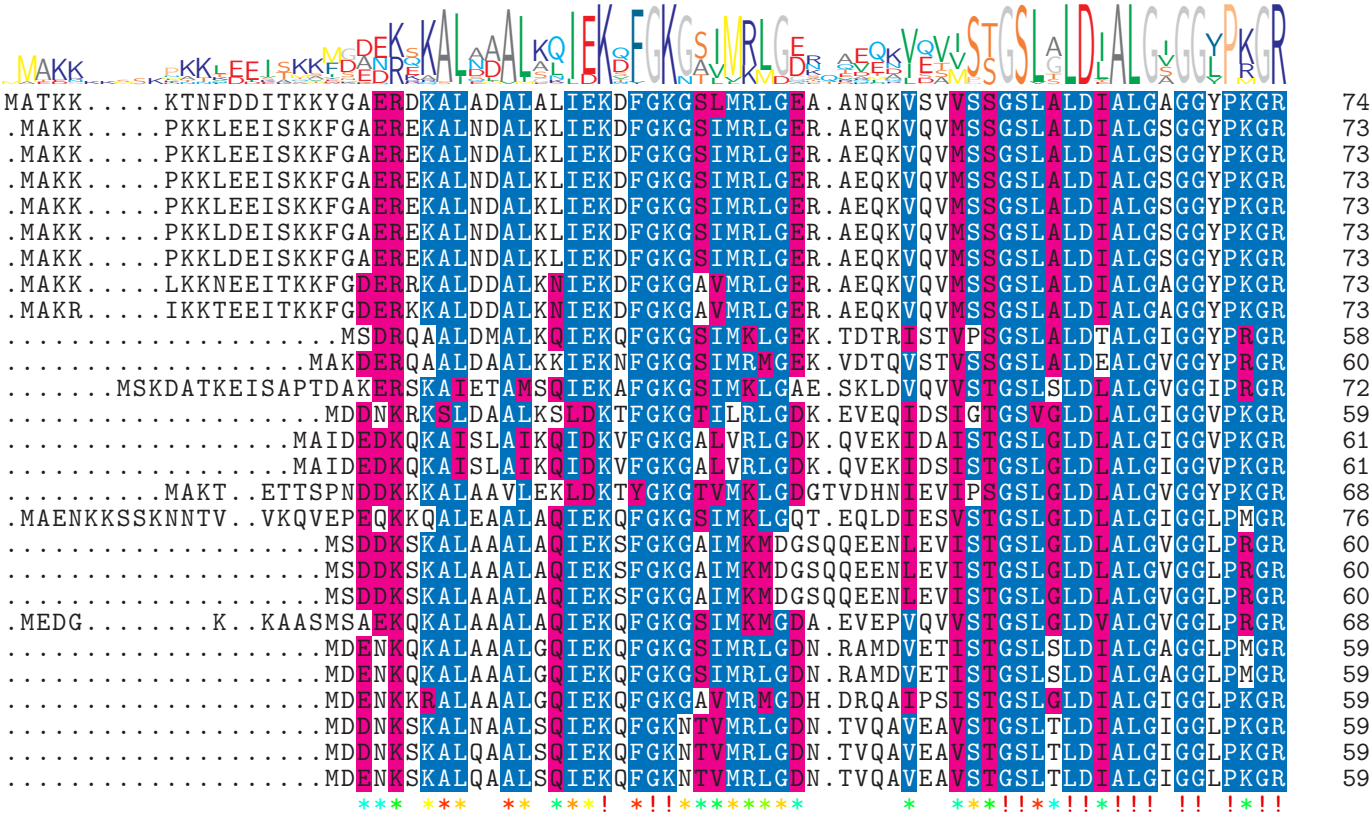


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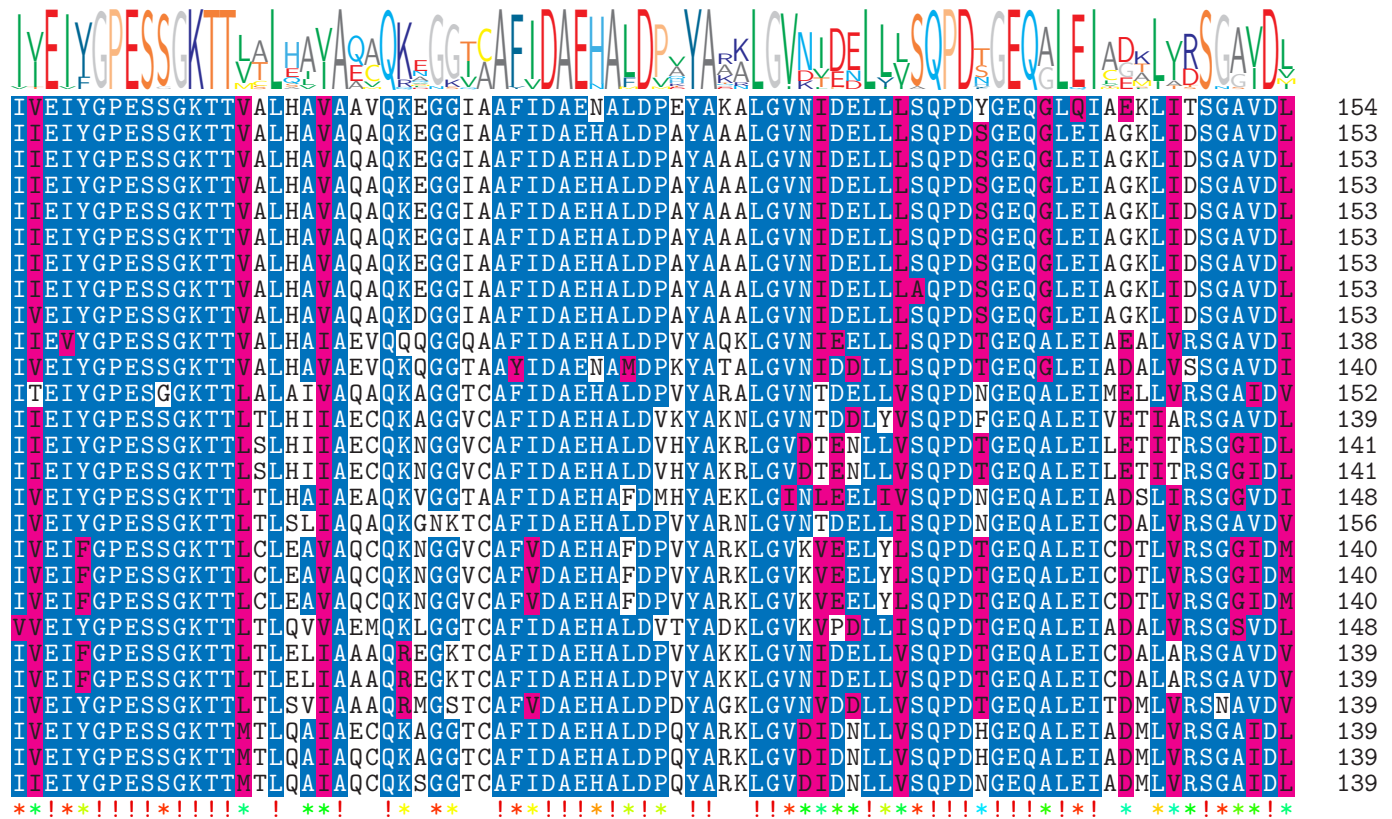
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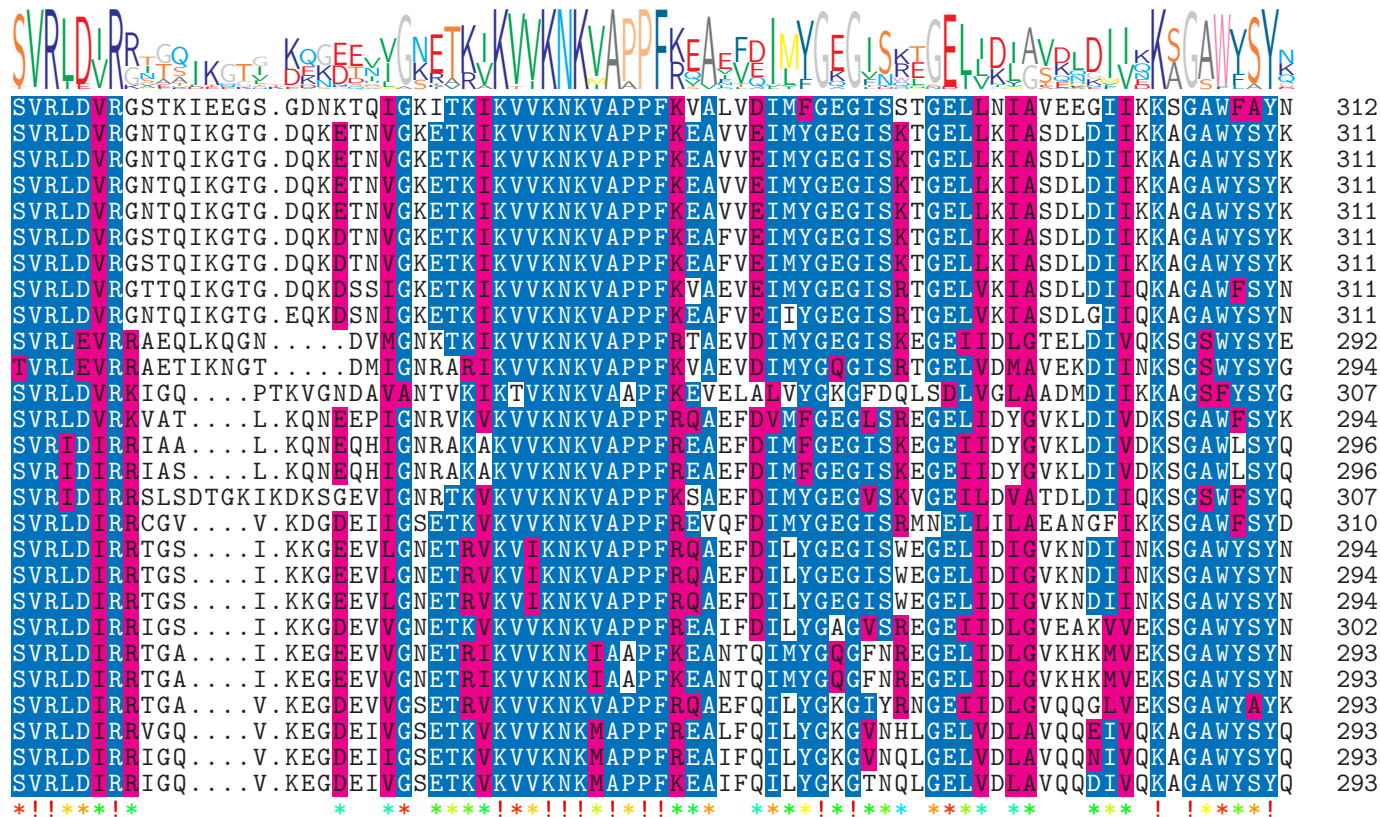
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consensus

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Sequence	Score
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DEKIGGGS E ENAKKYLAEHPEIF E IDKQVRSKFG L IDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGGS E ENAKKYLAEHPEIF E IDKQVRSKFG L IDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGGS E ENAKKYLAADNPEIF E IDHQVRVQFGL L IDGEEIAES....KKDE....AAQVDS..VNEEVPLDLGDELEIE	381
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logo

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consensus			

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