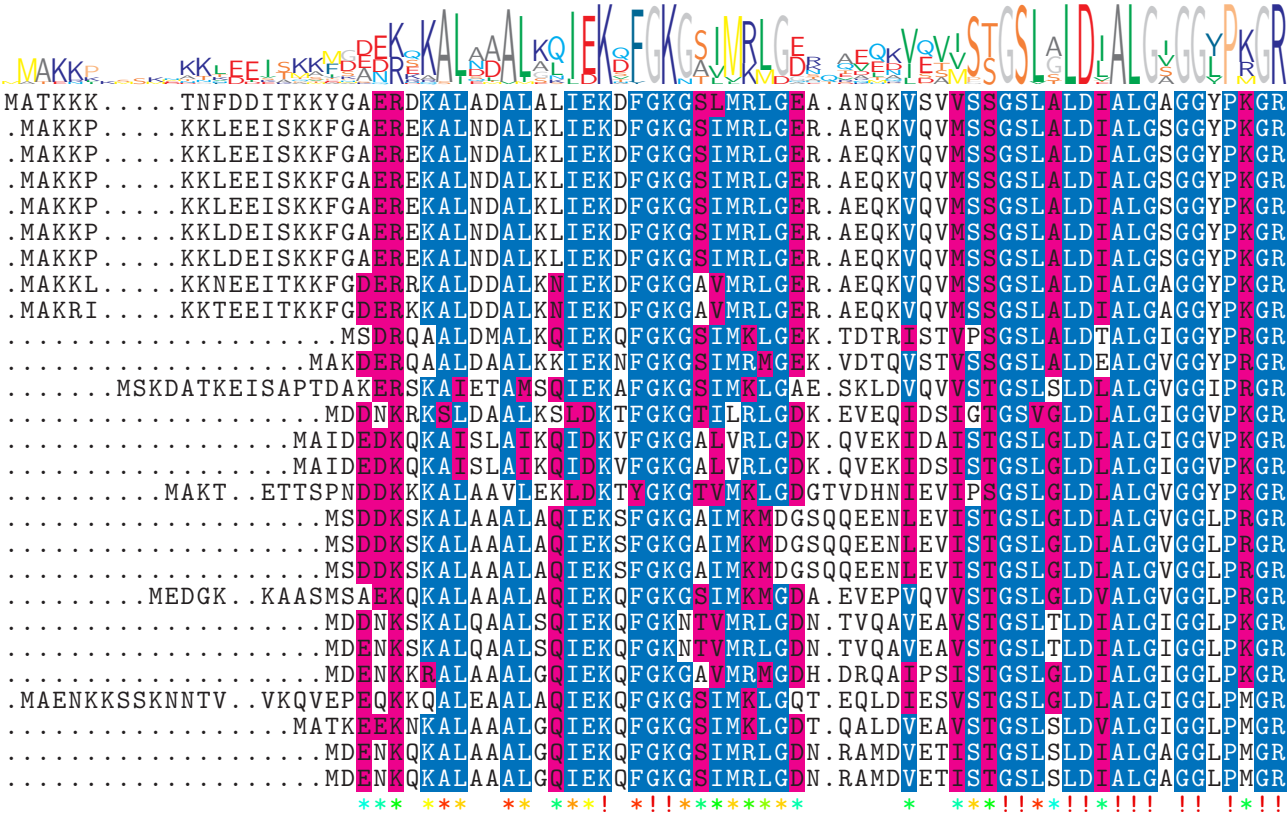


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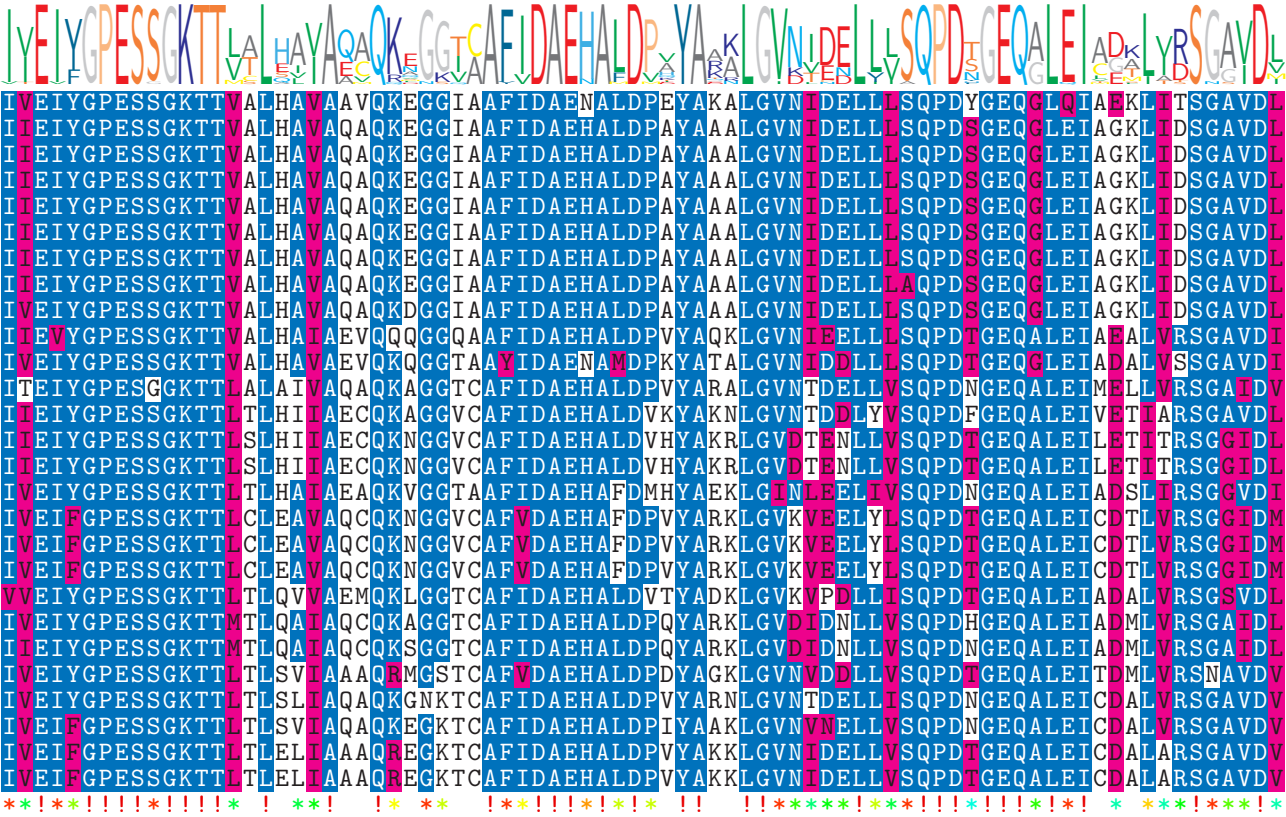
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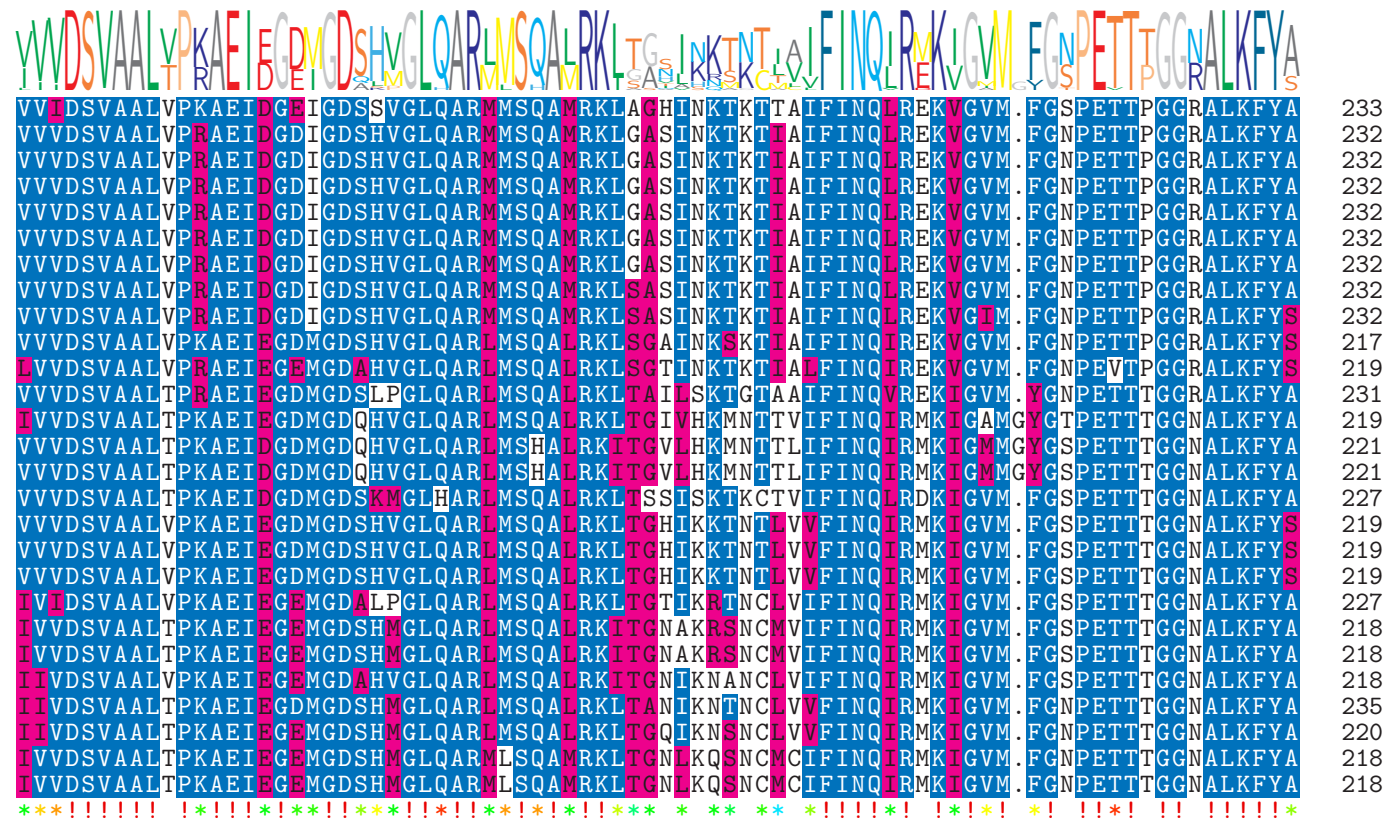


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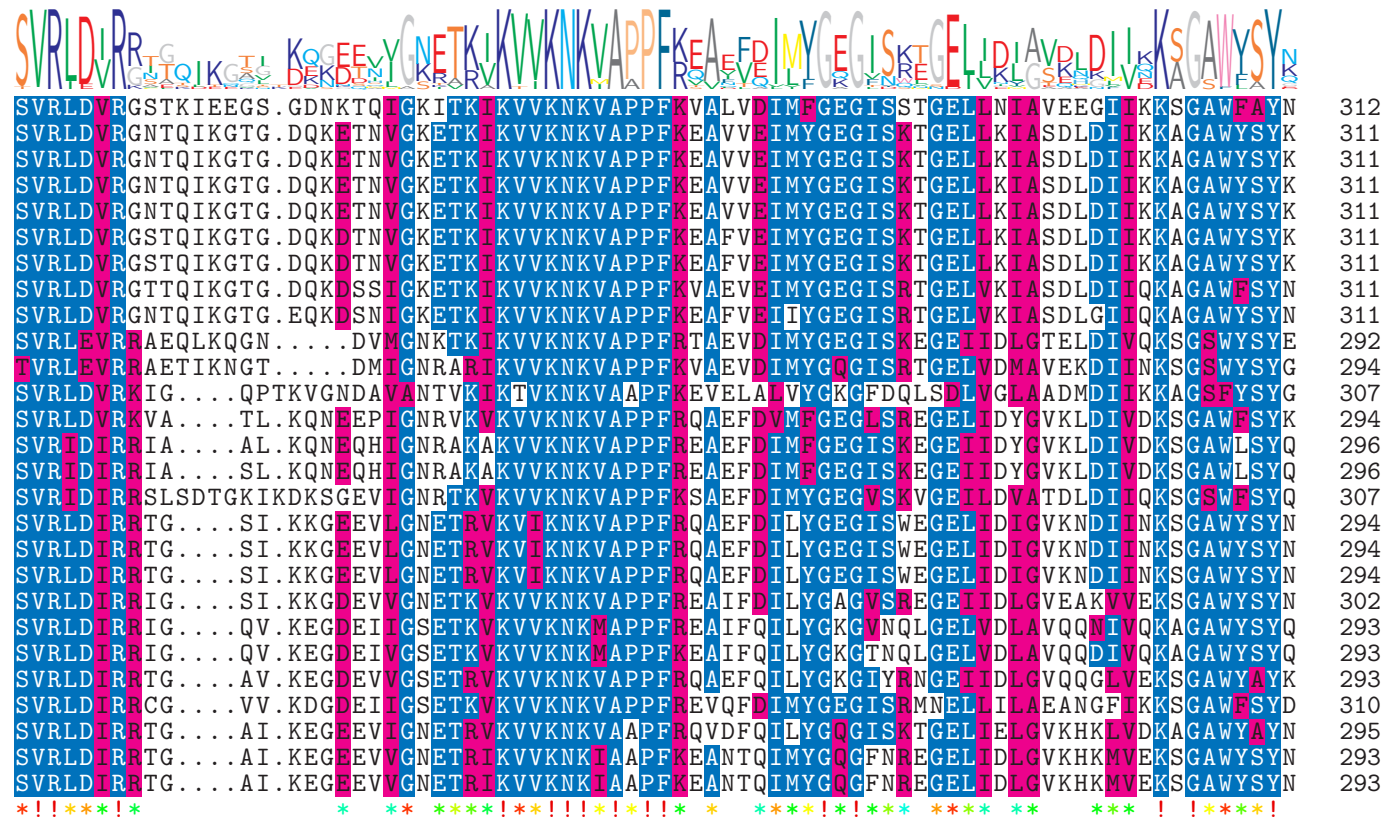
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Sequence	Score
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DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ KQ ¹⁰ VR ¹¹ SK ¹² FG ¹³ LIDGEEVSEQD ¹⁴ TENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ EH ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ KQ ¹⁰ VR ¹¹ SK ¹² FG ¹³ LIDGEEVSEQD ¹⁴ TENKKDE...PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGGGS ¹ EN ² AK ³ KYLA ⁴ DN ⁵ PE ⁶ IF ⁷ DE ⁸ ID ⁹ HK 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 RV ⁹ RV ¹⁰ KF ¹¹ GLLEESEESAMAVASE.....E.....TDDLALDLNNGIEIE	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 GLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EE ¹ RL ² GQ ³ GRE ⁴ NA ⁵ KQ ⁶ FL ⁷ KE ⁸ NK ⁹ DI ¹⁰ ML ¹¹ IQ ¹² EQ ¹³ IR ¹⁴ EH ¹⁵ YGLD ¹⁶ NNGVVQQQAET ¹⁷ QEEL ¹⁸ .EFE.E.....	348
SERIGGG ¹ RE ² NA ³ KNYLA ⁴ DH ⁵ ED ⁶ VE ⁷ DL ⁸ KV ⁹ RAA ¹⁰ YG ¹¹ ISD ¹² VPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGGG ¹ KE ² KTI ³ AY ⁴ IA ⁵ ER ⁶ PE ⁷ ME ⁸ QE ⁹ IT ¹⁰ DR ¹¹ VM ¹² AA ¹³ IR ¹⁴ AG ¹⁵ NAGEAPALAPAPAAPEAA.EA.....	363
DKKL ¹ GQ ² GRE ³ NS ⁴ KA ⁵ FL ⁶ KE ⁷ DN ⁸ PE ⁹ IA ¹⁰ DE ¹¹ IT ¹² KA ¹³ IQ ¹⁴ NS ¹⁵ MG ¹⁶ IE ¹⁷ GM ¹⁸ I.....SGSDEGEE.....	343
DKKL ¹ GQ ² GRE ³ NA ⁴ KALL ⁵ KED ⁶ KALA ⁷ DE ⁸ IT ⁹ TL ¹⁰ KI ¹¹ KE ¹² SI ¹³ GS ¹⁴ NEEI.....MPLPDEPLEEME.....	347
DKKL ¹ GQ ² GRE ³ NA ⁴ KALL ⁵ KED ⁶ KALA ⁷ DE ⁸ IT ⁹ TL ¹⁰ KI ¹¹ KE ¹² SI ¹³ GS ¹⁴ NEEI.....MPLPDEPLEEME.....	347
DTKL ¹ GQ ² GRD ³ TV ⁴ KEL ⁵ IK ⁶ DN ⁷ PE ⁸ LA ⁹ EEL ¹⁰ EAK ¹¹ IREKLQAK.....	343
GAKIGGK ¹ DN ² VR ³ VL ⁴ KN ⁵ PE ⁶ IS ⁷ DE ⁸ IDA ⁹ KI ¹⁰ RALN ¹¹ GV ¹² EM ¹³ HI.....TEGTQDET ¹⁴ DGE.....RPEE.....	348
GAKIGGK ¹ DN ² VR ³ VL ⁴ KN ⁵ PE ⁶ IS ⁷ DE ⁸ IDA ⁹ KI ¹⁰ RALN ¹¹ GV ¹² EM ¹³ HI.....TEGTQDET ¹⁴ DGE.....RPEE.....	348
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GE ¹ RIG ² GRD ³ NC ⁴ REF ⁵ LRE ⁶ NA ⁷ E ⁸ LA ⁹ RE ¹⁰ IE ¹¹ NK ¹² VRE ¹³ HL ¹⁴ GV ¹⁵ TP ¹⁶ MG.....AVTLAAEEVEED.....	352
GNKIGGG ¹ KNN ² VIR ³ YLE ⁴ EN ⁵ PE ⁶ QI ⁷ ST ⁸ EIE ⁹ AV ¹⁰ IREQLLTKASD.....QTAAHDET.EEEPDL...LES.....	349
GNKIGGG ¹ KNN ² VIR ³ YFE ⁴ ENN ⁵ QI ⁶ AE ⁷ IE ⁸ RN ⁹ IREQLLT ¹⁰ TGTN.....GAVQI.ED.EEEPDL...LLES.....	349
GSKIGGK ¹ ANA ² AK ³ YLE ⁴ EN ⁵ PE ⁶ IG ⁷ EQ ⁸ EIE ⁹ QQ ¹⁰ IR ¹¹ AK ¹² LL ¹³ TSAN.....TKATPVAEDLADADI.....	347
GTKIGGG ¹ KNN ² AVK ³ WL ⁴ KN ⁵ PE ⁶ IA ⁷ DK ⁸ LE ⁹ QE ¹⁰ IR ¹¹ N ¹² TLAS ¹³ NPNS.....VLKATASDSNDES ¹⁴ SV..EEFSEEEEL.....	372
GEKIGGK ¹ KANAMK ² WLE ³ EH ⁴ PE ⁵ EALA ⁶ LET ⁷ KL ⁸ RNELLAN ⁹ PEK.....VLAADIAEKN.ESST..GLEADY.....	354
GDKIGGG ¹ KANACK ² YL ³ KN ⁴ PE ⁵ IA ⁶ KT ⁷ LD ⁸ KK ⁹ REMLLN ¹⁰ PENM.....QLIAETSSAADDVEF..GAVPEEF.....	354
GDKIGGG ¹ KANACK ² YL ³ KN ⁴ PE ⁵ IA ⁶ KT ⁷ LD ⁸ KK ⁹ REMLLN ¹⁰ PENM.....QLIAETSSAADDVEF..GAVPEEF.....	354

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consensus			

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