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consensus

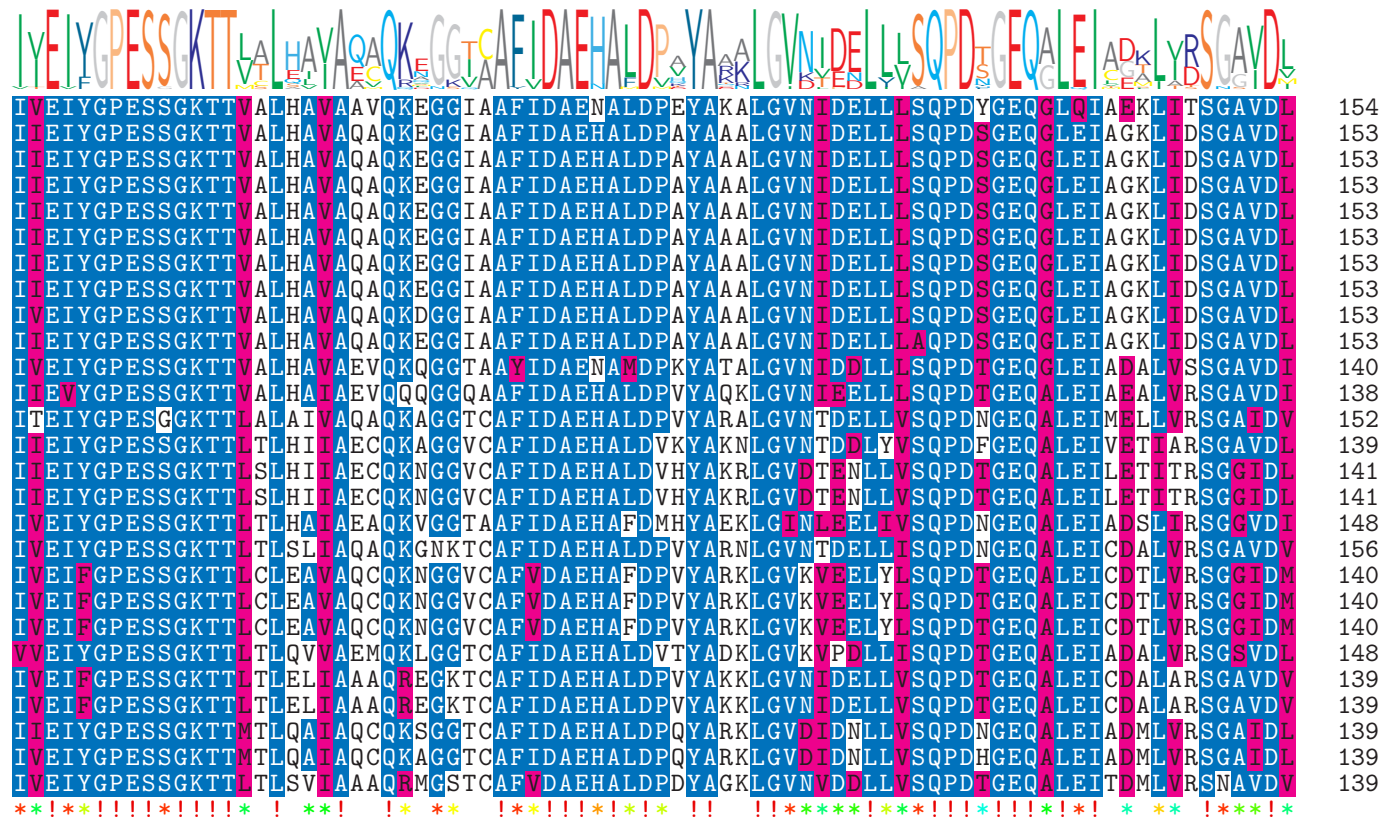
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consensus

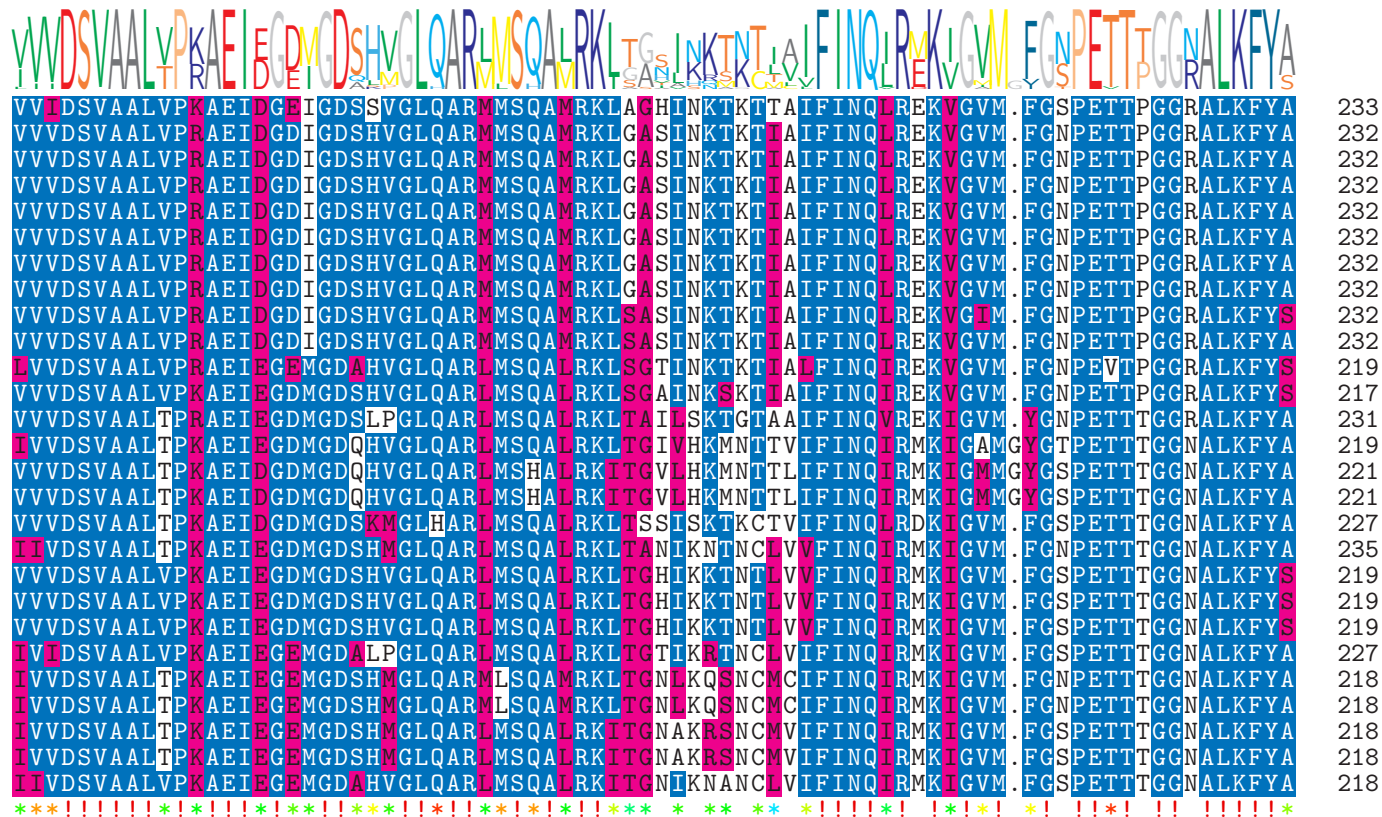
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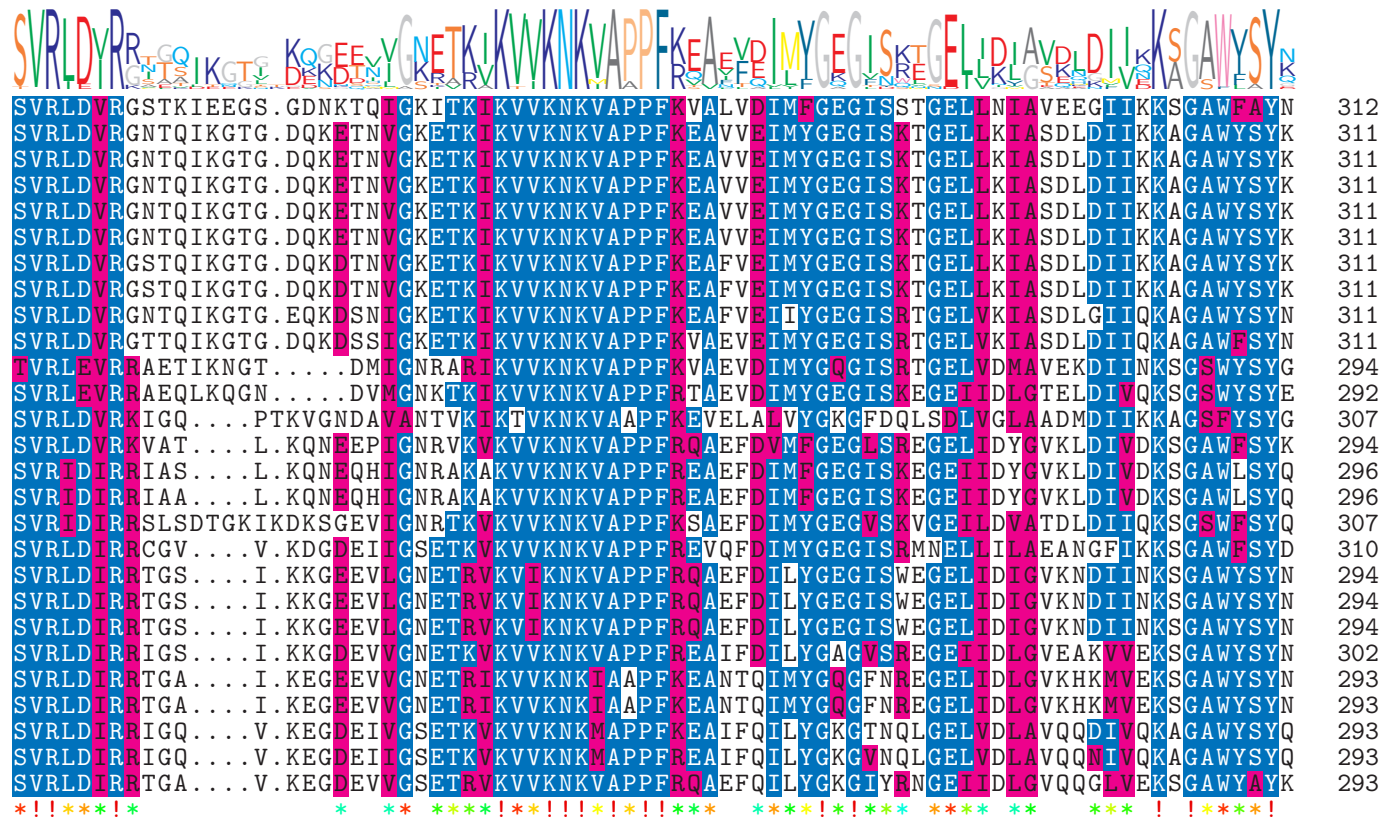
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consensus

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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
DEKIGGGS E NA K KYLA E HP E IF D EIDDKQVRSKF G LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGGS E NA K KYLA E HP E IF D EIDDKQVRSKF G LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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SERIGGG R EN A KNYL D H E D V E V RLKVR A E G ISDVPPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
EERLGGC R EN A K Q FL K EN K D I ML I IQ E IR E HY L DNNGVVQQAEETQEEL.EFE.E.....	348
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DK K LGGG R EN A K A L L K E D K A L A D E I TL K I K ES I GS N EEI....MPLPDEP..LEEME.....	347
DK K LGGG R EN A K A L L K E D K A L A D E I TL K I K ES I GS N EEI....MPLPDEP..LEEME.....	347
DT K LGGG R D T V K EL I K D N P E L A E E L E A K I RE K L Q A.....	343
GT K IGGG K NAV K WL K EN P E I A D K L E I Q I RT L AS N PN SVLKATASDSNDESD..VEEFSEEL.....	372
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logo

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consensus			

- non conserved
- similar
- ≥ 50% conserved