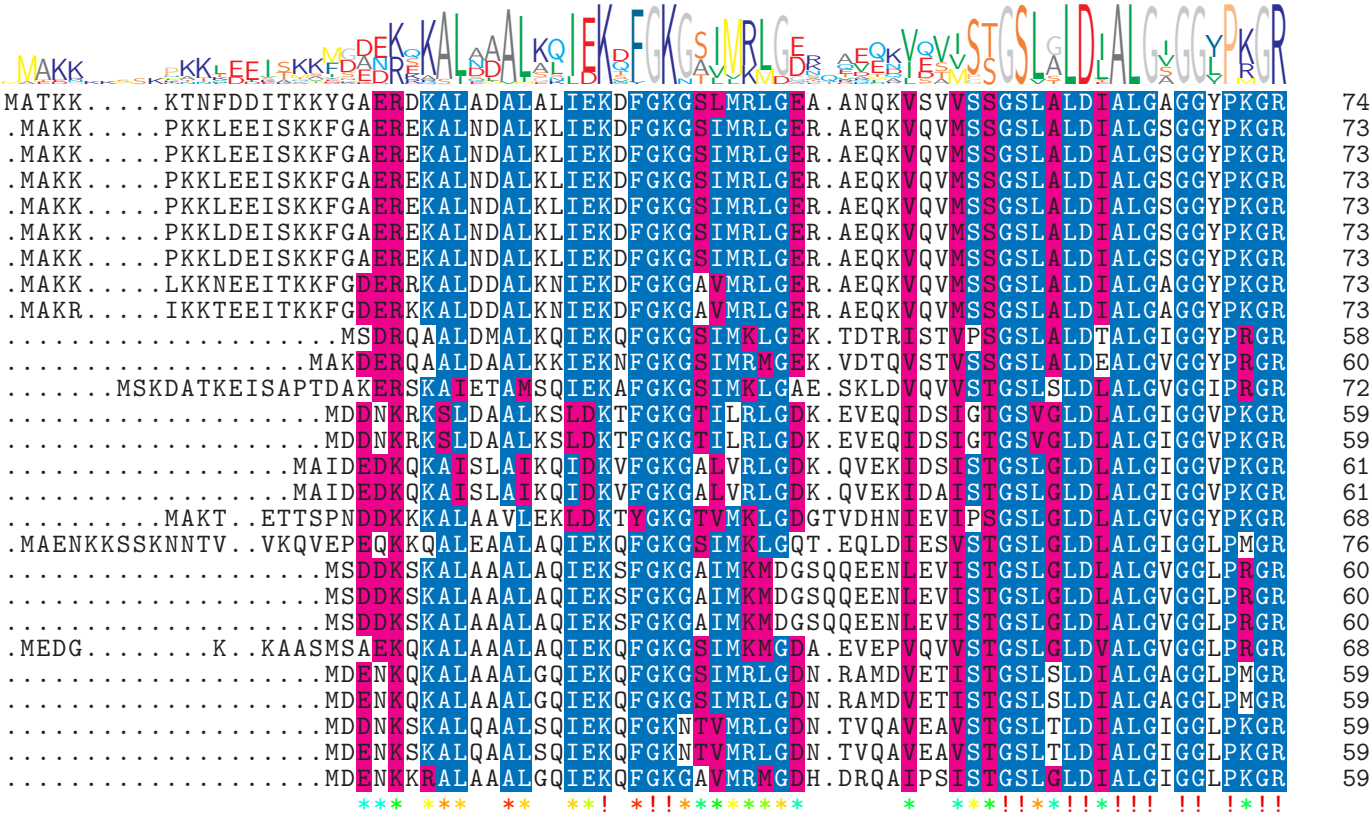


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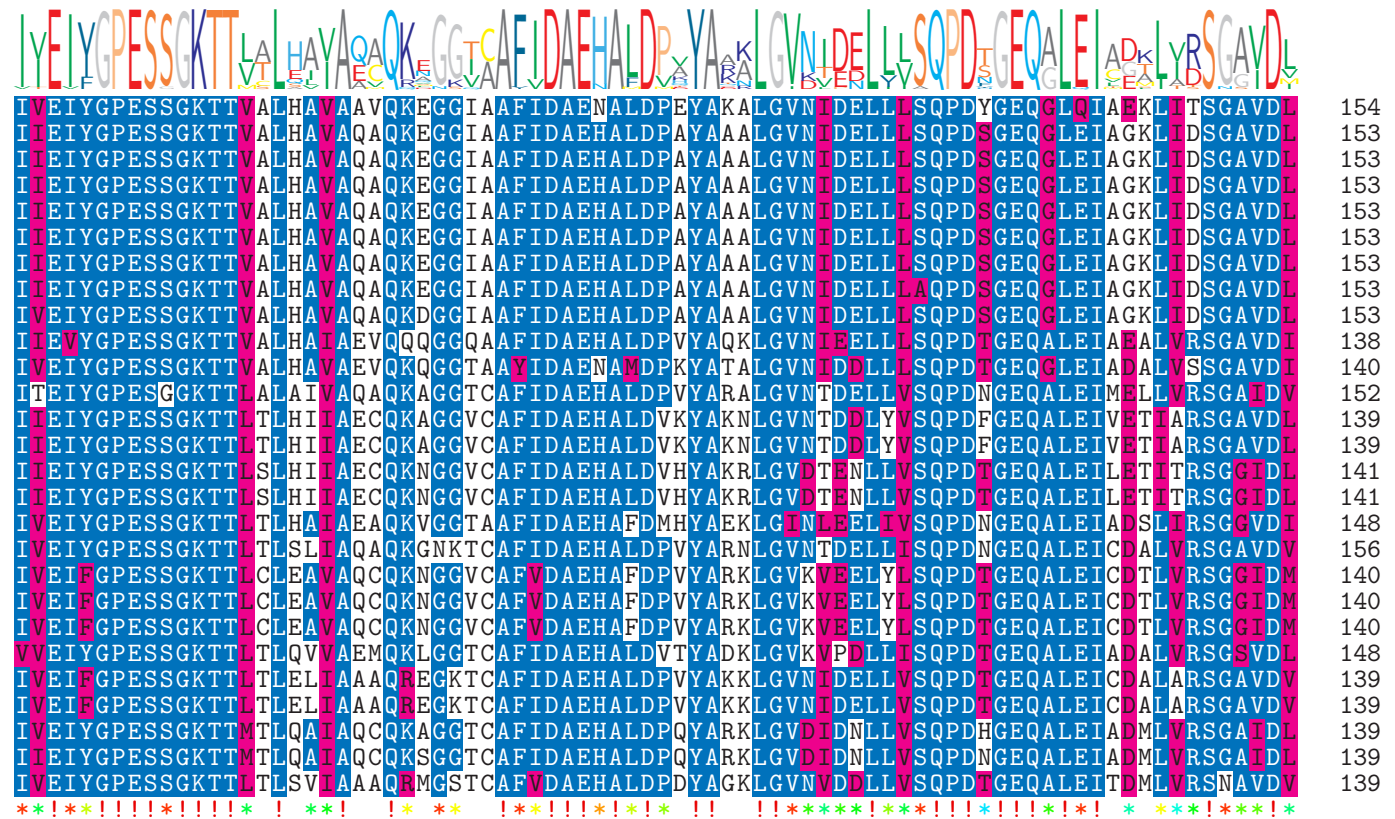
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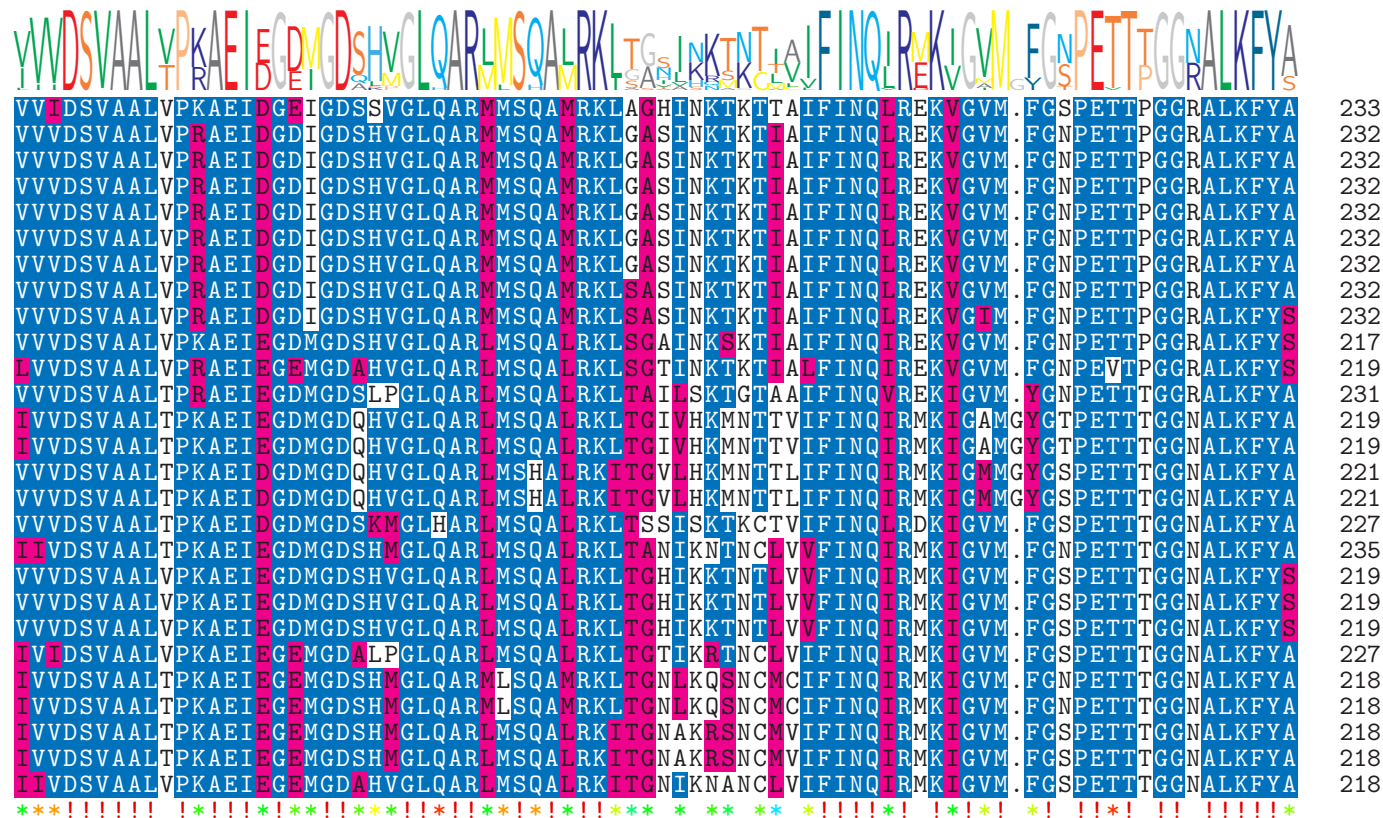
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consensus	*!!**	!	*	*	*	*	*	*****!!*!!*!!*!!*!!*	*	*	*****!!*!!**	*****	***	***	!!*!!**

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consensus

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GEKIGQGSNAKKRYLADHPQLFDEIDRKVRVKFGLLEESEESAAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSNAKKFLADNPFIIDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQAEETQEEL.EFE.E.....	348
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DERIGQGEKTIATYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAPEAAE..A.....	363
DKKLGGGRENSKAFLKENPEIADAITKAIQNSMGIEGMI....SGSEDD..EGE..E.....	343
DKKLGGGRENSKAFLKENPEIADAITKAIQNSMGIEGMI....SGSEDD..EGE..E.....	343
DKKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDE..PLE..EME.....	347
DKKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDE..PLE..EME.....	347
DTKLGGGRDTV KELIKDNPELAELEAKIREKLQAK.....	343
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consensus			

- X non conserved
- X similar
- X ≥ 50% conserved