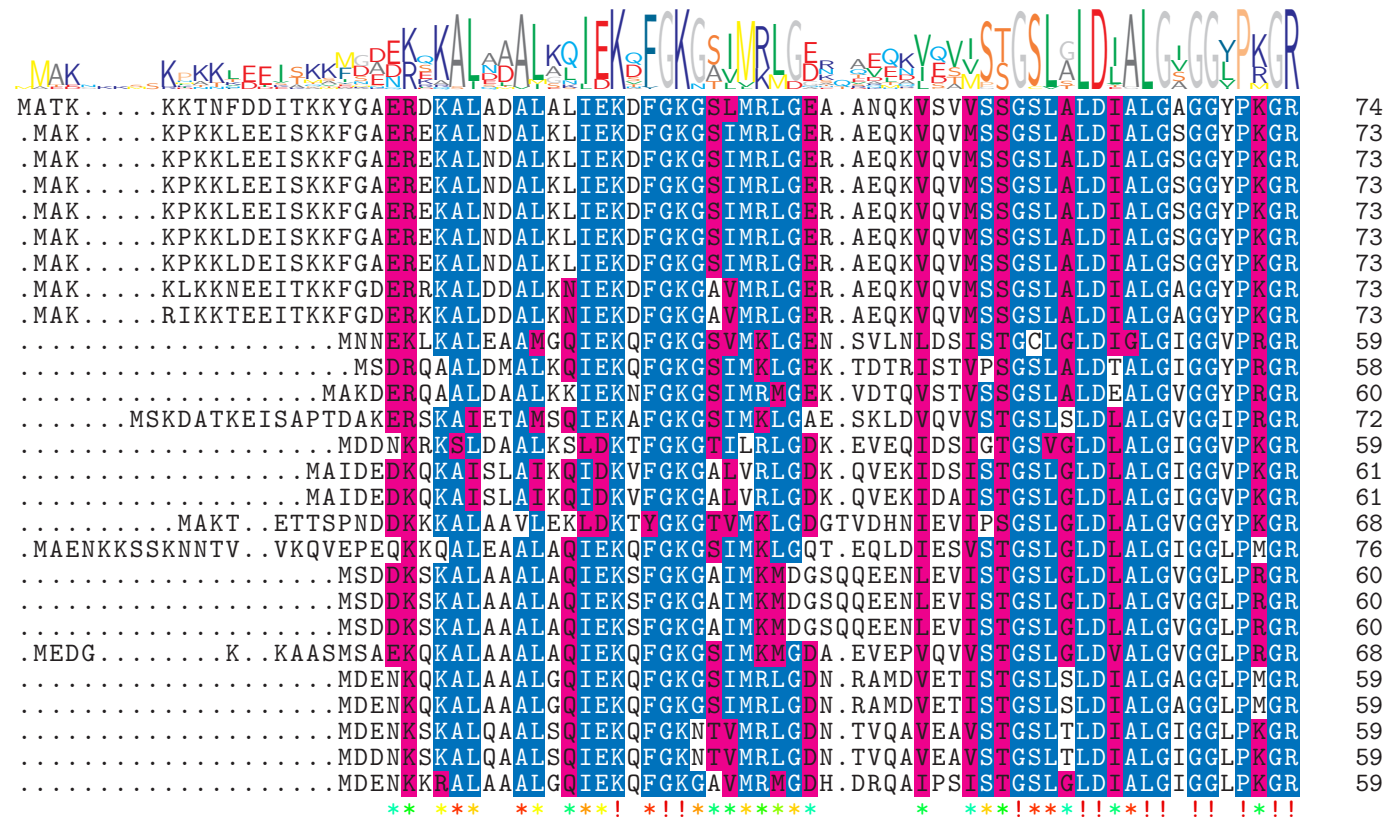


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consensus

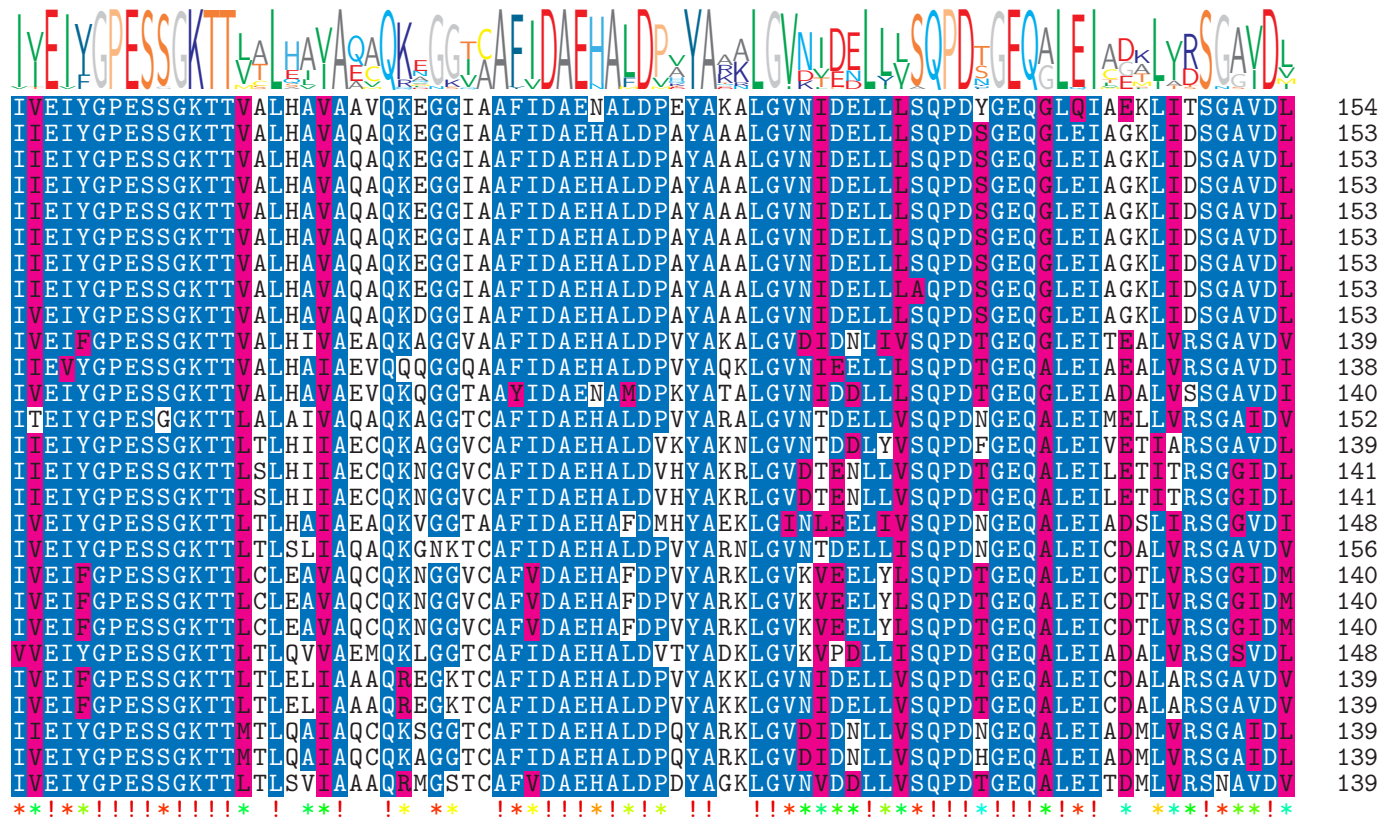
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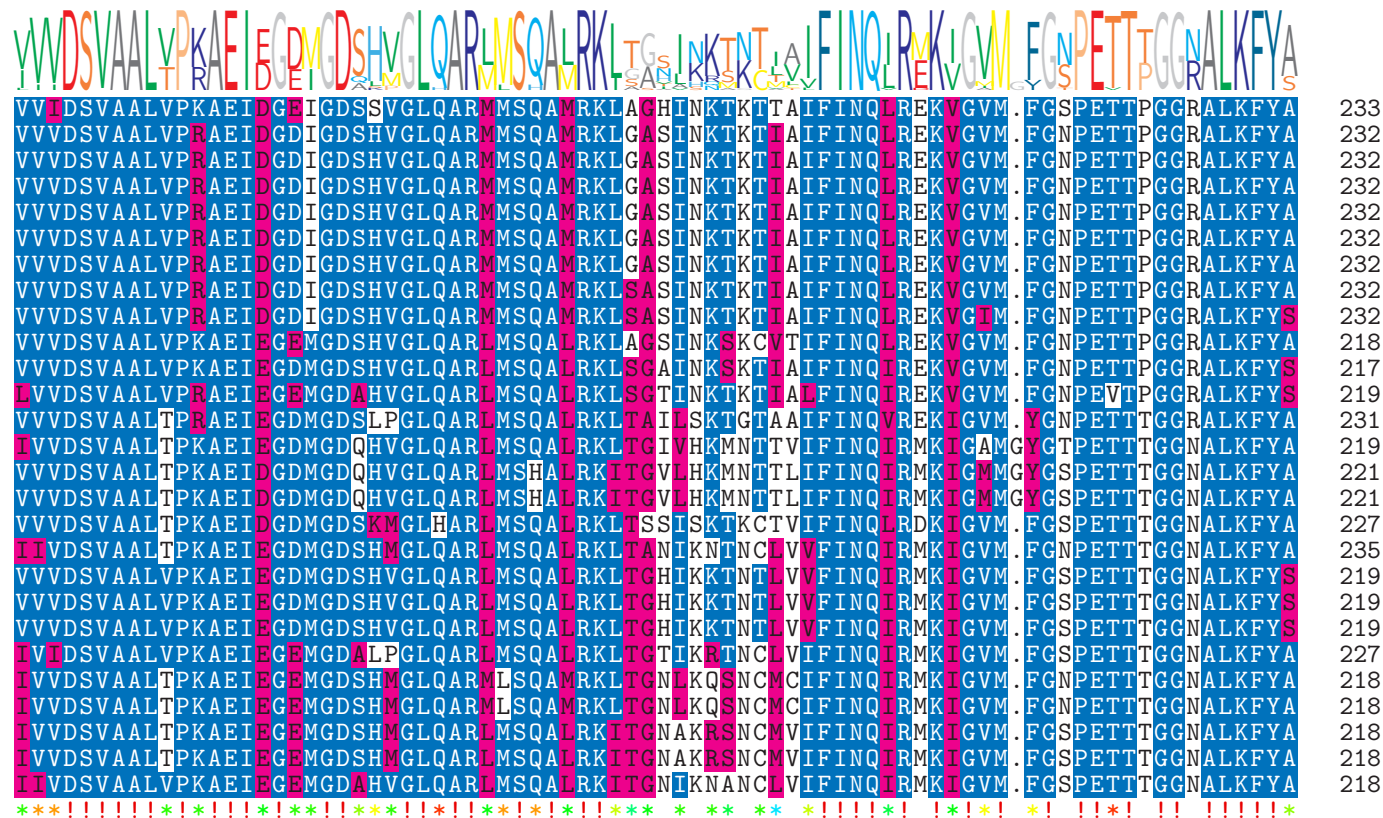
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consensus

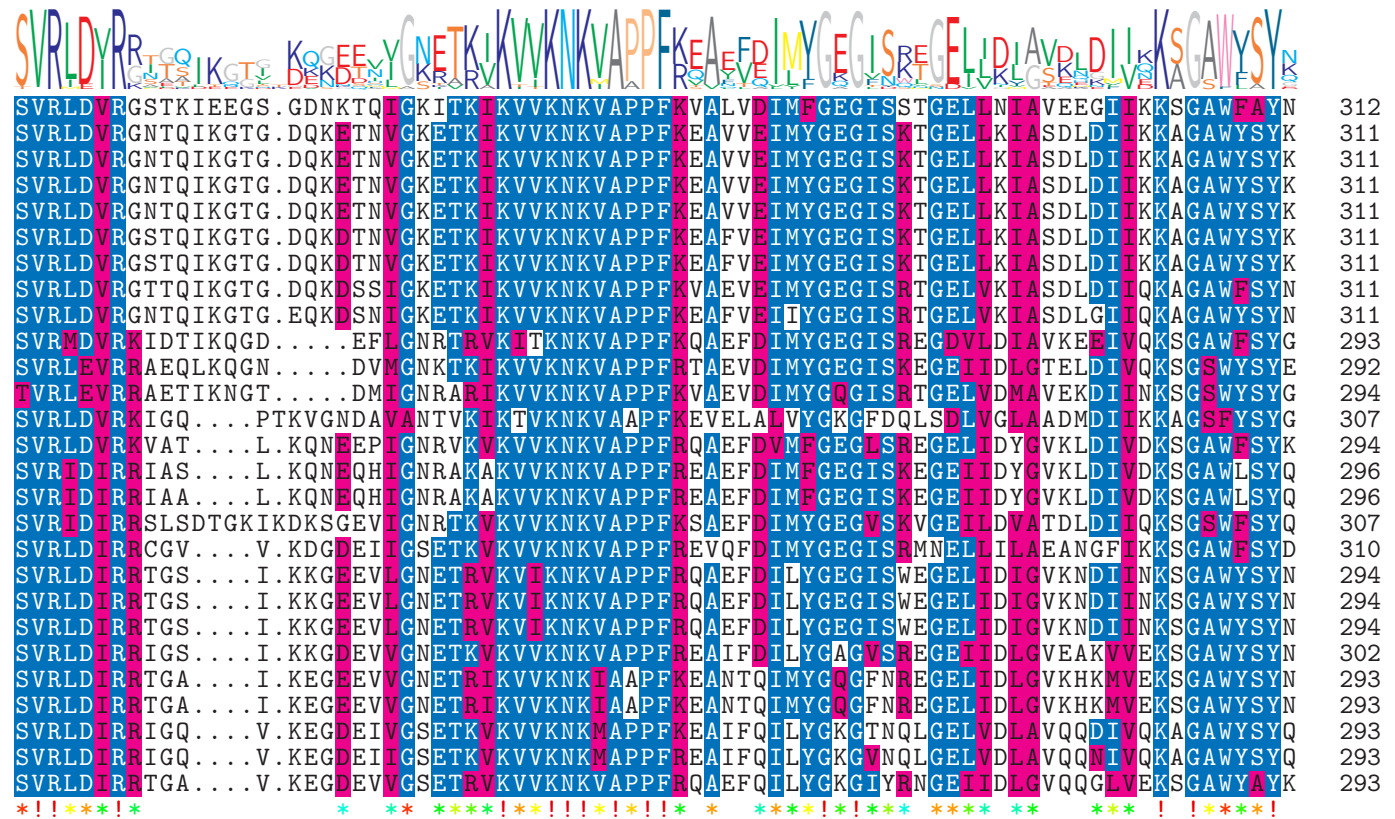
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consensus

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consensus

Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE PKKEEA . . VNEEVPLDLGDELEIE	385
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EERLGGGRENAKQFLKENKIDMLMIQEIQIREHYGLDNNGVVQQAAETQEEL . EFE . E	348
SERIGGGRENAKNYLADHEDVEVRLKVRVQYGLISDVPEEDLPTTEDE . QI . NILPDD STEE	355
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logo

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consensus			

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