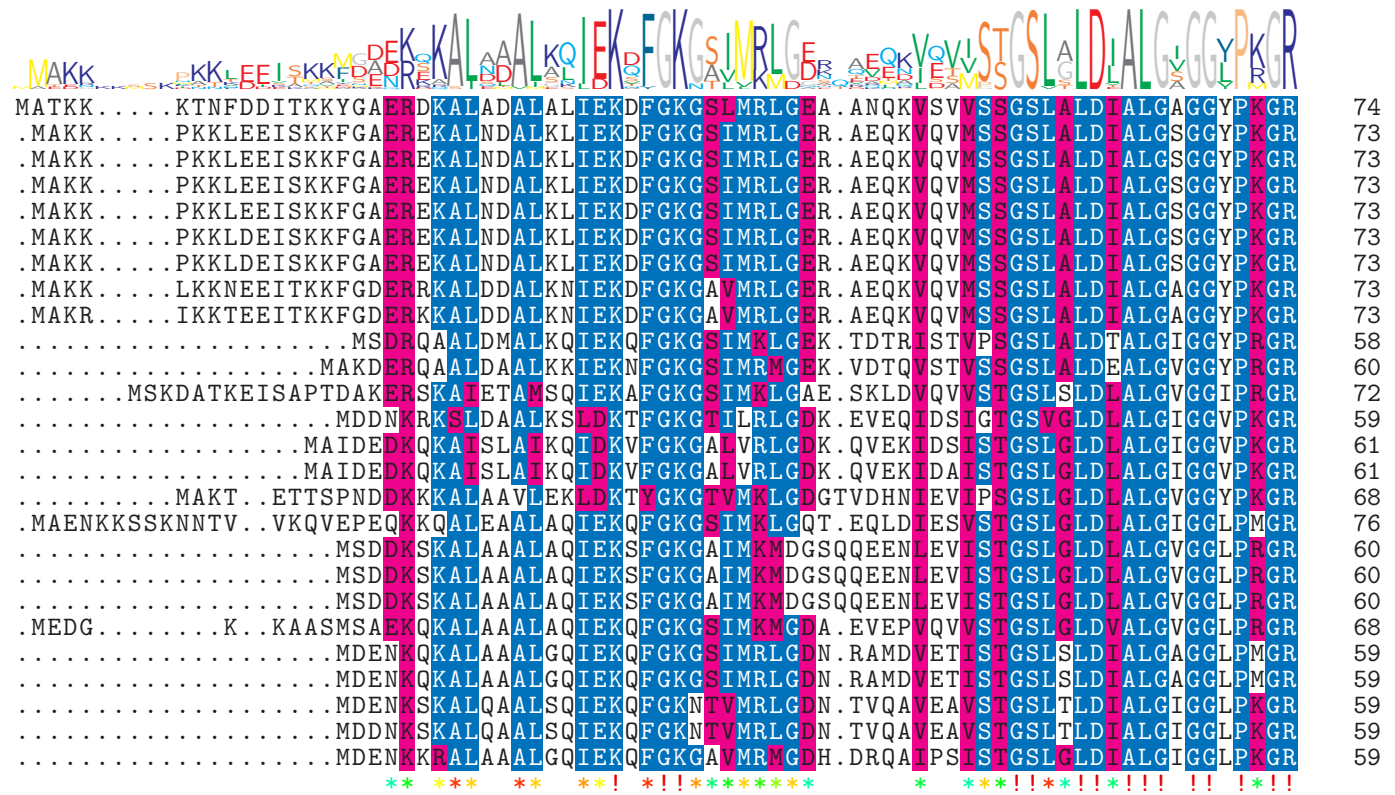


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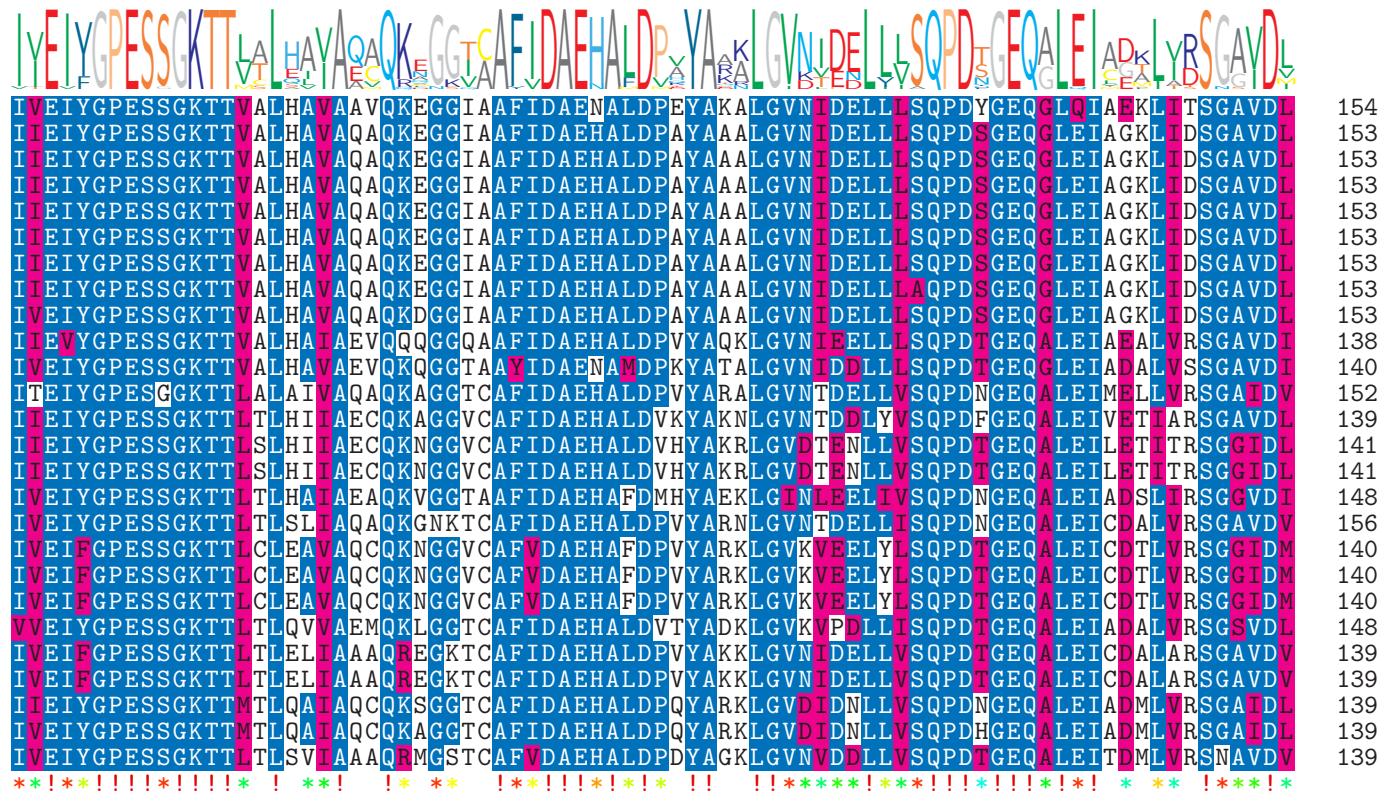
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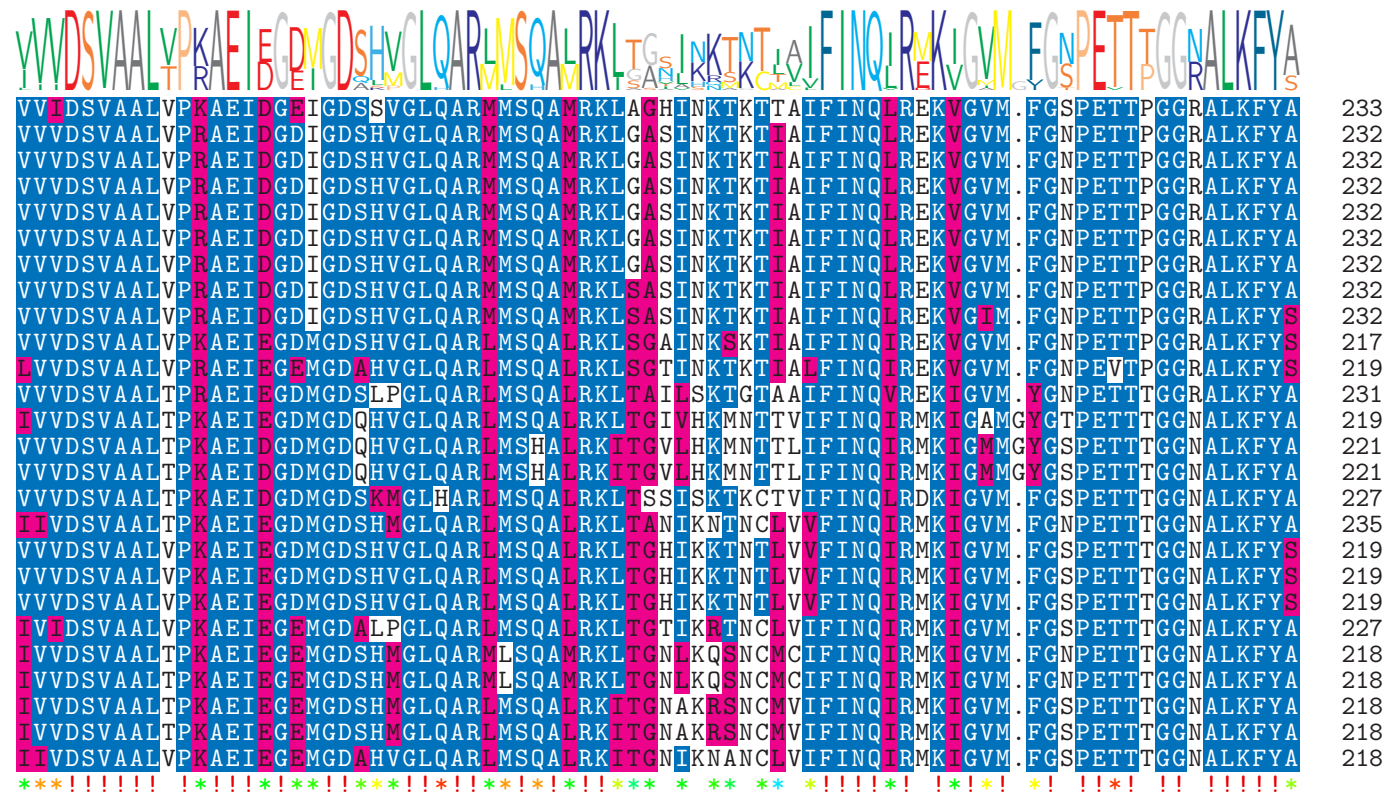
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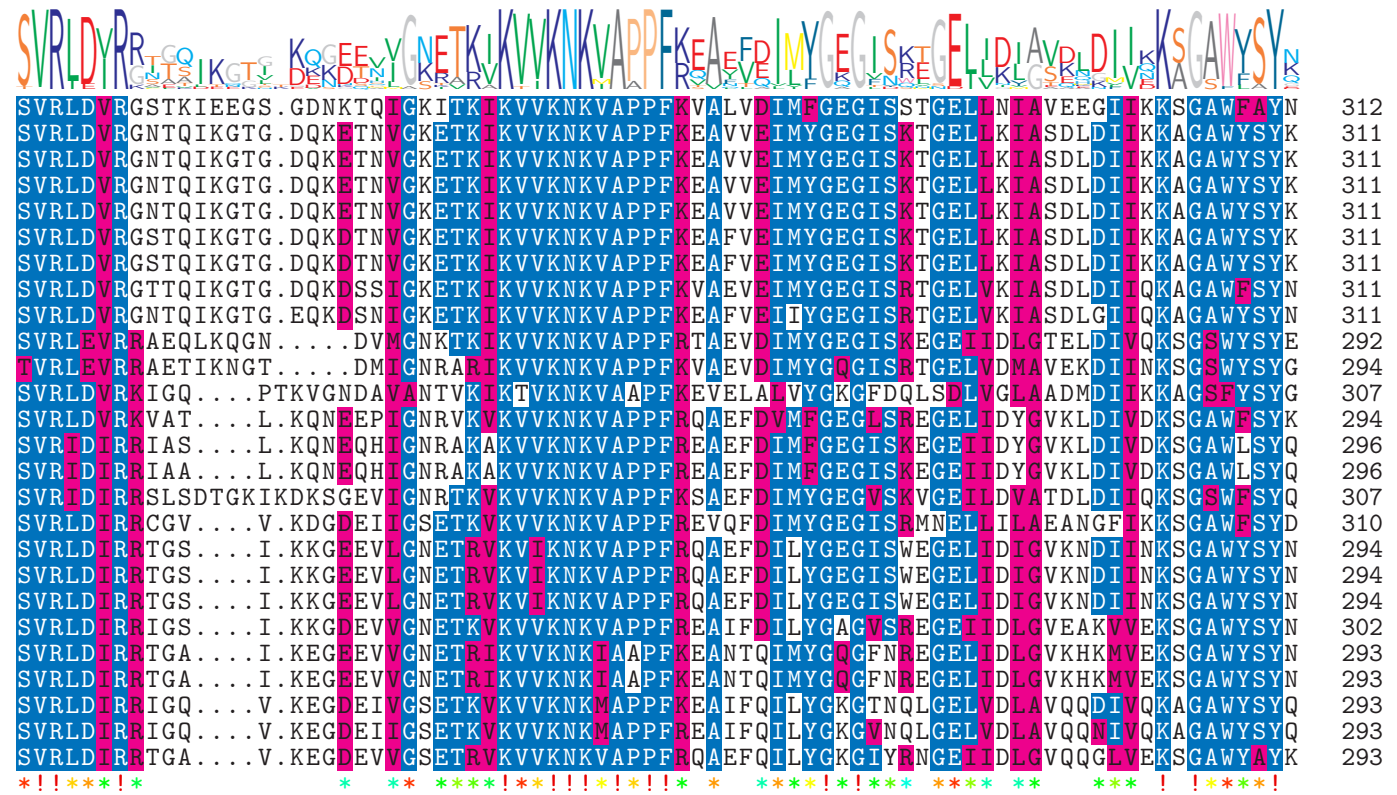
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Sequence	Score
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DEKIGQGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EI ⁺ FD ⁺ EI ⁺ DK ⁺ QV ⁺ RS ⁺ KF ⁺ GL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EI ⁺ FD ⁺ EI ⁺ DK ⁺ QV ⁺ RS ⁺ KF ⁺ GL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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GEKIGQGS ⁺ EN ⁺ AK ⁺ KFL ⁺ AD ⁺ NP ⁺ EI ⁺ FD ⁺ DI ⁺ DH ⁺ KV ⁺ RV ⁺ QY ⁺ GL ⁺ IEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EER ⁺ L ⁺ GGG ⁺ REN ⁺ AK ⁺ Q ⁺ FL ⁺ KEN ⁺ KD ⁺ I ⁺ ML ⁺ MI ⁺ QE ⁺ QI ⁺ RE ⁺ HY ⁺ GL ⁺ DNNGVVQQQAETQEEL.EFE.E.....	348
SERIGQGS ⁺ REN ⁺ AK ⁺ NYL ⁺ AD ⁺ HED ⁺ VE ⁺ DEV ⁺ LR ⁺ VRA ⁺ AY ⁺ GI ⁺ SDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGQGE ⁺ KTI ⁺ AY ⁺ IA ⁺ ER ⁺ PE ⁺ ME ⁺ QEI ⁺ IR ⁺ DR ⁺ VMA ⁺ AI ⁺ RAG ⁺ NAGEAP ⁺ AI ⁺ PAP ⁺ APAA ⁺ EAAA.....	363
DKKLGGG ⁺ RE ⁺ NS ⁺ KA ⁺ FL ⁺ KEN ⁺ PE ⁺ IA ⁺ DE ⁺ IT ⁺ KAI ⁺ QNS ⁺ MG ⁺ IEGMI....SGSEDDE..GEE.....	343
DKKLGGG ⁺ RE ⁺ NA ⁺ KALL ⁺ KE ⁺ DK ⁺ AL ⁺ ADE ⁺ IT ⁺ LKI ⁺ KES ⁺ IG ⁺ SNEEI....MPLPDEP..LEEME.....	347
DKKLGGG ⁺ RE ⁺ NA ⁺ KALL ⁺ KE ⁺ DK ⁺ AL ⁺ ADE ⁺ IT ⁺ LKI ⁺ KES ⁺ IG ⁺ SNEEI....MPLPDEP..LEEME.....	347
DTKLGGGRD ⁺ TV ⁺ KEL ⁺ IKD ⁺ NP ⁺ EL ⁺ AEE ⁺ LE ⁺ AK ⁺ IRE ⁺ KLQAK.....	343
GTKIGGG ⁺ KNN ⁺ AV ⁺ KWL ⁺ KEN ⁺ PE ⁺ IA ⁺ DK ⁺ LE ⁺ QEI ⁺ IR ⁺ NT ⁺ LA ⁺ SN ⁺ PNS.....VLKATASDSNDESD..VEEFSEEEEL.....	372
GAKIGGG ⁺ KDN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PE ⁺ IS ⁺ DE ⁺ IDA ⁺ KI ⁺ RAL ⁺ NC ⁺ VE ⁺ MHI....TEGTQDETD.....GERPEE.....	348
GAKIGGG ⁺ KDN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PE ⁺ IS ⁺ DE ⁺ IDA ⁺ KI ⁺ RAL ⁺ NC ⁺ VE ⁺ MHI....TEGTQDETD.....GERPEE.....	348
GAKIGGG ⁺ KDN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PE ⁺ IS ⁺ DE ⁺ IDA ⁺ KI ⁺ RAL ⁺ NC ⁺ VE ⁺ MHI....TEGTQDETD.....GERPEE.....	348
GERIGGG ⁺ RD ⁺ NC ⁺ REF ⁺ LRE ⁺ NA ⁺ EL ⁺ ARE ⁺ IE ⁺ NK ⁺ VR ⁺ RE ⁺ HL ⁺ CV ⁺ TPMG....AV.TLAEVEED.....	352
GDKIGGG ⁺ KAN ⁺ ACK ⁺ YL ⁺ KEN ⁺ PE ⁺ IA ⁺ KT ⁺ LD ⁺ KKL ⁺ RE ⁺ MLL ⁺ NP ⁺ EN ⁺ M....QLIAETSSAADDVE..FGAVPEEF.....	354
GDKIGGG ⁺ KAN ⁺ ACK ⁺ YL ⁺ KEN ⁺ PE ⁺ IA ⁺ KT ⁺ LD ⁺ KKL ⁺ RE ⁺ MLL ⁺ NP ⁺ EN ⁺ M....QLIAETSSAADDVE..FGAVPEEF.....	354
GNKIGGG ⁺ KNN ⁺ VIR ⁺ YFE ⁺ ENN ⁺ QIA ⁺ EI ⁺ ERN ⁺ IRE ⁺ QLLT ⁺ TGTN....GAVQ.I.EDEEEPD....LLES.....	349
GNKIGGG ⁺ KNN ⁺ VIR ⁺ YLE ⁺ EN ⁺ PQI ⁺ STE ⁺ IE ⁺ AV ⁺ IRE ⁺ QLLT ⁺ TKASD....QTAA.HDETEEEDPD....LLES.....	349
GSKIGGG ⁺ KANA ⁺ AK ⁺ YLE ⁺ EN ⁺ PI ⁺ GQEI ⁺ EQ ⁺ IR ⁺ AK ⁺ LL ⁺ VSAN....TKATPVAEDLADAD....I.....	347

logo

NTDB id 600 KW2 RS01765 WP 011675413.1	...	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	IEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	IEE	388
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consensus

 non conserved
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 ≥ 50% conserved