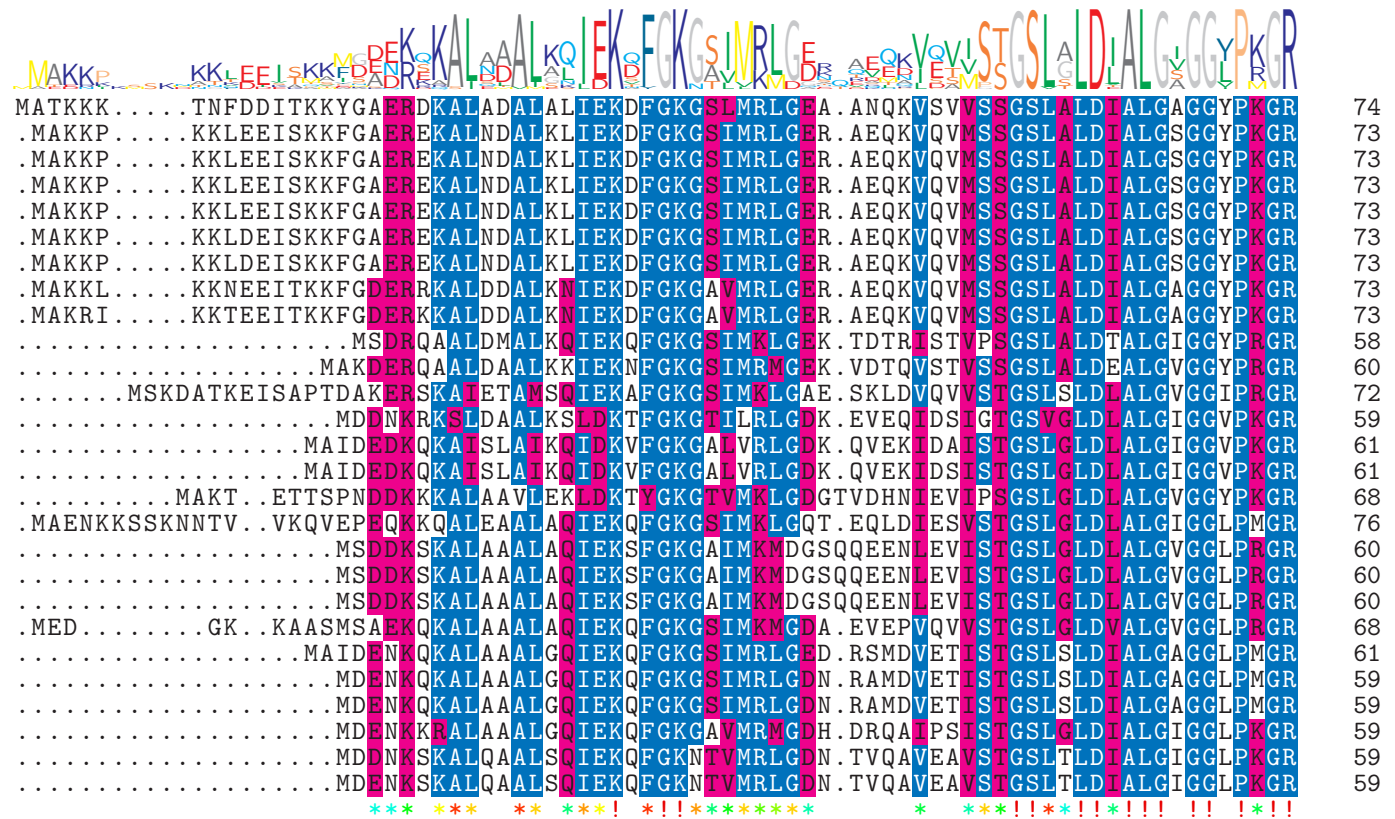


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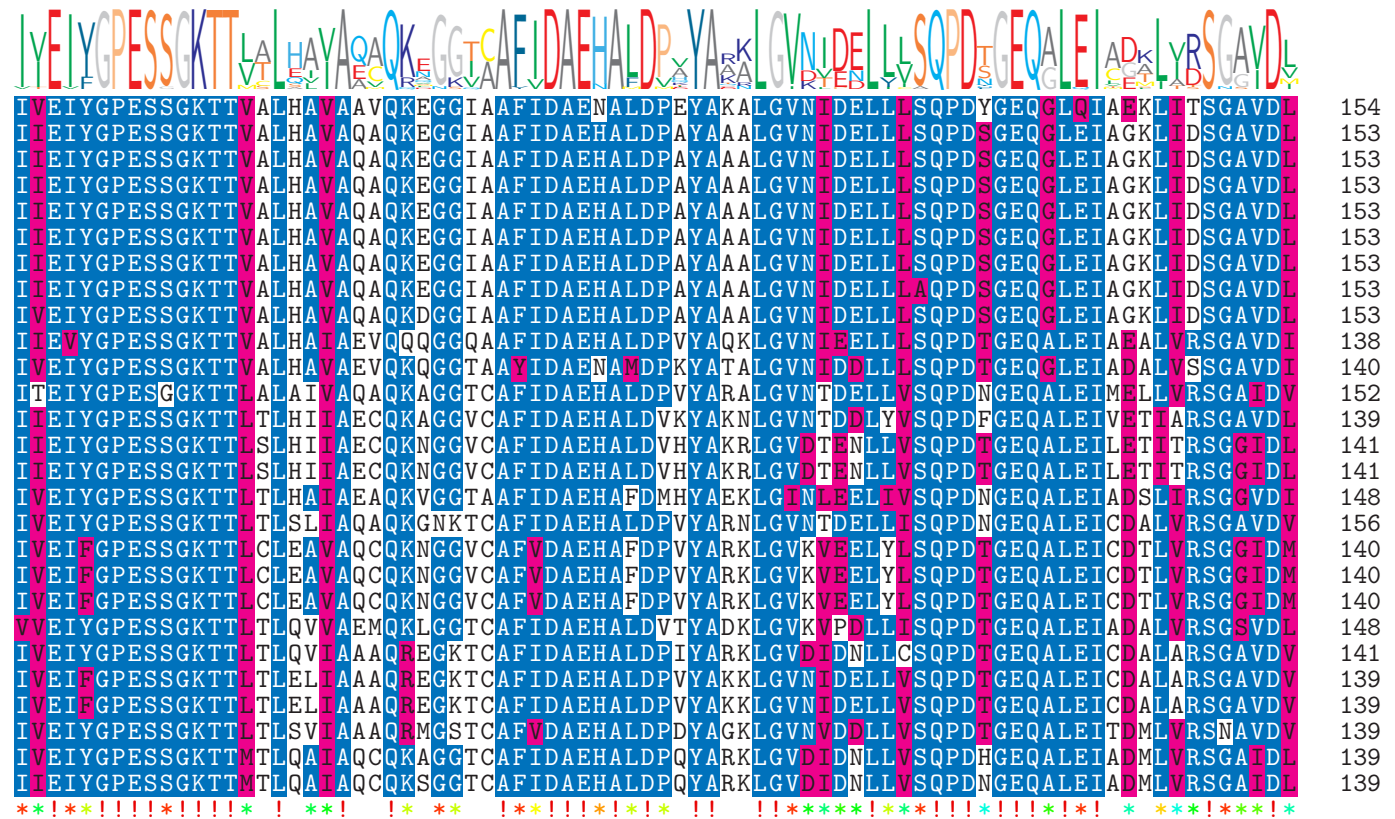
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consensus

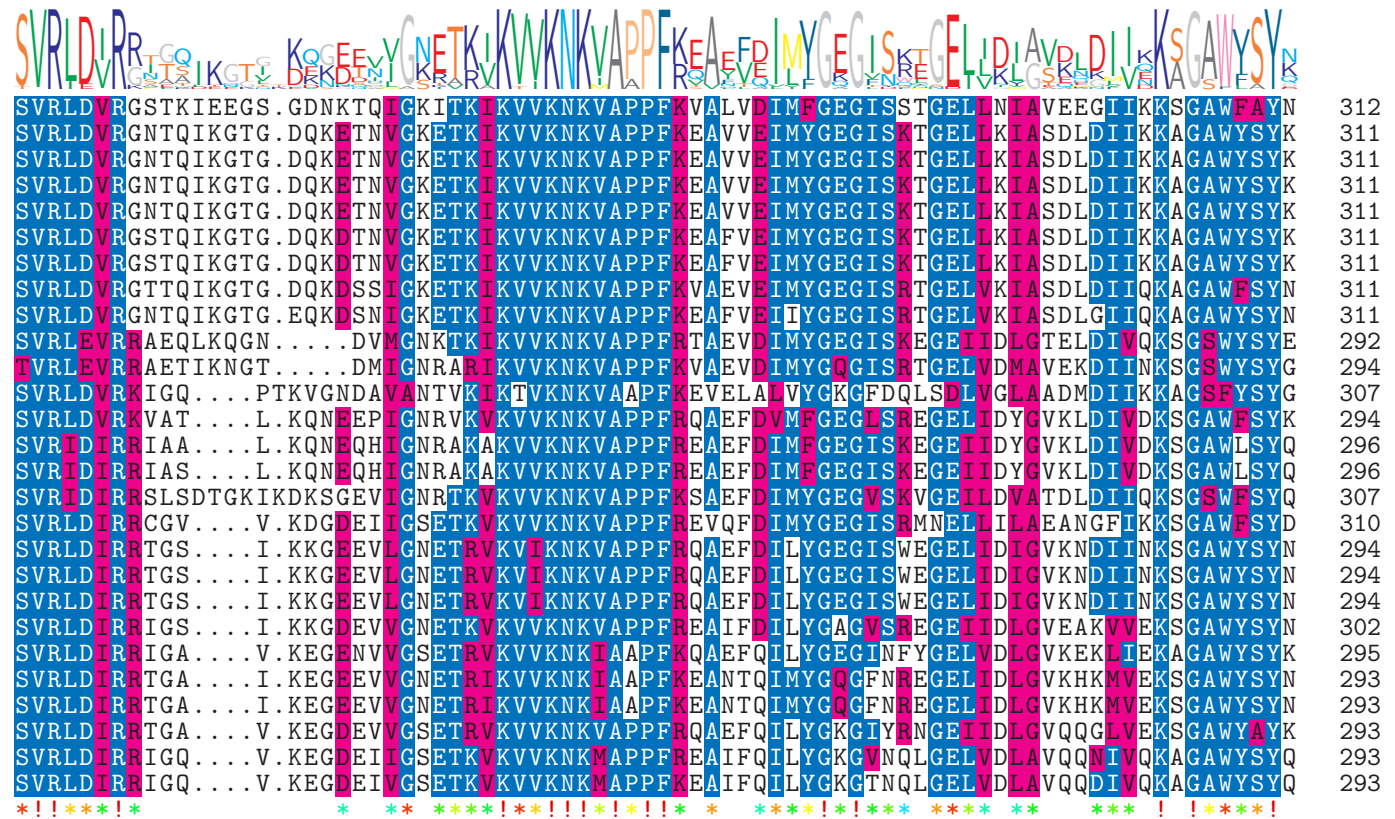
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consensus

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consensus

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Sequence	Score
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EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL . EFE . E	348
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consensus

 non conserved
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 ≥ 50% conserved