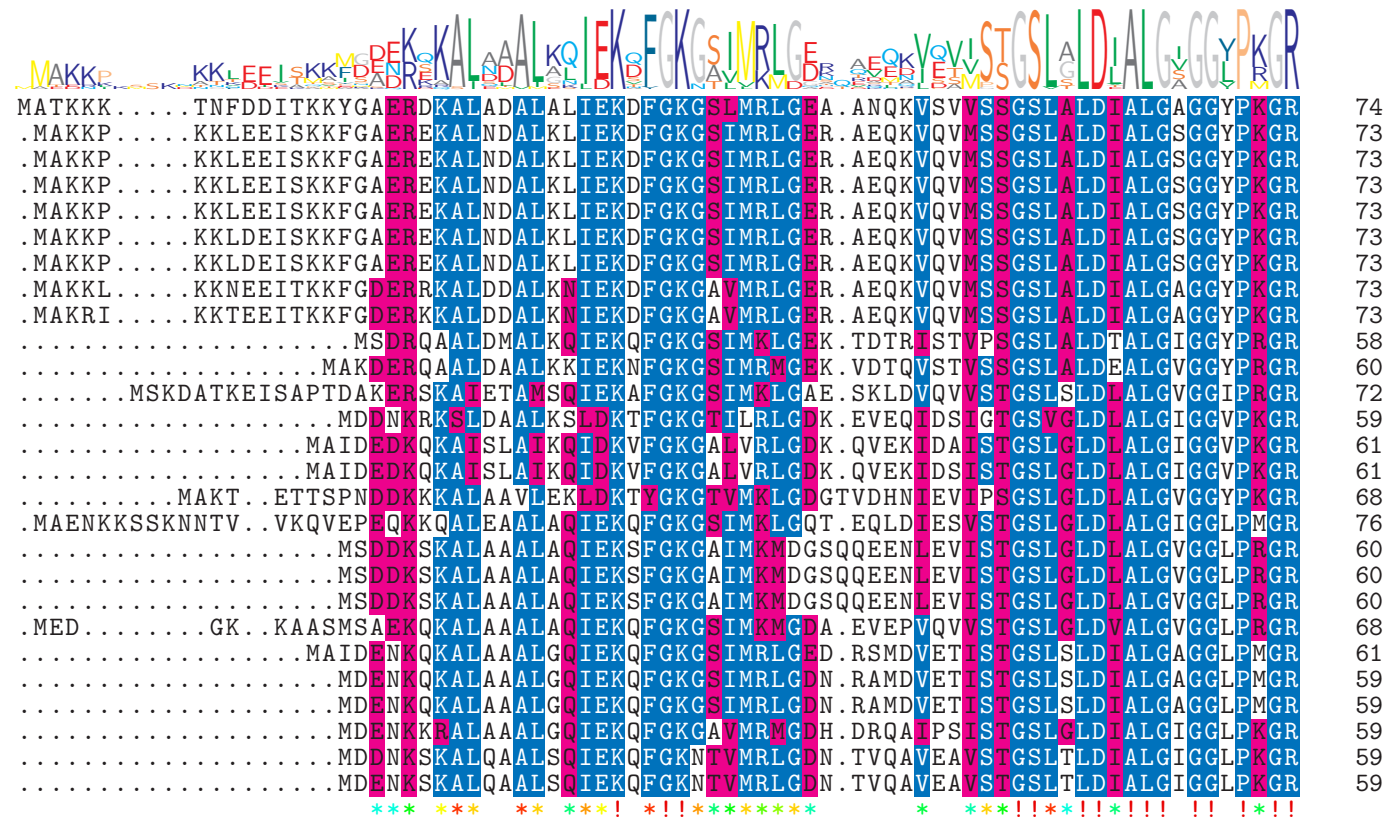


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consensus

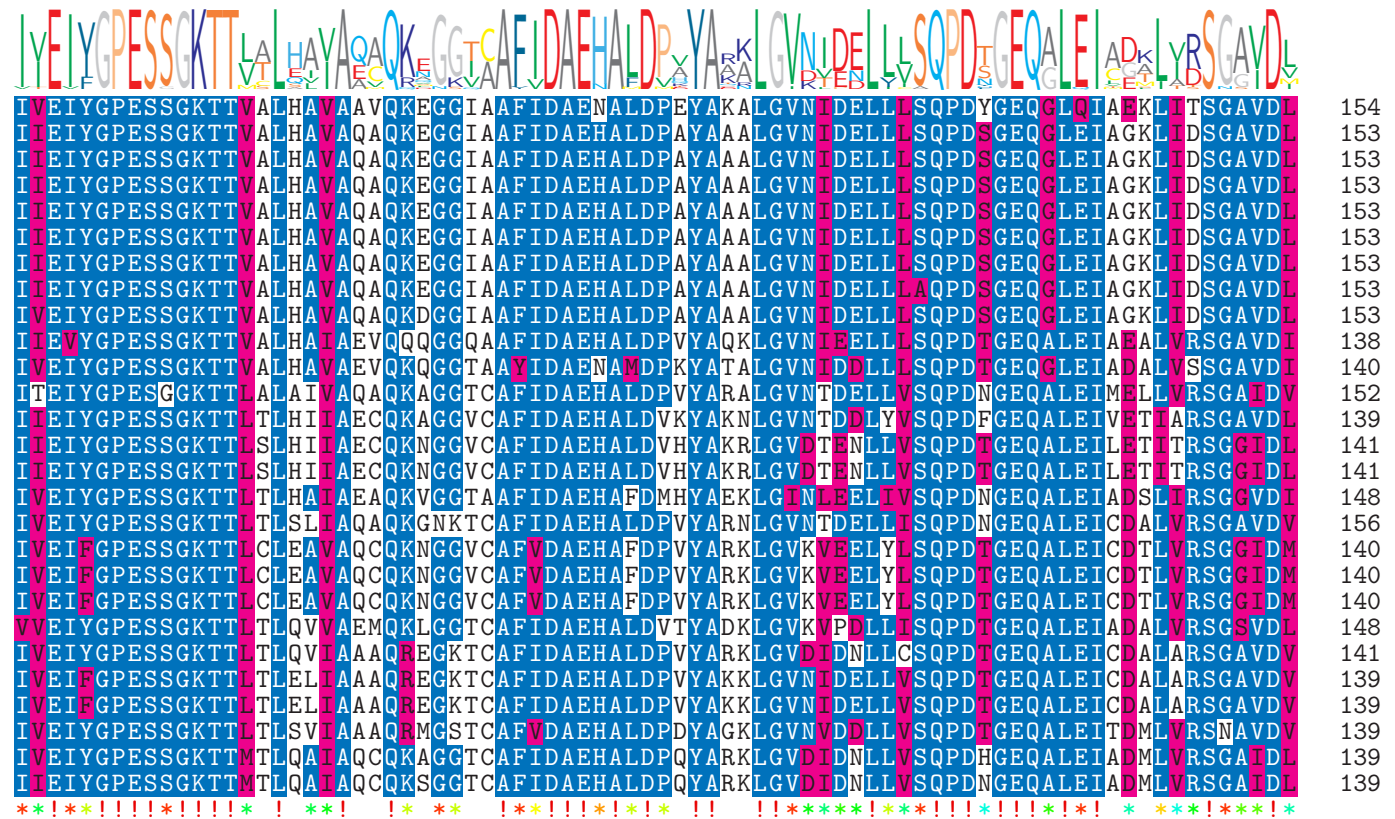
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consensus

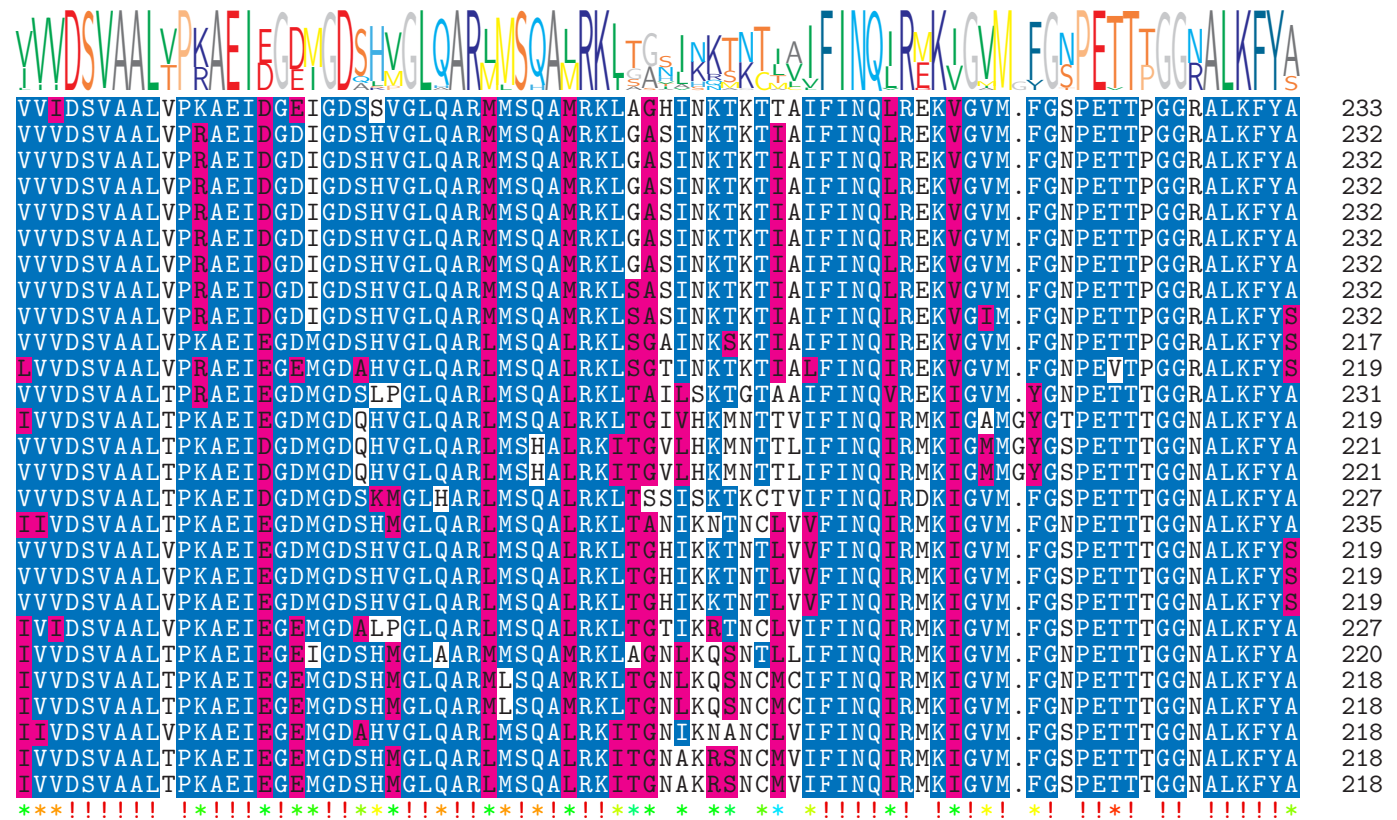
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consensus

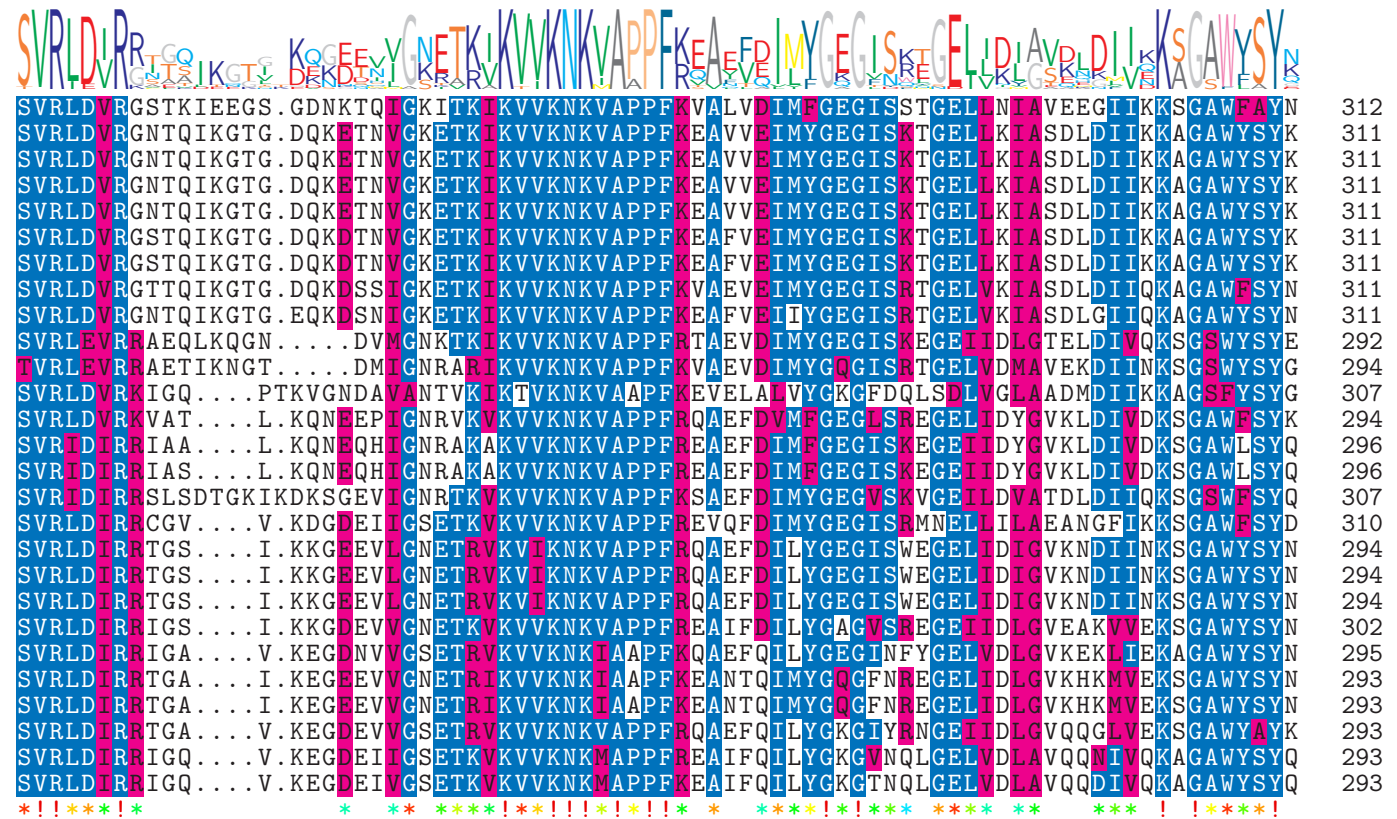
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consensus

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consensus

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GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
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DERIGGQKEKTIAYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAA...EA.....	363
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GSKIGGQKANAACKYLEENPEIGQIEEQQIRAKLLVTSA....NTKATPVAEDL..ADADI.....	347
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logo

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consensus

 non conserved
 similar
 ≥ 50% conserved