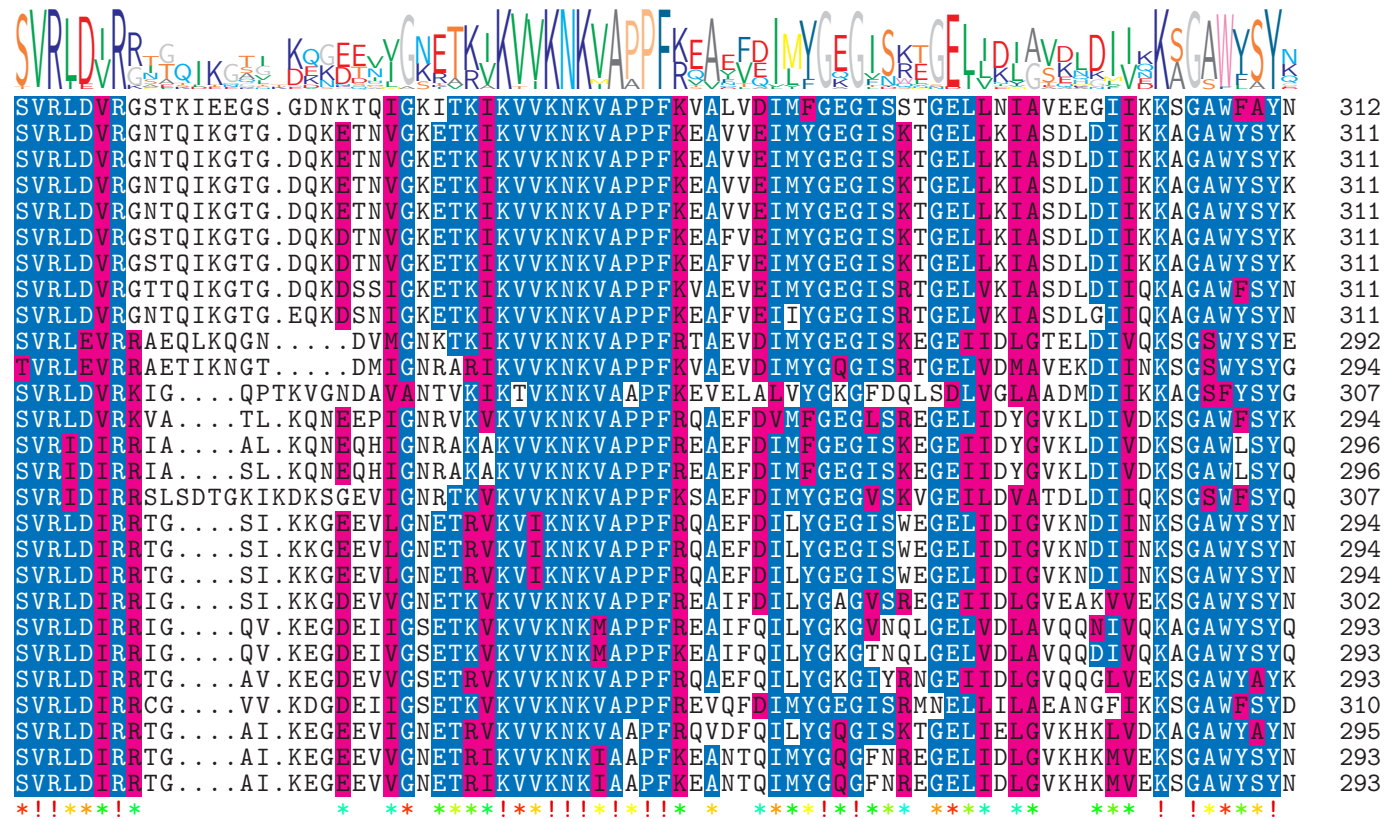
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Sequence	Score
DEKIGGGA ¹ E ² KA ³ KN ⁴ YL ⁵ KE ⁶ HQ ⁷ DV ⁸ FD ⁹ EID ¹⁰ HK ¹¹ VRAA ¹² HG ¹³ LDDSEVAETEEETTAS ¹⁴ KT ¹⁵ KAKAKKEEK ¹⁶ AVETEEIELE ¹⁷ D.....	387
DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ KQ ¹⁰ VR ¹¹ SK ¹² FG ¹³ LIDGEEVSEQD ¹⁴ TENKKDE...PKKEEA...VNEEVPLDLGDELEIE	385
DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ KQ ¹⁰ VR ¹¹ SK ¹² FG ¹³ LIDGEEVSEQD ¹⁴ TENKKDE...PKKEEA...VNEEVPLDLGDELEIE	385
DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ KQ ¹⁰ VR ¹¹ SK ¹² FG ¹³ LIDGEEVSEQD ¹⁴ TENKKDE...PKKEEA...VNEEVPLDLGDELEIE	385
DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ DN ⁵ PE ⁶ VFA ⁷ EID ⁸ HQ ⁹ VR ¹⁰ VQ ¹¹ FG ¹² LIDGEEAAE...SKKDE...VAQVDS...VNEEVTLDLGDLEIE	381
DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ DN ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ HQ ¹⁰ VR ¹¹ VQ ¹² FG ¹³ LIDGEEIAE...SKKDE...AAQVDS...VNEEVTLDLGDLEIE	381
GEKIGGGS ¹ EN ² AK ³ RYLA ⁴ DHP ⁵ QL ⁶ FD ⁷ EID ⁸ DR ⁹ KV ¹⁰ RV ¹¹ KF ¹² GLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGGGS ¹ EN ² AK ³ KFLA ⁴ DN ⁵ PE ⁶ IF ⁷ DD ⁸ ID ⁹ HK ¹⁰ VR ¹¹ VQ ¹² YGLIEEDNTEEKQSSKEK.....ETDEK...ADKNLVLELDDTIELE	382
EE ¹ RL ² GGGREN ³ AK ⁴ QFL ⁵ KN ⁶ KDIM ⁷ LI ⁸ QEQ ⁹ IREHY ¹⁰ GLD ¹¹ NNGVVQQQAET ¹² QEEL ¹³ .EFE.E.....	348
SERIGGGREN ¹ AK ² NYLA ³ DH ⁴ ED ⁵ VE ⁶ VR ⁷ LK ⁸ VRAA ⁹ YG ¹⁰ ISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGGGKE ¹ KTI ² AY ³ IA ⁴ ER ⁵ PE ⁶ ME ⁷ QEI ⁸ DR ⁹ VM ¹⁰ AA ¹¹ IRAGNAGEAPALAPAPAAPEAA.EA.....	363
DKKLGGGREN ¹ SKA ² FL ³ KN ⁴ PE ⁵ IA ⁶ DE ⁷ IT ⁸ KA ⁹ IQNSM ¹⁰ G ¹¹ IEGMI.....SGSEDDGEE.....	343
DKKLGGGREN ¹ AK ² ALL ³ KED ⁴ KALA ⁵ DE ⁶ IT ⁷ LK ⁸ IKESIG ⁹ SNEEI....MPLPDEPLEEME.....	347
DKKLGGGREN ¹ AK ² ALL ³ KED ⁴ KALA ⁵ DE ⁶ IT ⁷ LK ⁸ IKESIG ⁹ SNEEI....MPLPDEPLEEME.....	347
DTKLGGGRD ¹ TV ² KEL ³ IK ⁴ DN ⁵ PE ⁶ LAE ⁷ ELE ⁸ AK ⁹ IREKLQAK.....	343
GAKIGGGKD ¹ NVR ² VWL ³ KN ⁴ PE ⁵ IS ⁶ DE ⁷ IDA ⁸ KI ⁹ RALN ¹⁰ GVEMHI....TEGTQDET ¹¹ DGE.....RPEE.....	348
GAKIGGGKD ¹ NVR ² VWL ³ KN ⁴ PE ⁵ IS ⁶ DE ⁷ IDA ⁸ KI ⁹ RALN ¹⁰ GVEMHI....TEGTQDET ¹¹ DGE.....RPEE.....	348
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GERIGGGRD ¹ NC ² REFL ³ RENA ⁴ ELARE ⁵ IE ⁶ NKV ⁷ REHL ⁸ GVTPMG....AVTLAAEEVEED.....	352
GNKIGGGK ¹ NNV ² IRY ³ LEEN ⁴ PQ ⁵ ISTE ⁶ IE ⁷ AV ⁸ IREQLLTKASD....QTAAHDET.EEEPDL....LES.....	349
GNKIGGGK ¹ NNV ² IRY ³ FEENN ⁴ QIA ⁵ EIE ⁶ IRN ⁷ IREQLLTGTN....GAVQI.ED.EEEPDL....LLES.....	349
GSKIGGGKANA ¹ AAK ² YLEEN ³ PEIG ⁴ QIE ⁵ IEQQ ⁶ IRAKLL ⁷ VTAN....TKATPVAEDLADADI.....	347
GTKIGGGK ¹ NNAV ² KWL ³ KN ⁴ PE ⁵ IA ⁶ DK ⁷ LQE ⁸ IRNTL ⁹ ASNPNS....VLKATASDSNDES ¹⁰ SDV...EEFSEEL.....	372
GEKIGGGKAN ¹ AMK ² WLEEH ³ PEE ⁴ ALALE ⁵ TK ⁶ LRNELLAN ⁷ PEK....VLAADIAEKN.ESST...GLEADY.....	354
GDKIGGGKAN ¹ ACK ² YL ³ KN ⁴ PE ⁵ IAKT ⁶ LDK ⁷ KLREMLLN ⁸ PENM....QLIAETSSAADDVEF...GAVPEEF.....	354
GD ¹ KIGGGKAN ² ACK ³ YL ⁴ KN ⁵ PE ⁶ IAKT ⁷ LDK ⁸ KLREMLLN ⁹ PENM....QLIAETSSAADDVEF...GAVPEEF.....	354

logo

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consensus			

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