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consensus

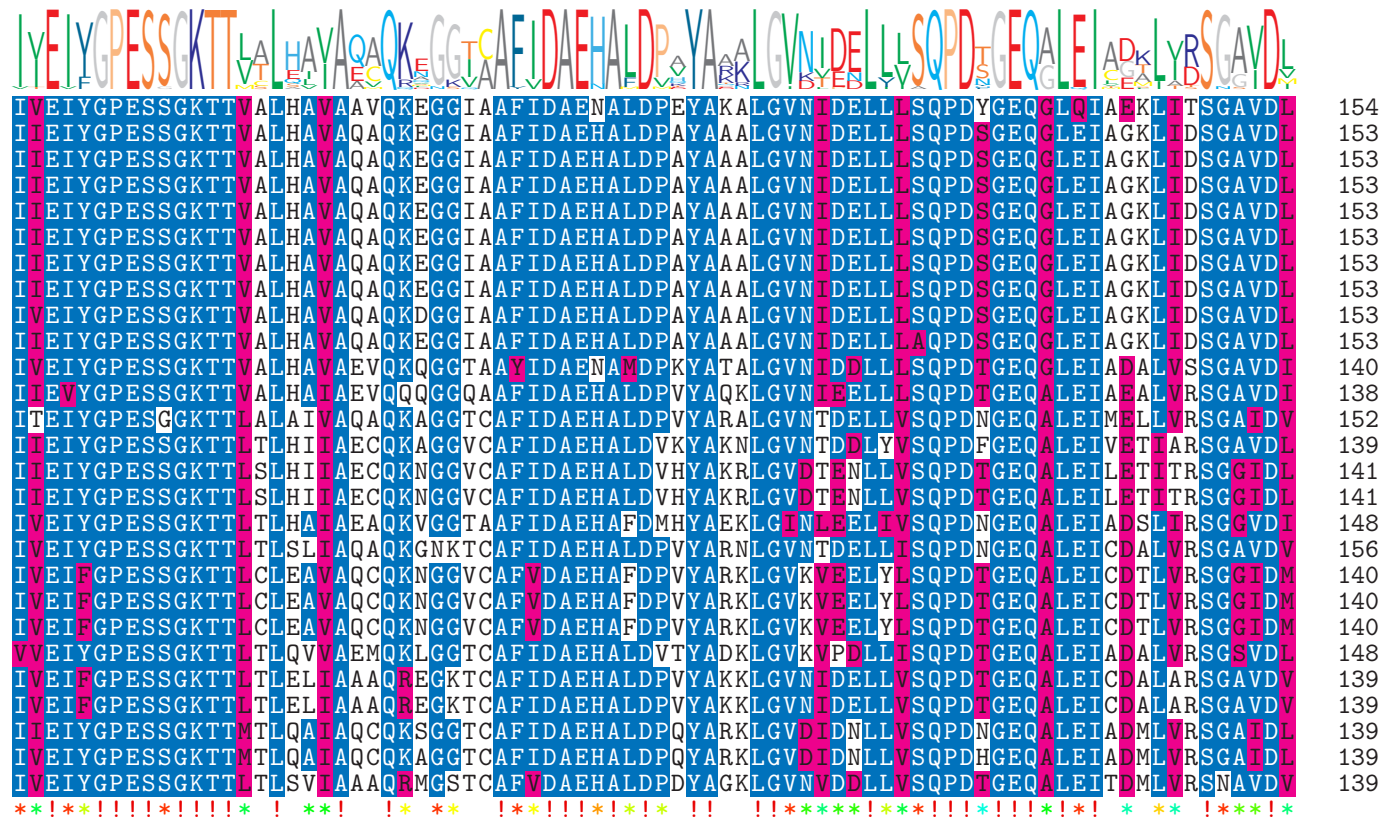
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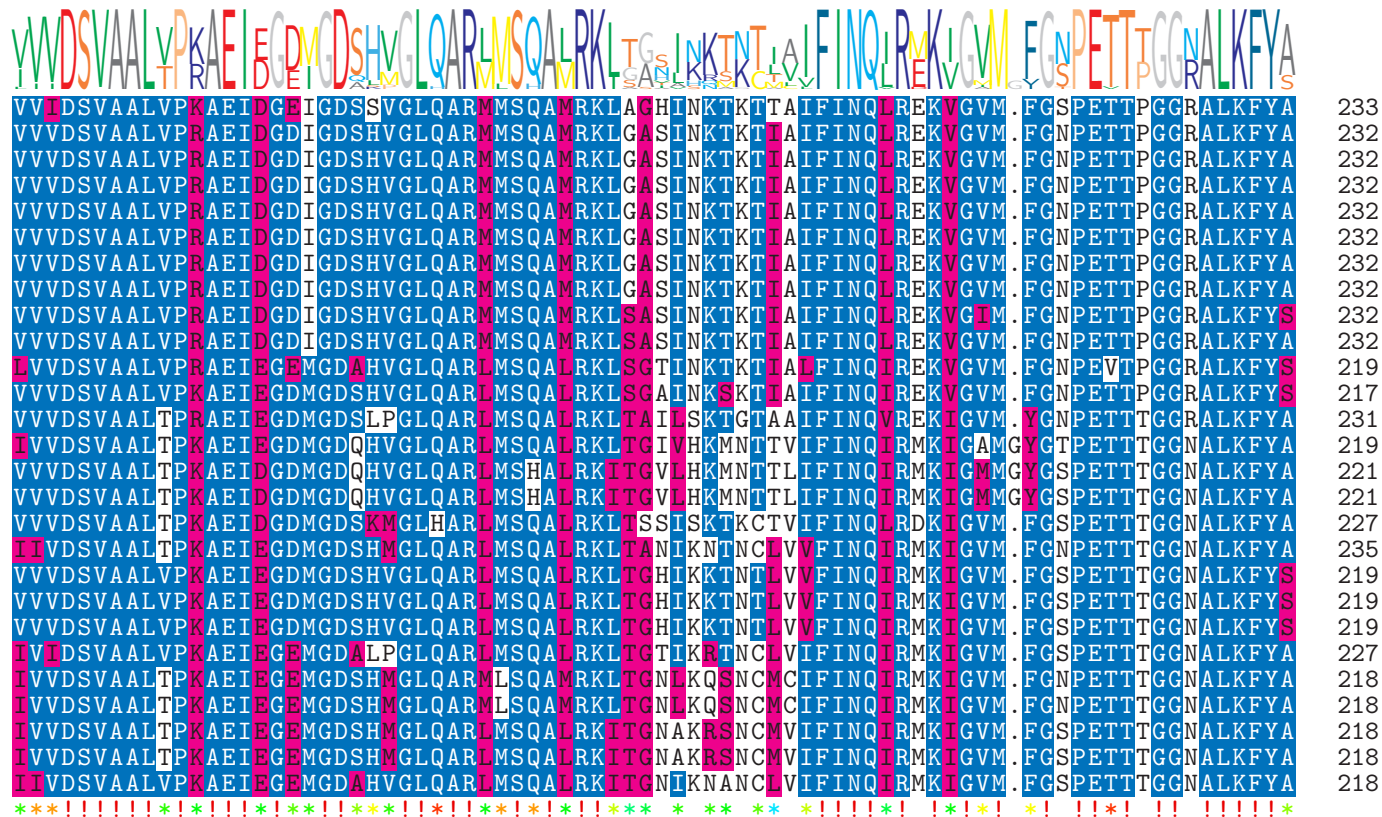
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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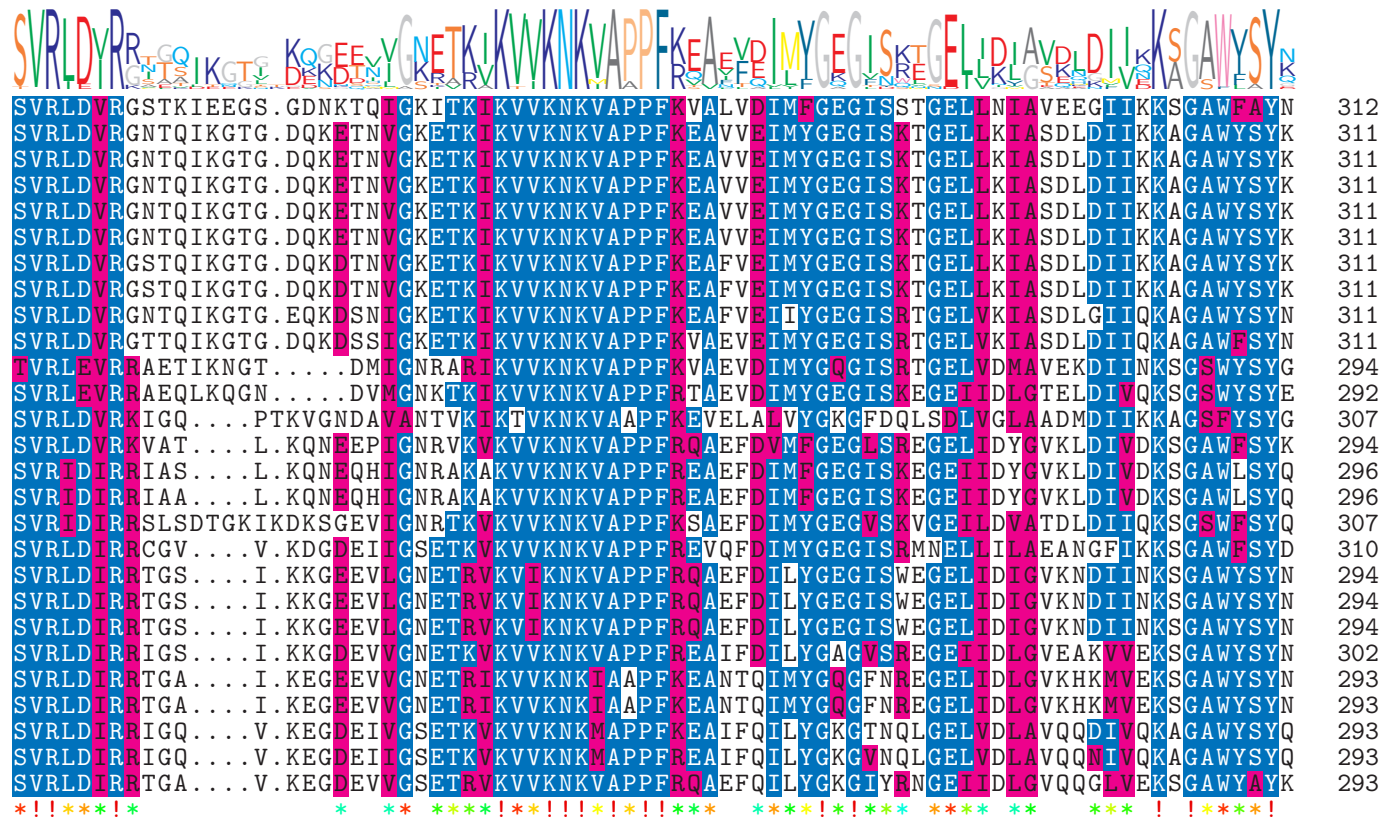
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consensus

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Sequence	Score
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DEKIGGGS E NA K KYLA E HP E IF D EIDDKQVRSKF G LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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DEKIGGGS E NA K KYLA E HP E IF D EIDDKQVRSKF G LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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GEKIGGGS E NA K RYL D HP Q L F DEIDRKVVRV K FLLEESEEEESAMAVASE.....E.....TDDLALDLNNGIEIE	377
SERIGGG R EN A KNYL D H E D V E V RLKVR A AG I SDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
EER L GGC R EN A K Q FL K EN K D I ML I Q E IR E HY L DNNGVVQQAEETQEEL.EFE.E.....	348
DERIGGG E K T I A Y I A R P E M E Q E IR D RVMA A IR A G N AGEAPALAPAPA A E A E A	363
DK K LGG R EN S K A FL K EN P E I A D E I TK A I Q NS M G I EGMI....SGSEDDE..GEE.....	343
DK K LGG R EN A K A LL K ED K AL A D E IT L K I ES I GS N EEI....MPLPDEP..LEEME.....	347
DK K LGG R EN A K A LL K ED K AL A D E IT L K I ES I GS N EEI....MPLPDEP..LEEME.....	347
DT K LGG R DT V K E L I DN P E L A E E L E A K I RE K L Q A.....	343
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GN K IGGG K NN V IR Y FE E NN Q IA E E I ERN I RE Q LL T T G TN....GAVQ.I.EDEE E PD....L L ES.....	349
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NTDB id 1205 PSJM300 05545 AFN77183.1	...	347	
consensus			

- non conserved
- similar
- ≥ 50% conserved