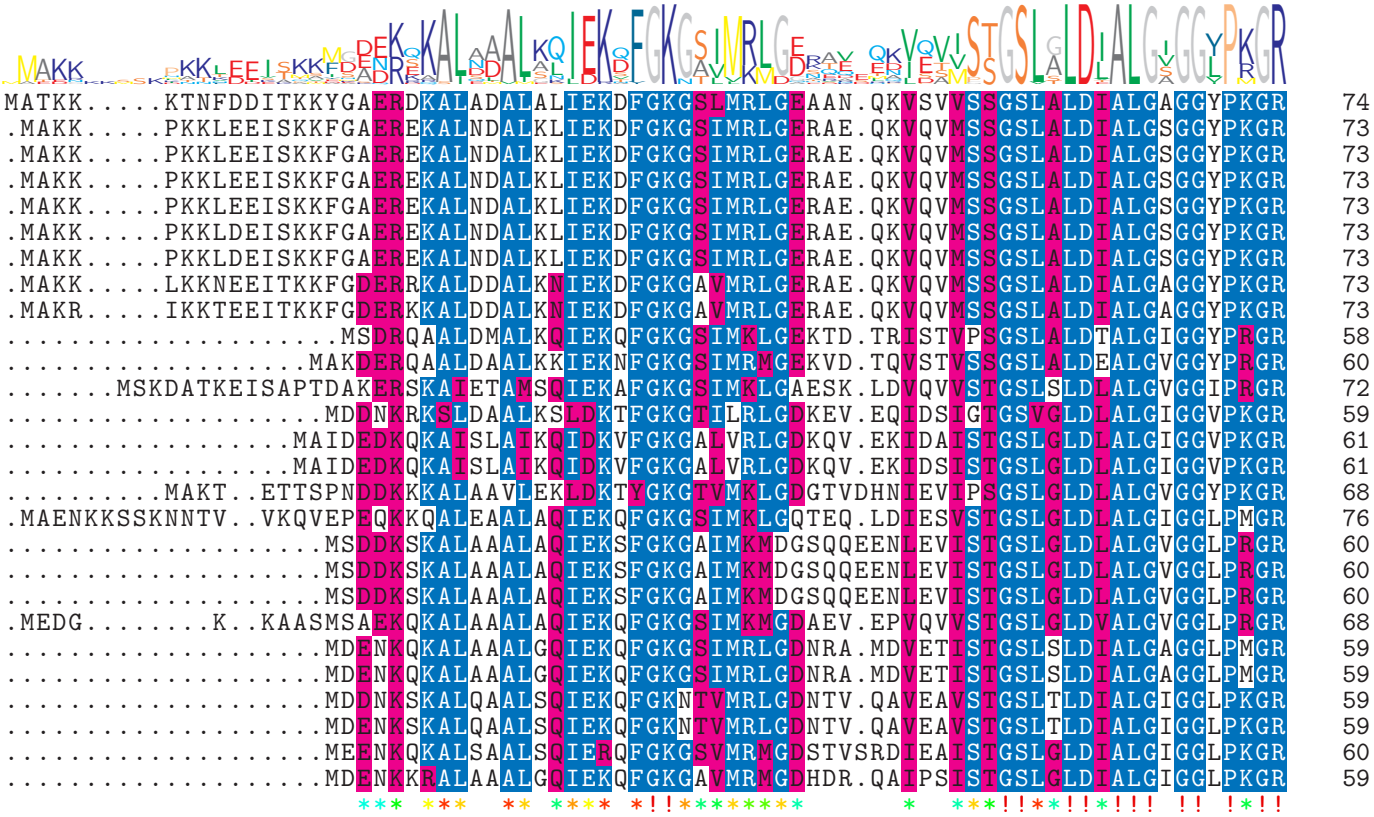


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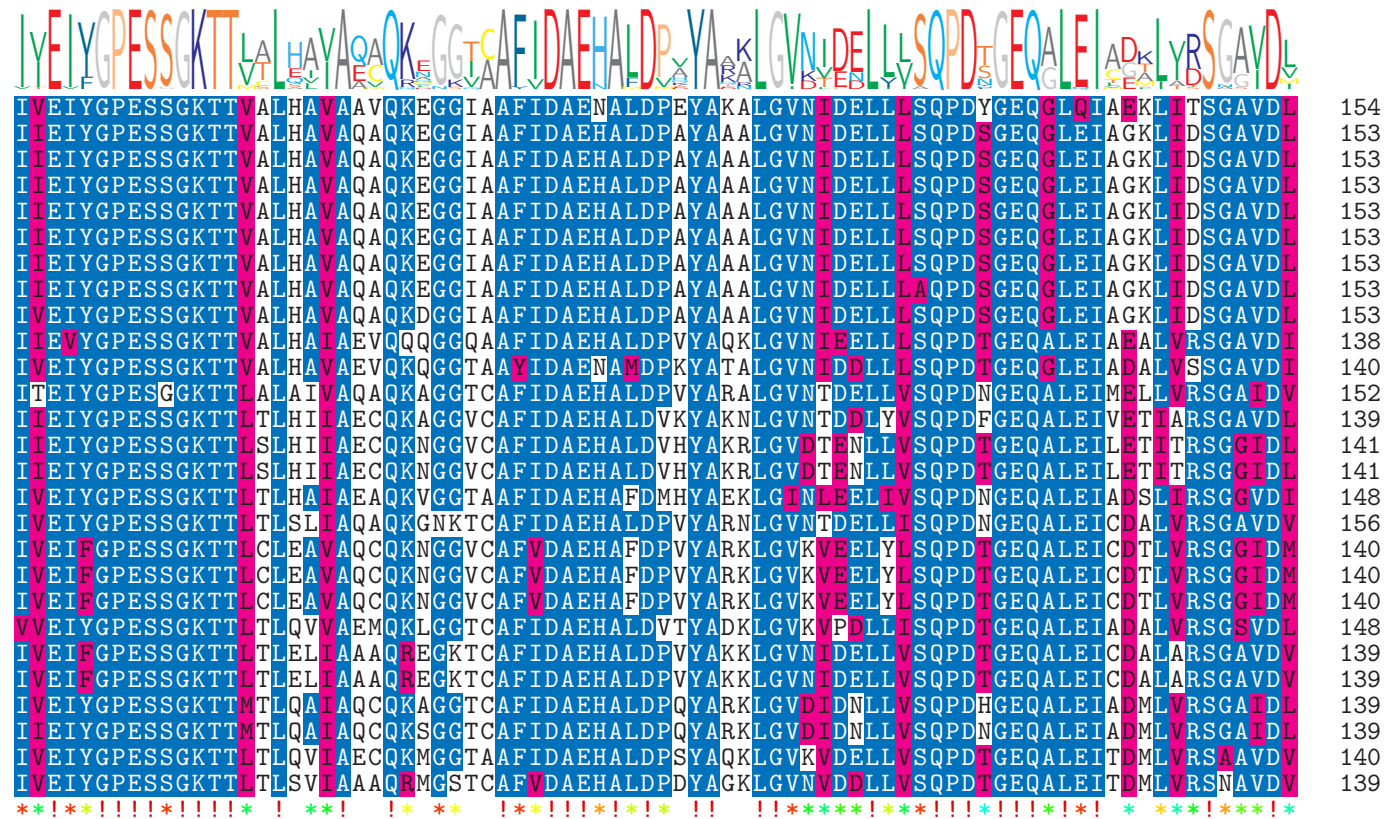
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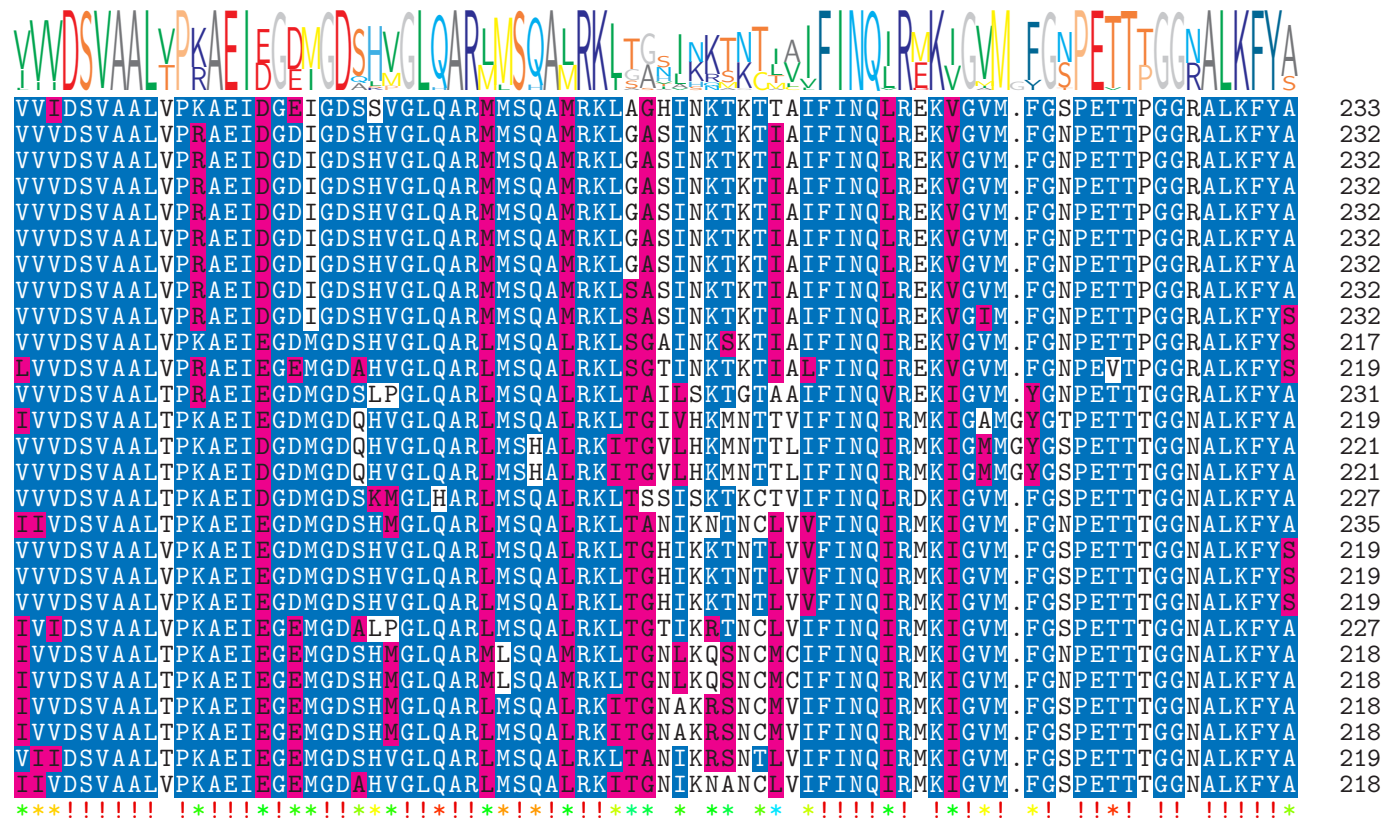
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DKKLGGGRENAAKALLKEDKALADEITLKIKESIGSNEEIM PLPD . EPLE . EME	347
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GNKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTASDQ TAAHDET . EE . EPDLL . ES	349
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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved