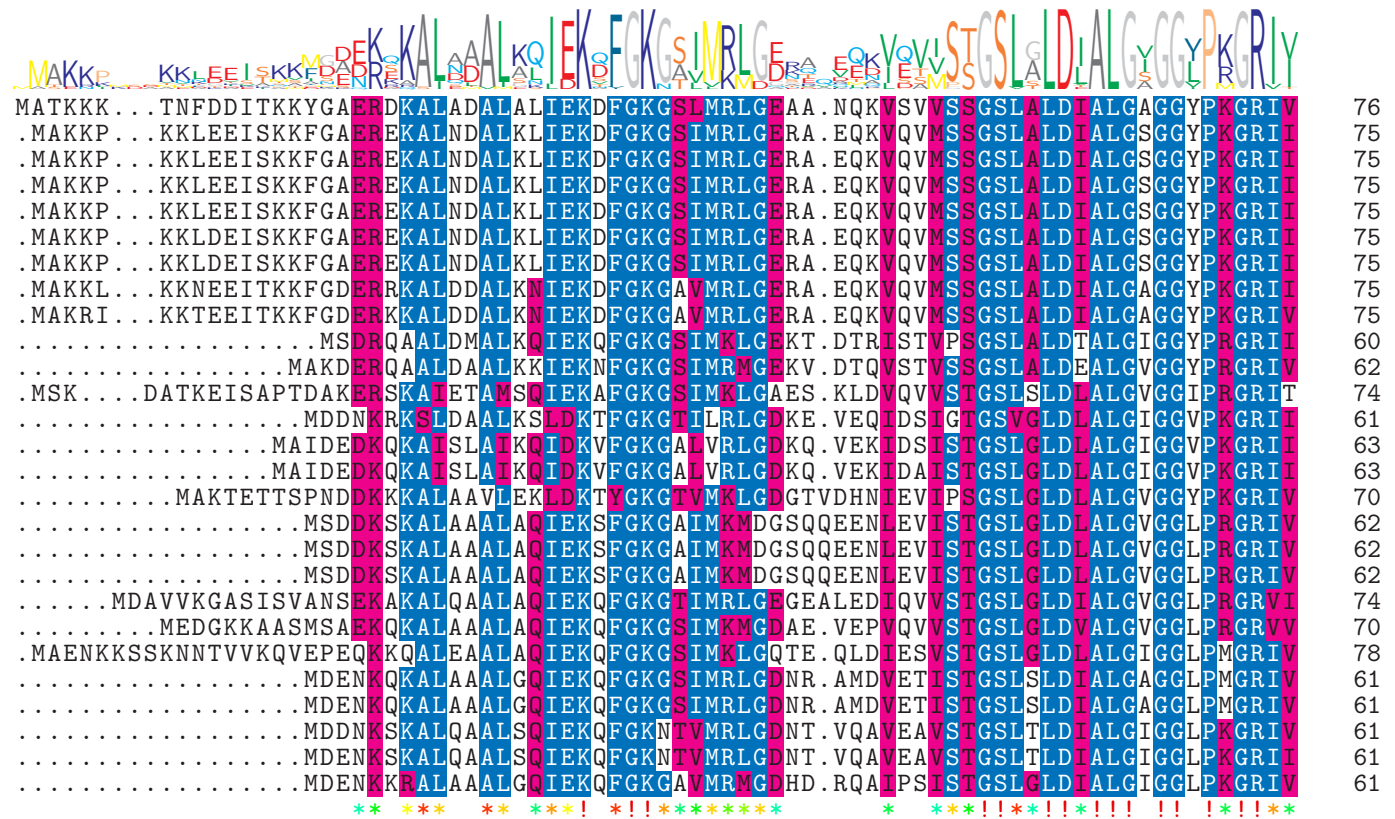


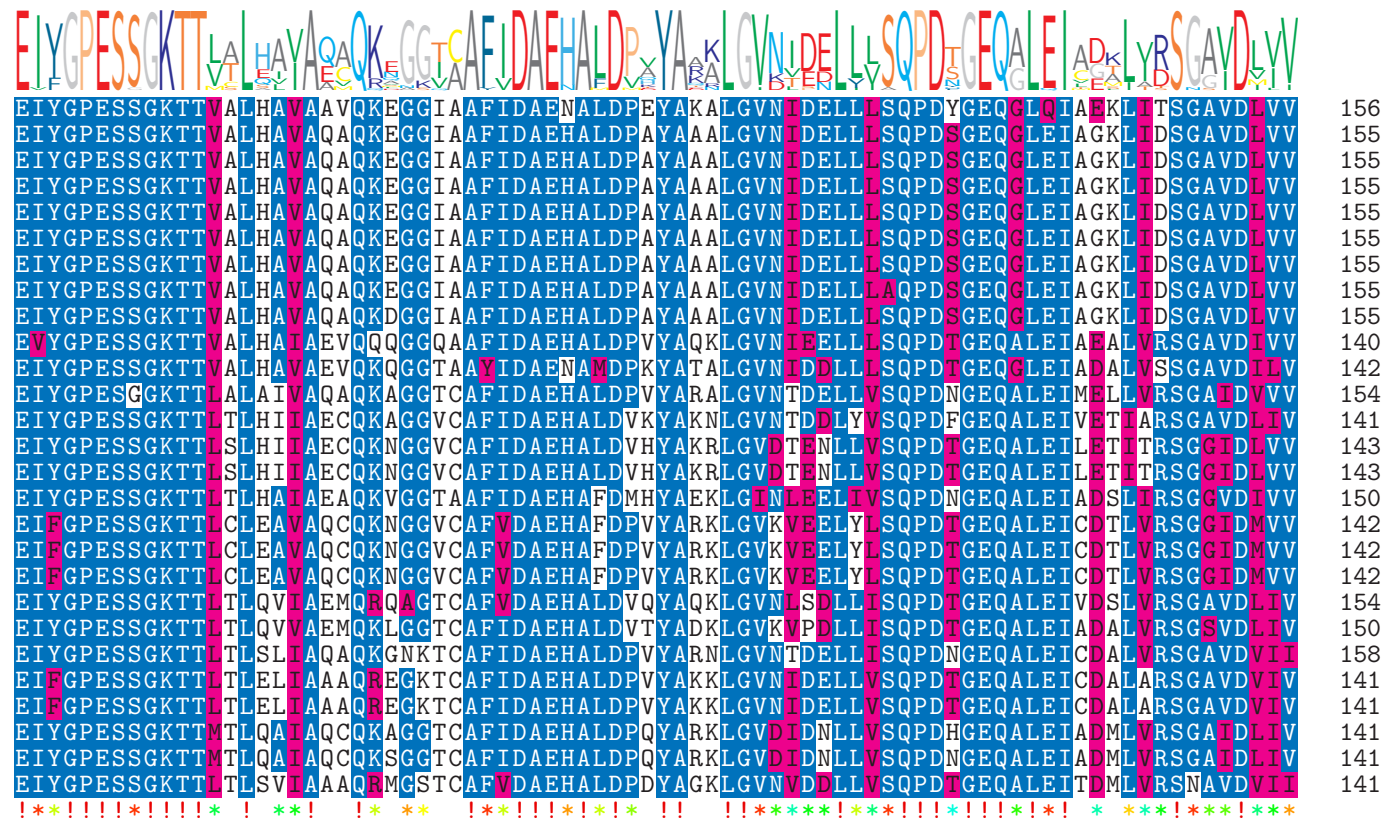
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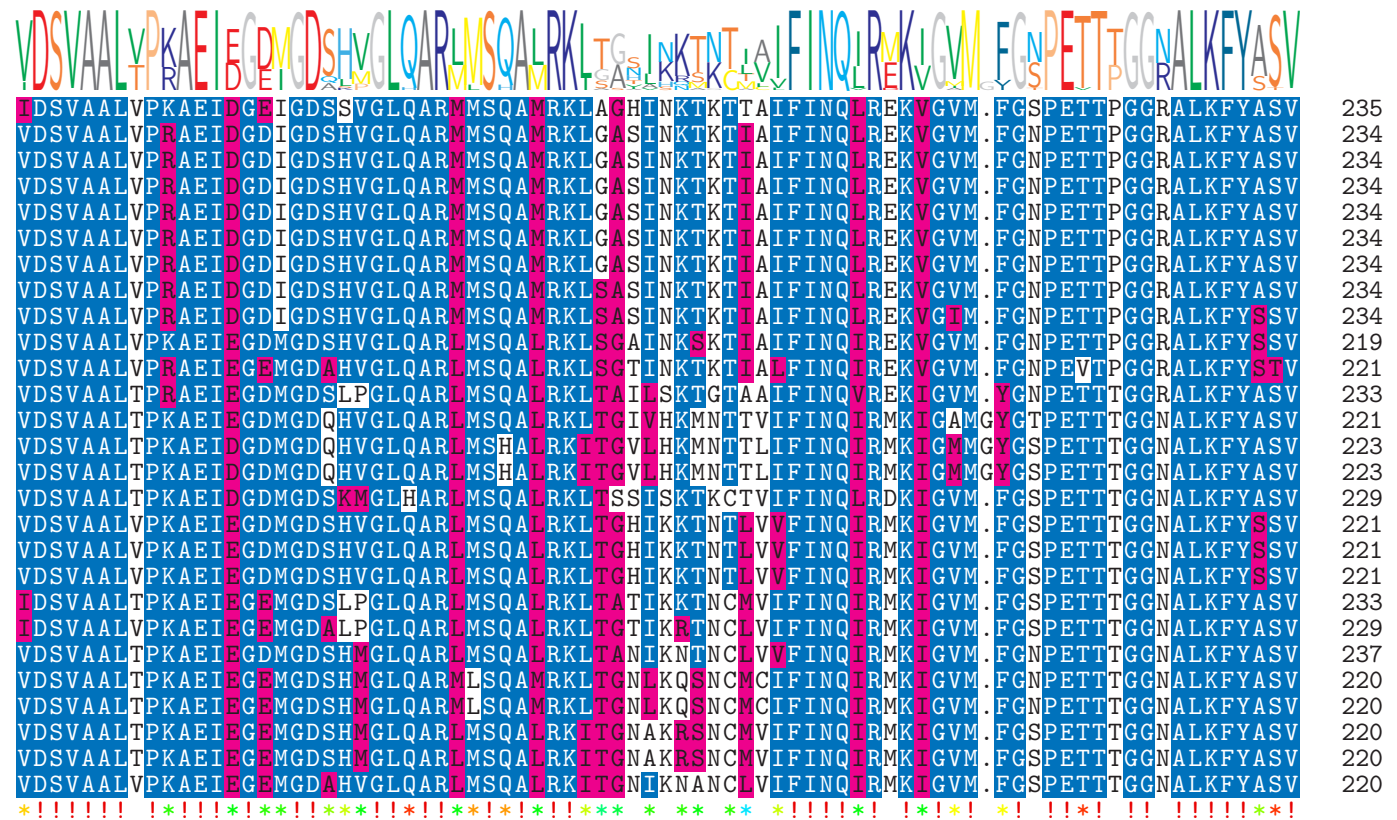
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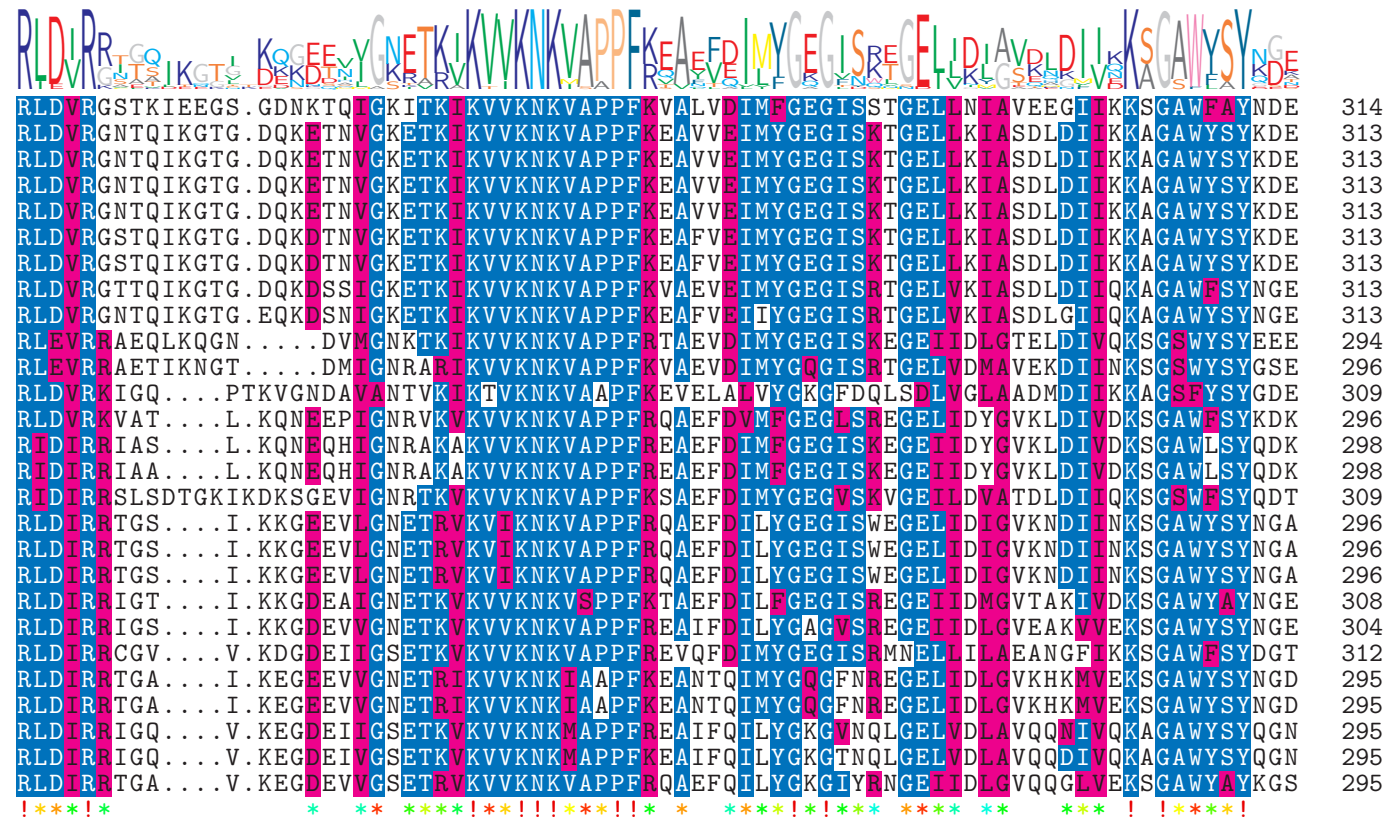
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consensus

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KIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQD TENKKDE....PKKKEA..VNEEVPLDLGDELEIEIE	387
KIGQGSSENAKKYLAHNPEVFAEIDHQVRVQFGLIDGEEAAE....SKKDE....VAQVDS..VNEEVTLDLGDLEIEIE	383
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KIGQGSSENAKRYLADHPQLFDEIDRKVRVKFGLLEESSEESAMAVASE.....E.....TDDLALDLNGIEIED	378
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logo

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consensus

 non conserved
 similar
 ≥ 50% conserved