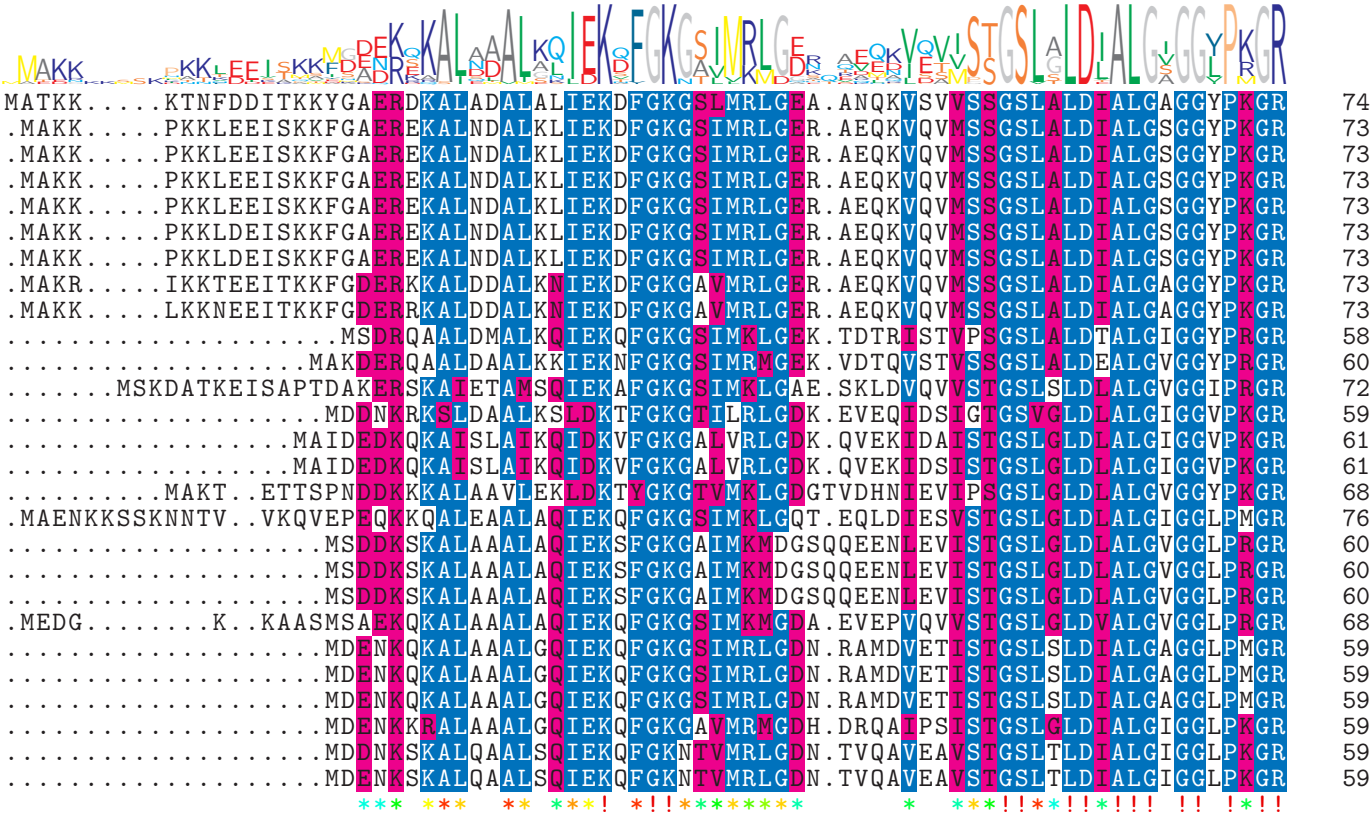


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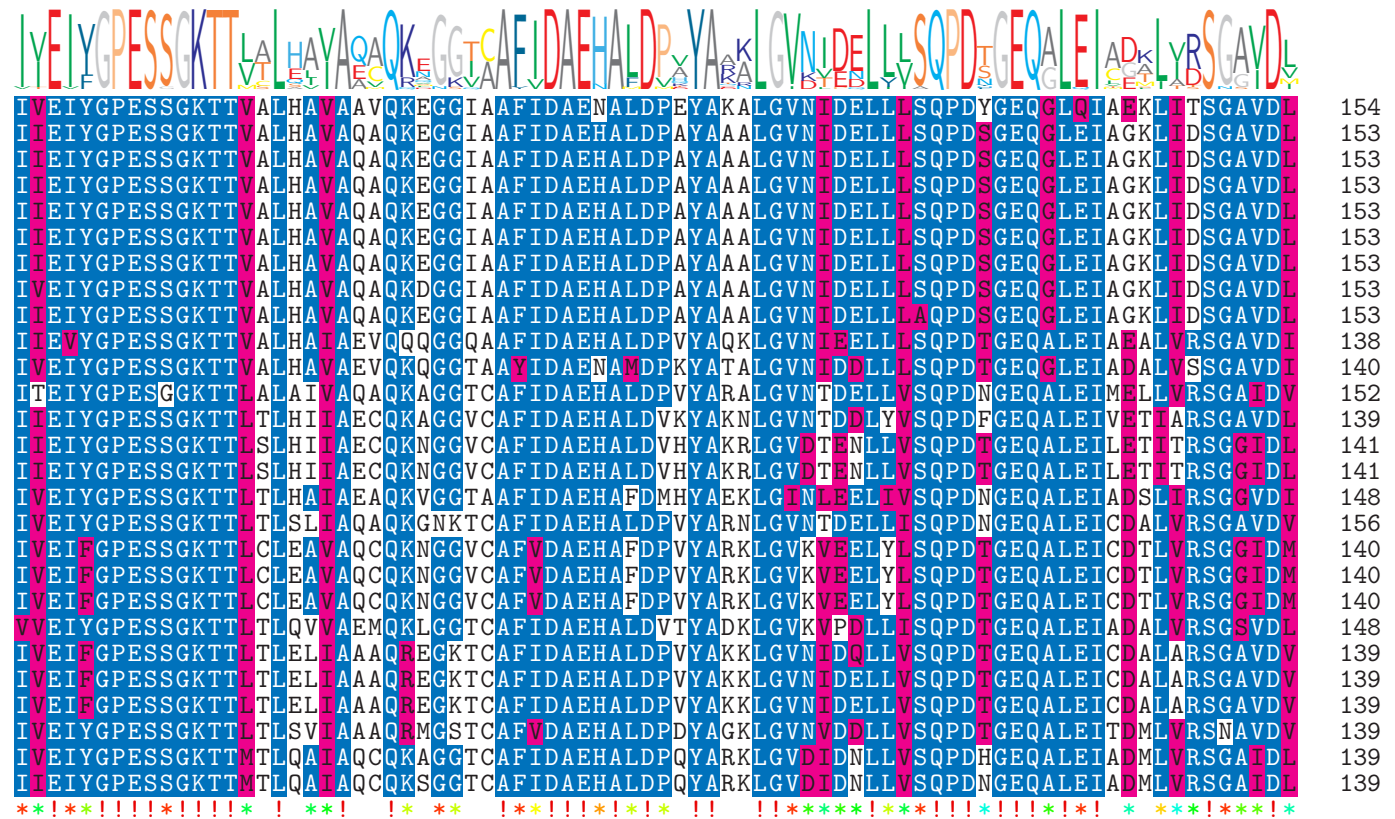
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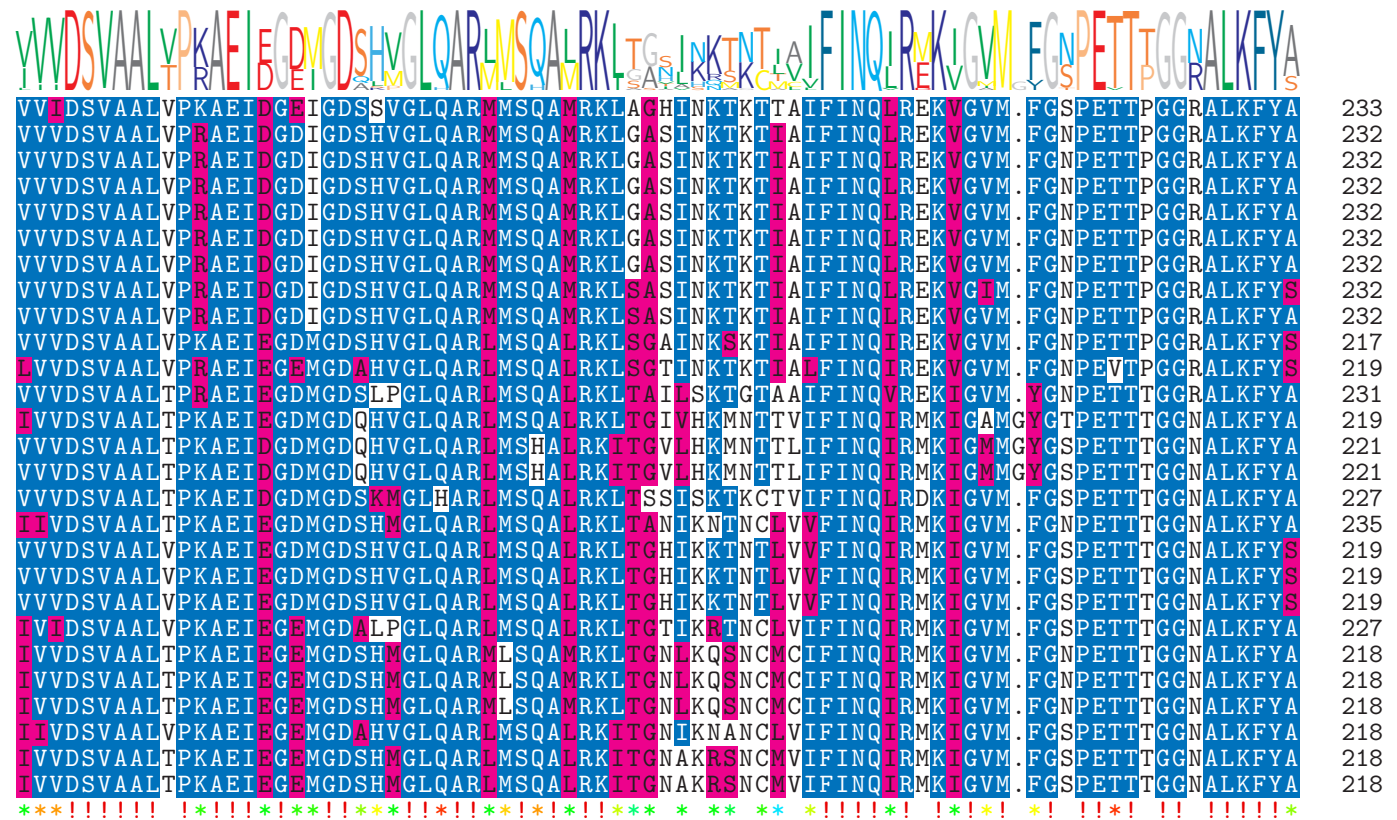
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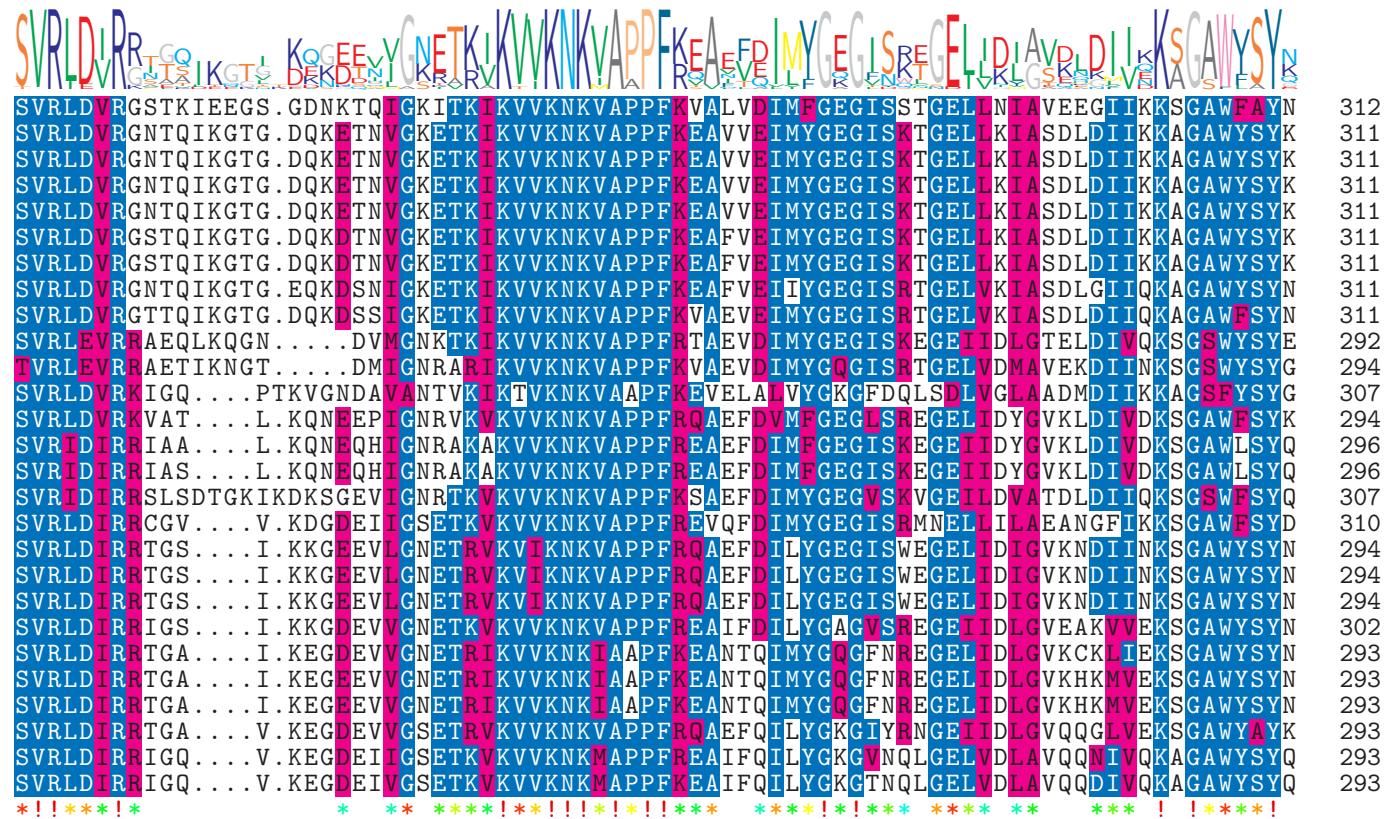
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consensus

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GEKIGQGSENAKFLADNPVFDIDHKVVRVQGLIEEDNTEEEKQSSKEK ETDEK . . ADKNLVLELDDTTIELE	382
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE E TDDLALDLNNGIEIE	377
EERLGGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL . EFE . E	348
SERIGGGRENAKNYLADHEDVEDEVRLKVRVQGLSDVPEEDLPTTEDE . QI . NILPDDST EE	355
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GSKIGQKKANAACKYLEENPEIGQIEEQIRAKLLVTSANTK . . ATPVAEDLA . . DADI	347
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GNKIGQKKNNVIRIYFENNQIAEIERNIREQLLTGTNGA . . VQ . I . EDEE . . EPDLLS	349

logo

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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved