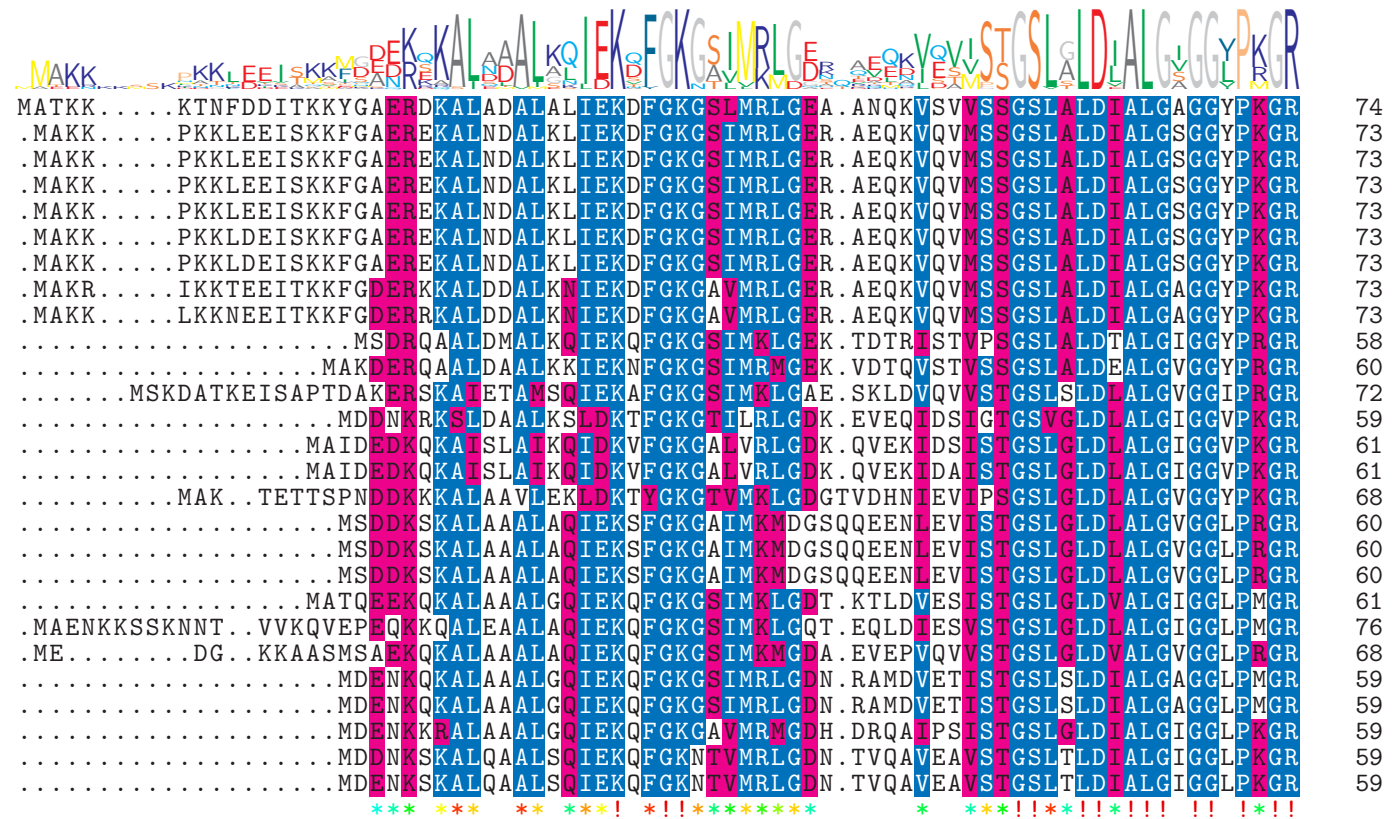


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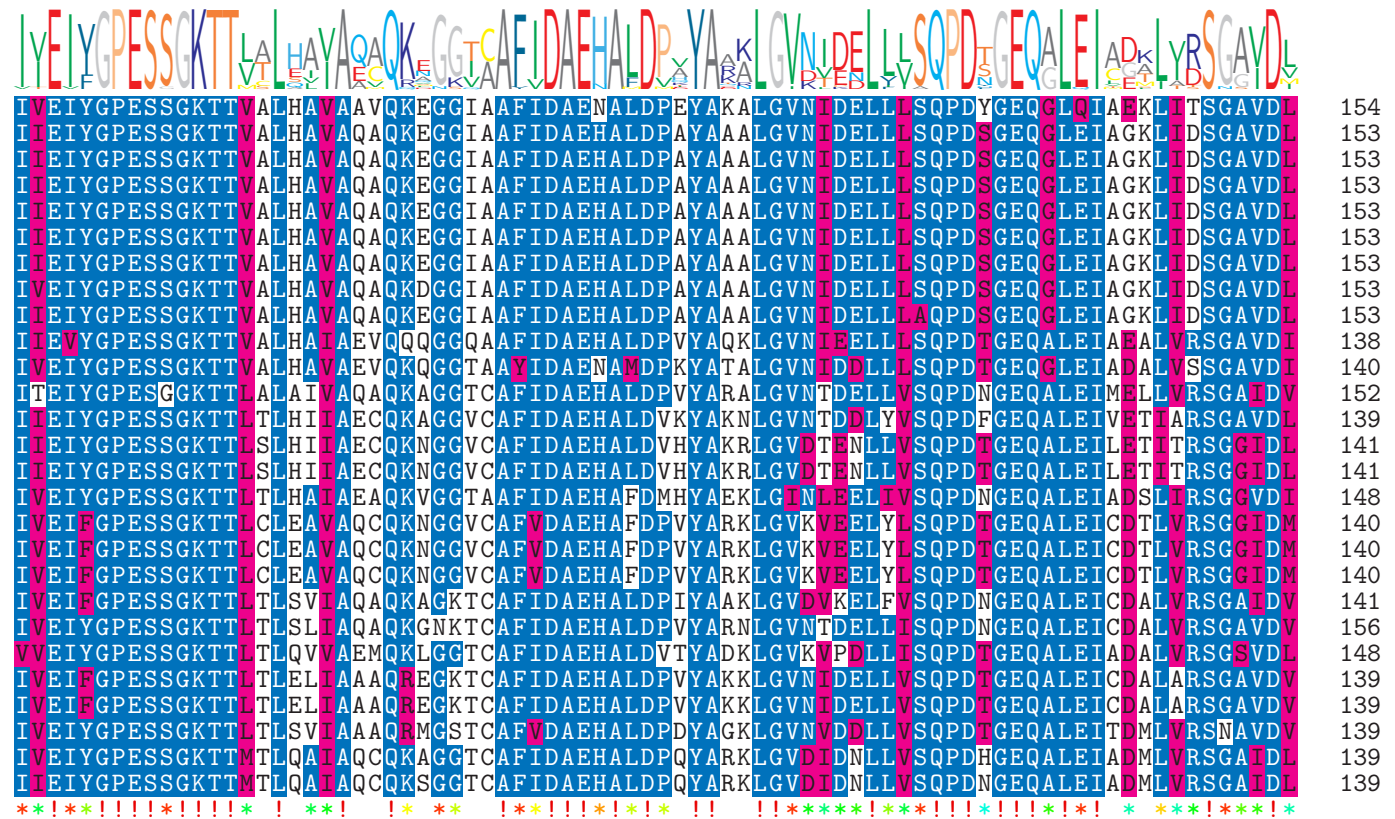
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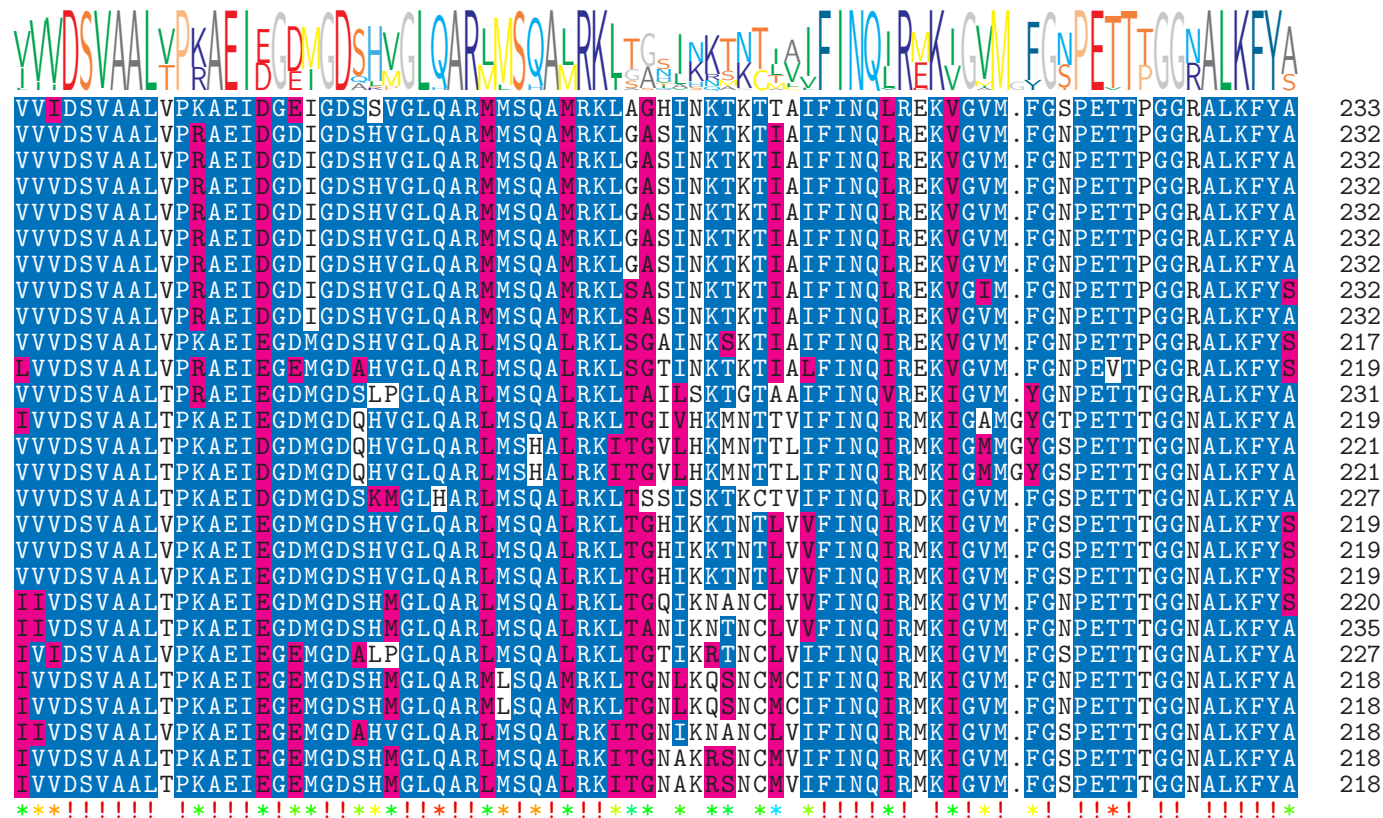
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consensus

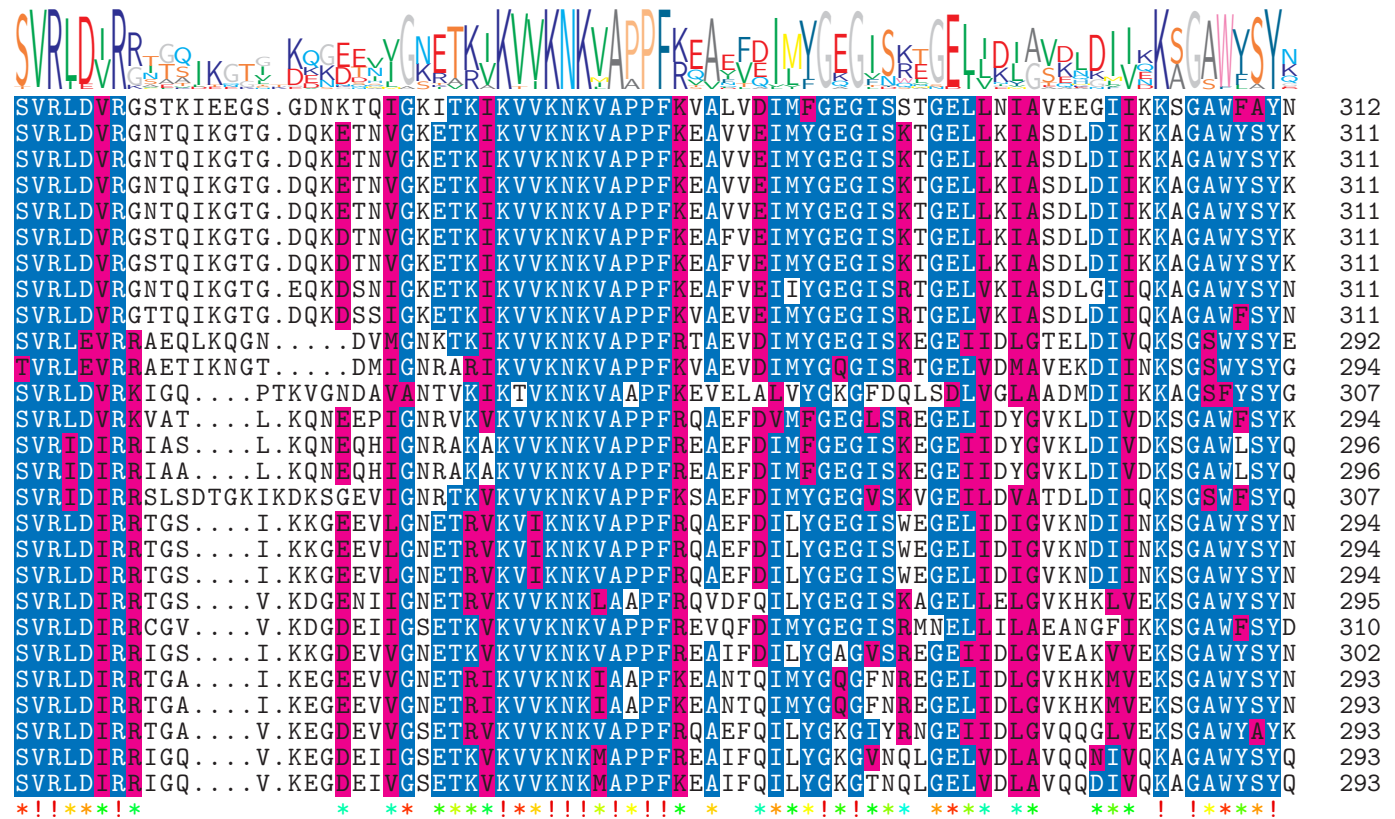
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consensus

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GEKIGQGS ENAKRYLA DHPQLFDEIDRKVRVKFGLLEESEESAM.AVASE.....E.....TDDLALDLDNGIEI 376
EERLGQGR ENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQA.EETQEEL.EFE.E..... 348
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DERIGQG KEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAEA..... 363
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consensus					

-  non conserved
-  similar
-  ≥ 50% conserved