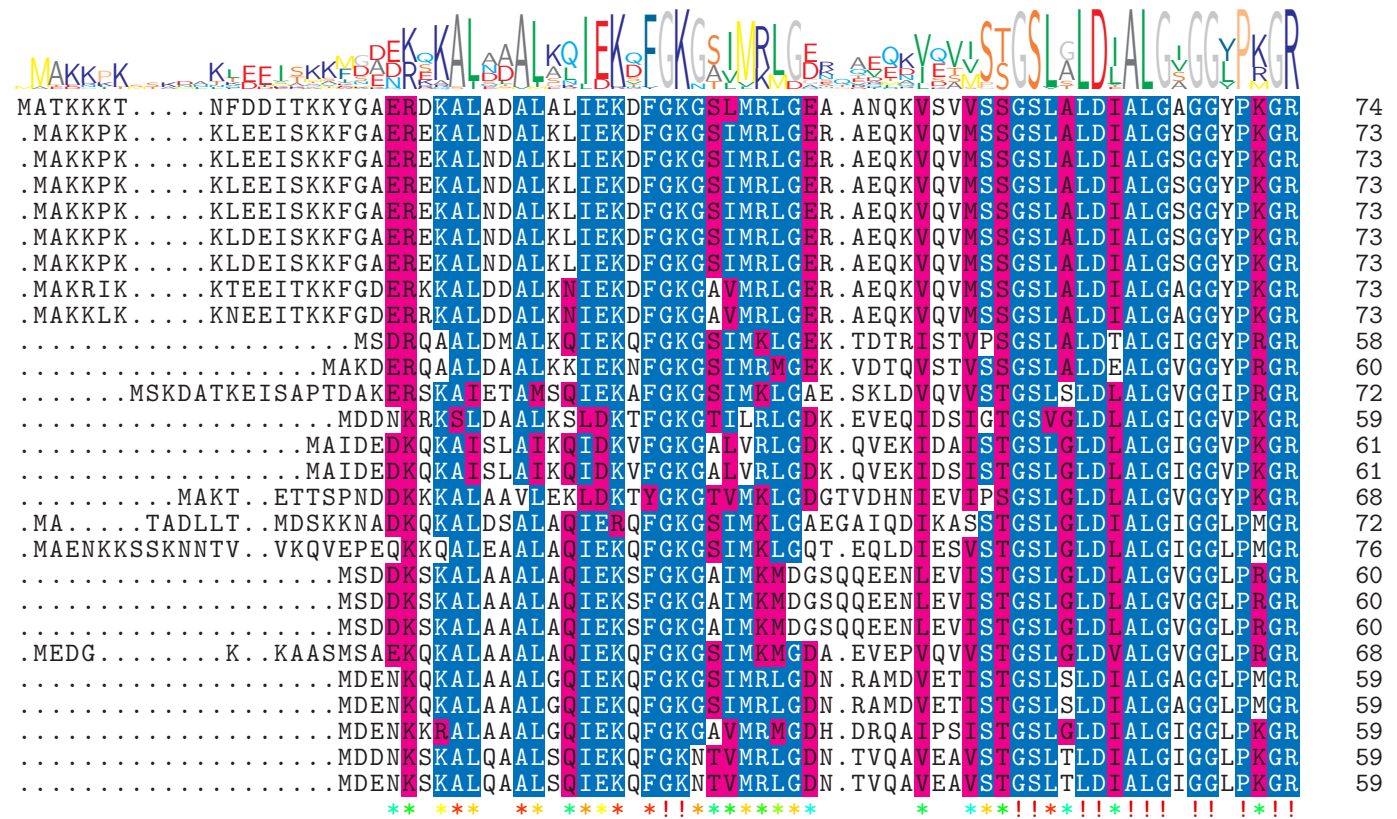


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consensus

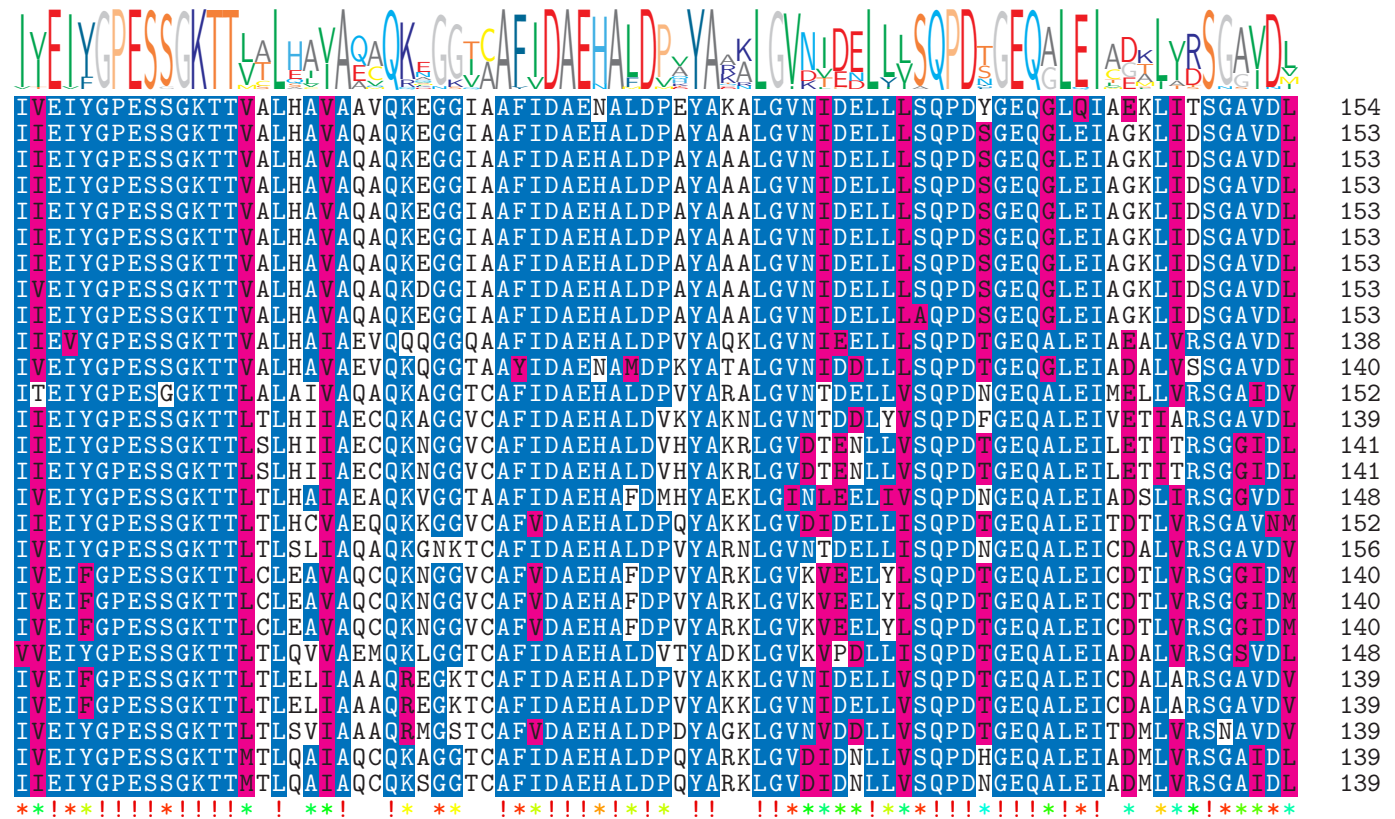
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consensus

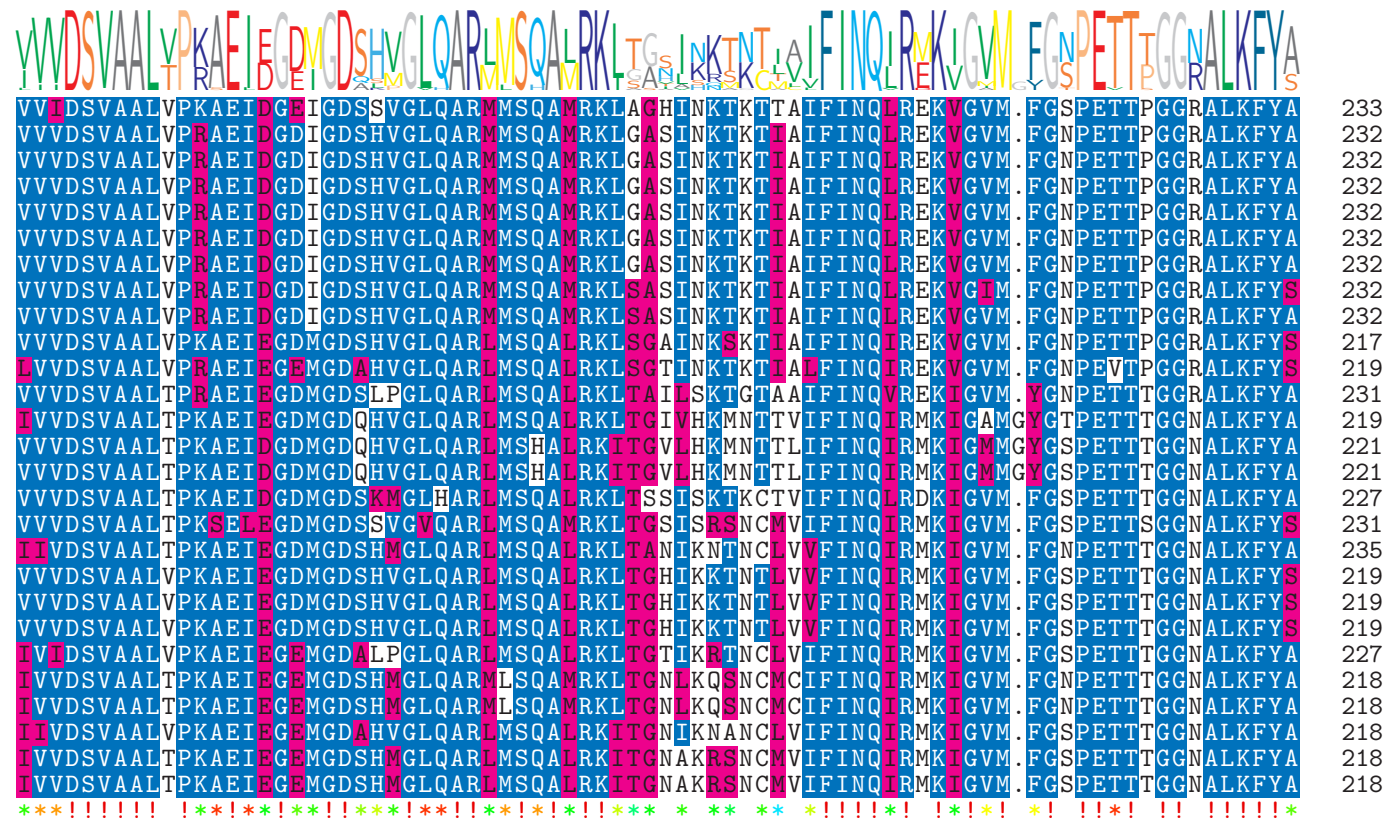
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

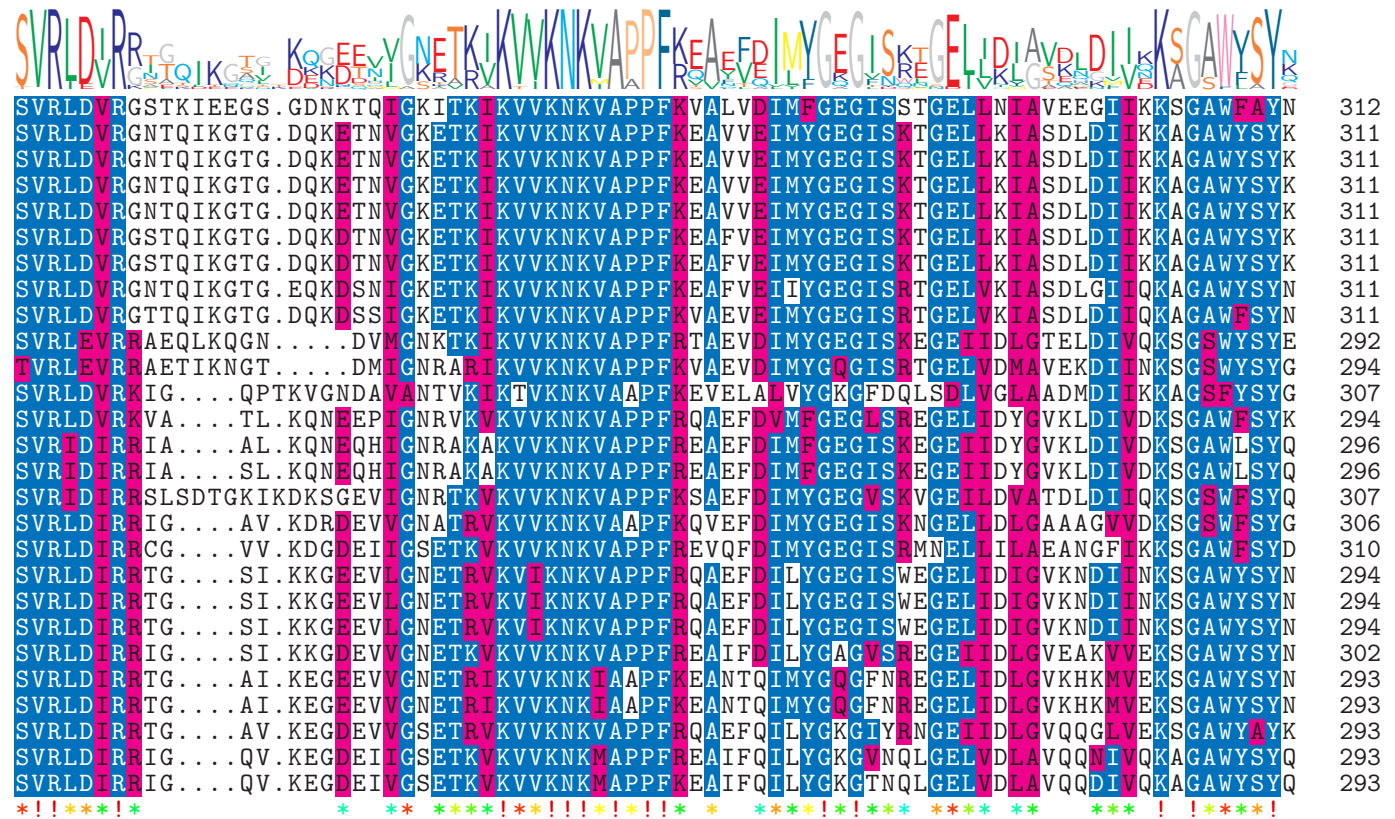
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consensus

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consensus

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Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK.KDE...PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK.KDE...PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGSENAKKYLAHNPEVFAEIDHQVRVQFGLIDGEEAAE...SK.KDE...VAQVDS..VNEEVPLDLGDELEI	380
DEKIGQGSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAE...SK.KDE...AAQVDS..VNEEVPLDLGDELEI	380
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQFGLIEEDNTEEEKSSKE.K...ETDEK..ADKNLVLELDDTIEL	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEEESAMAVAS.E...E...TDDLALDLNGIEI	376
EERLGGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQ.EEL.EFE.E...	348
SERIGGGRENAKNYLADHEDVEDEVRLKVRVQFGLSDVPEEDLPTTEDE..QI.NILPDDST...EE...	355
DERIGGQKEKTIATYIAERPMEQEITDRVMAAIRAGNAGEAPALAPAPAAPEAA...E...A...	363
DKKLGGGRENKSKAFLKENPEIADKLEITKATQNSMGTGEMISG..SEDDEGEE...	343
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DKKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEIMP..LPDEPLEEM.....E...	347
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DERIGGQRENKQFLRDNPKIAFAIEDKIRASHGLDFDMPE..GADA...ADDEDILEA...	359
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GNKIGGQKNNVIRIYFENNQIAEIERNIREQLLTTGTNGA..VQ.I.EDEEEPDLL..LE.....S...	349

logo

		ELEE	
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NTDB id 1074 ABD1 RS10195 WP 000344167.1		349	
consensus			

-  non conserved
-  similar
-  ≥ 50% conserved