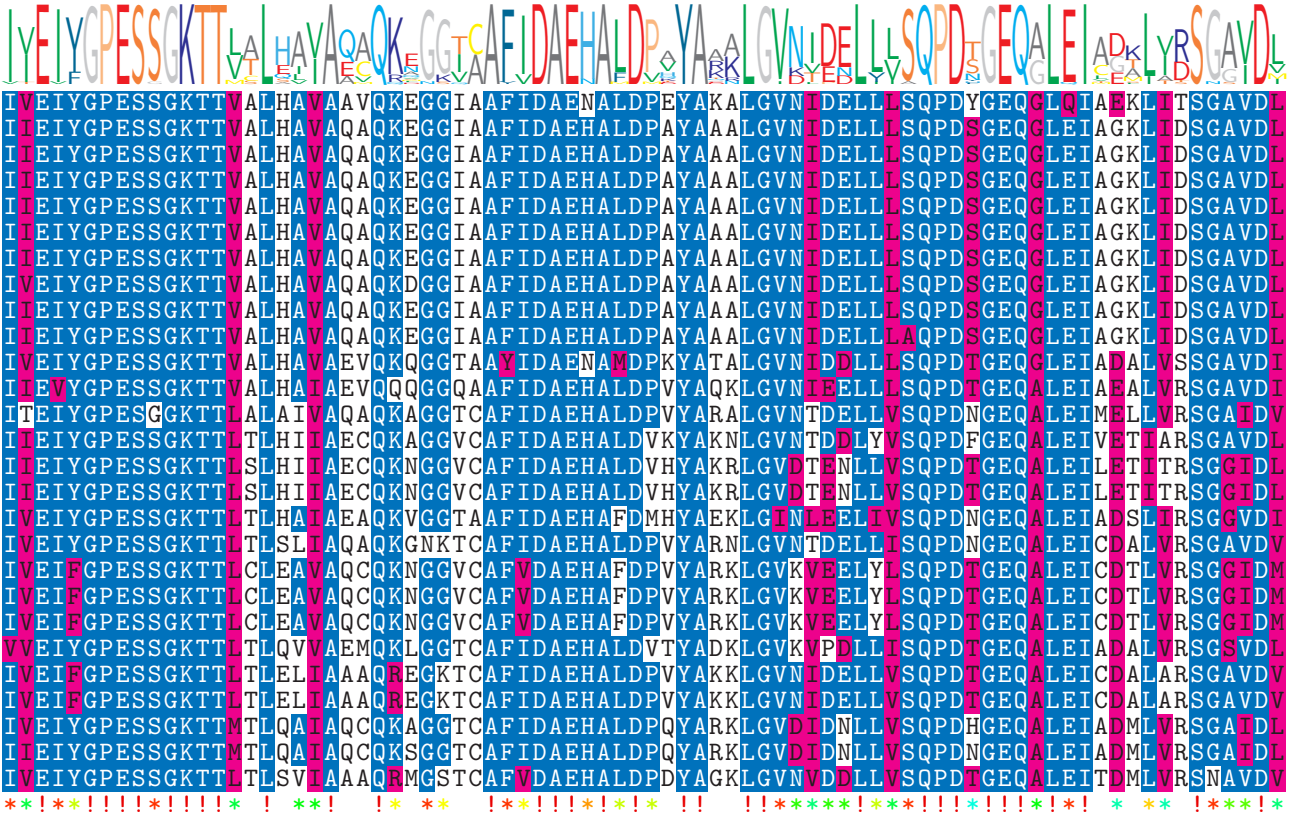


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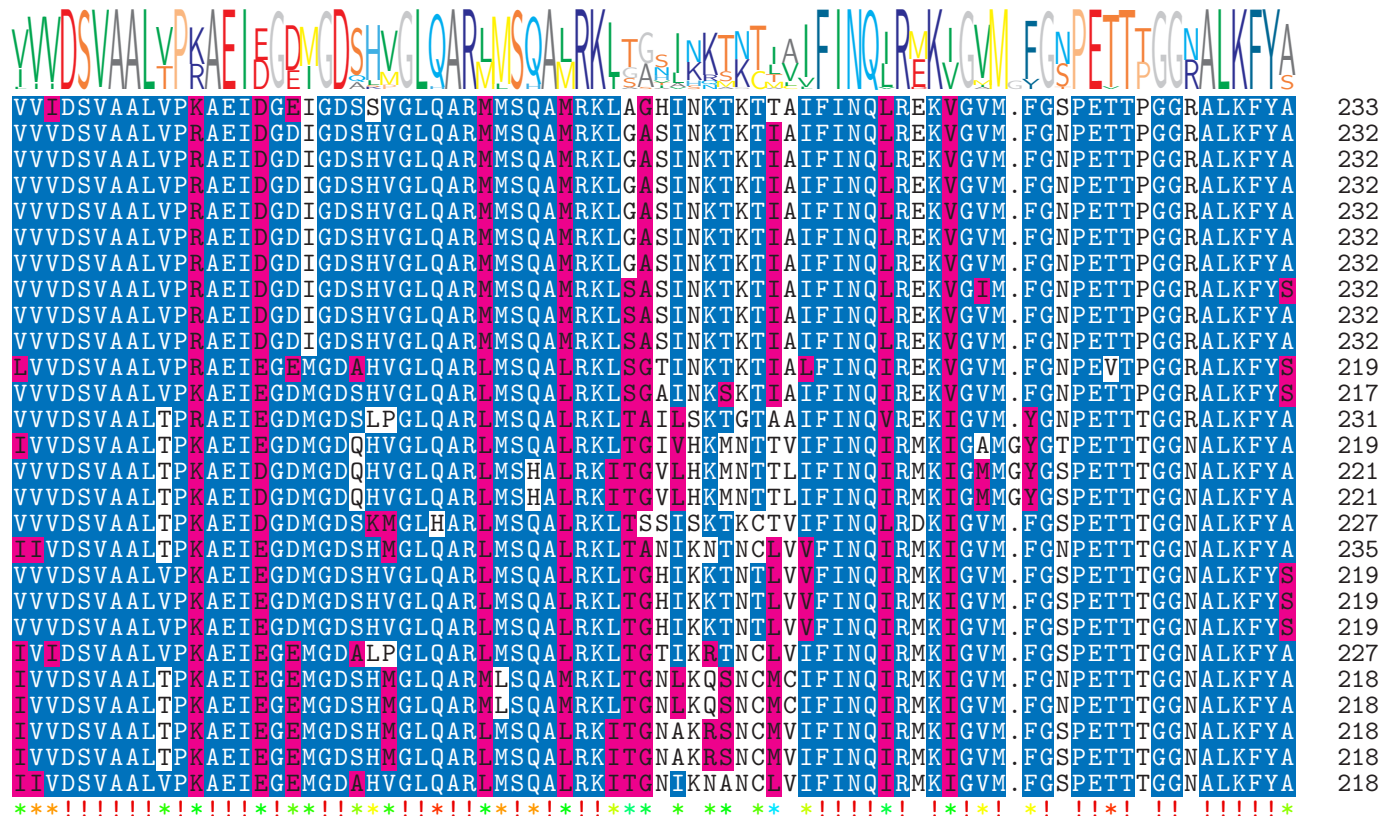


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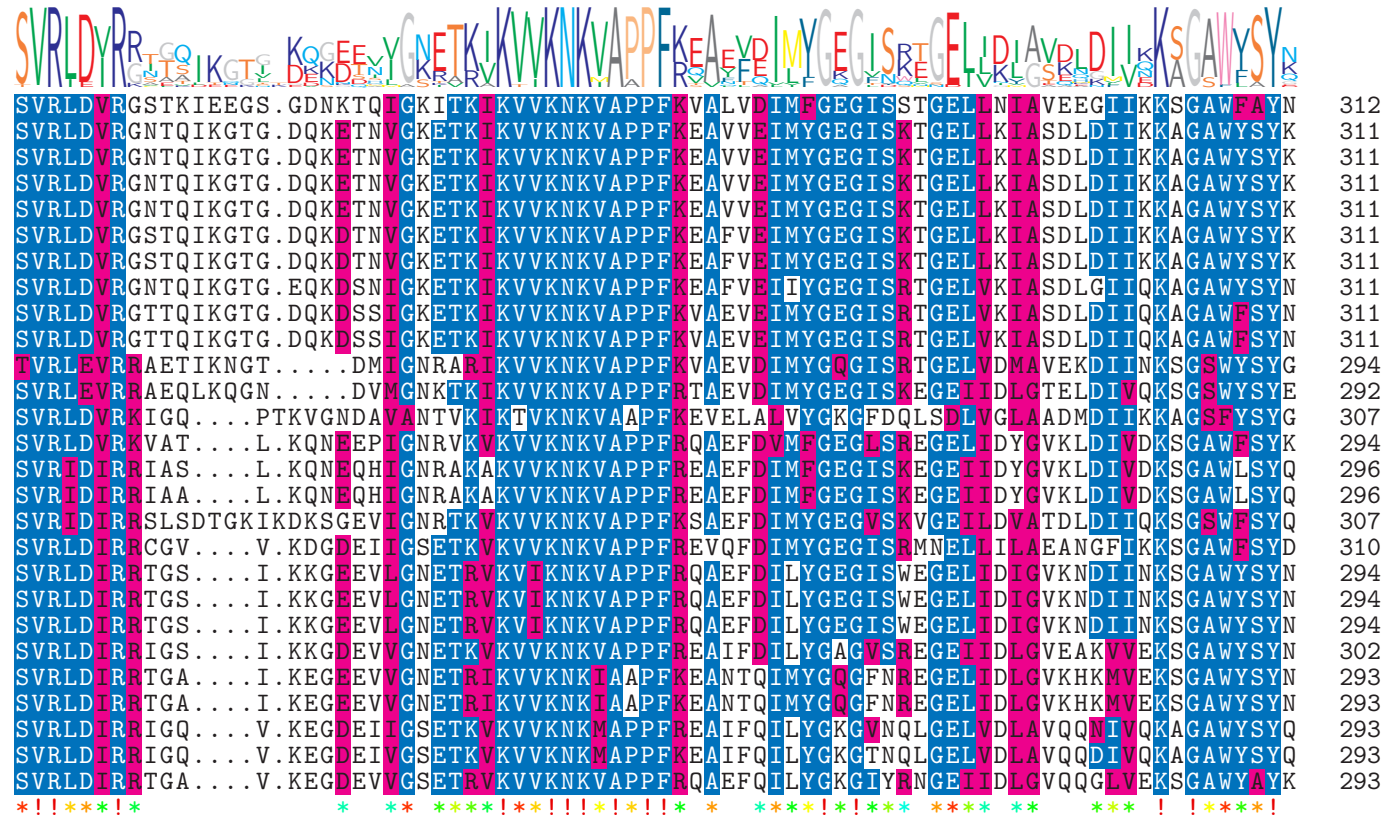
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consensus

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SERIGQG R ENAKNYL A D H DE V ED V RLKVR A AY G ISDVPEEDLPTTEDE.QIN.....ILPDDSTEE.	355
EE R L G Q G RENA K FL N ED I ML I TEQ I REHY G L D NNGVVQQQA E ET Q EELE.....FE.E.....	348
DE R IG Q Q E K T I A Y I A E R P E M E Q E I R D R V MA A I R A G E A P A L A P A A E E AEA.....	363
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DK K L G Q G RENA K ALL K ED K AL A D E IT L K I KE S IG S NE E I.....MPLPDE..PL.....EEME	347
DK K L G Q G RENA K ALL K ED K AL A D E IT L K I KE S IG S NE E I.....MPLPDE..PL.....EEME	347
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consensus			

- non conserved
- similar
- ≥ 50% conserved