

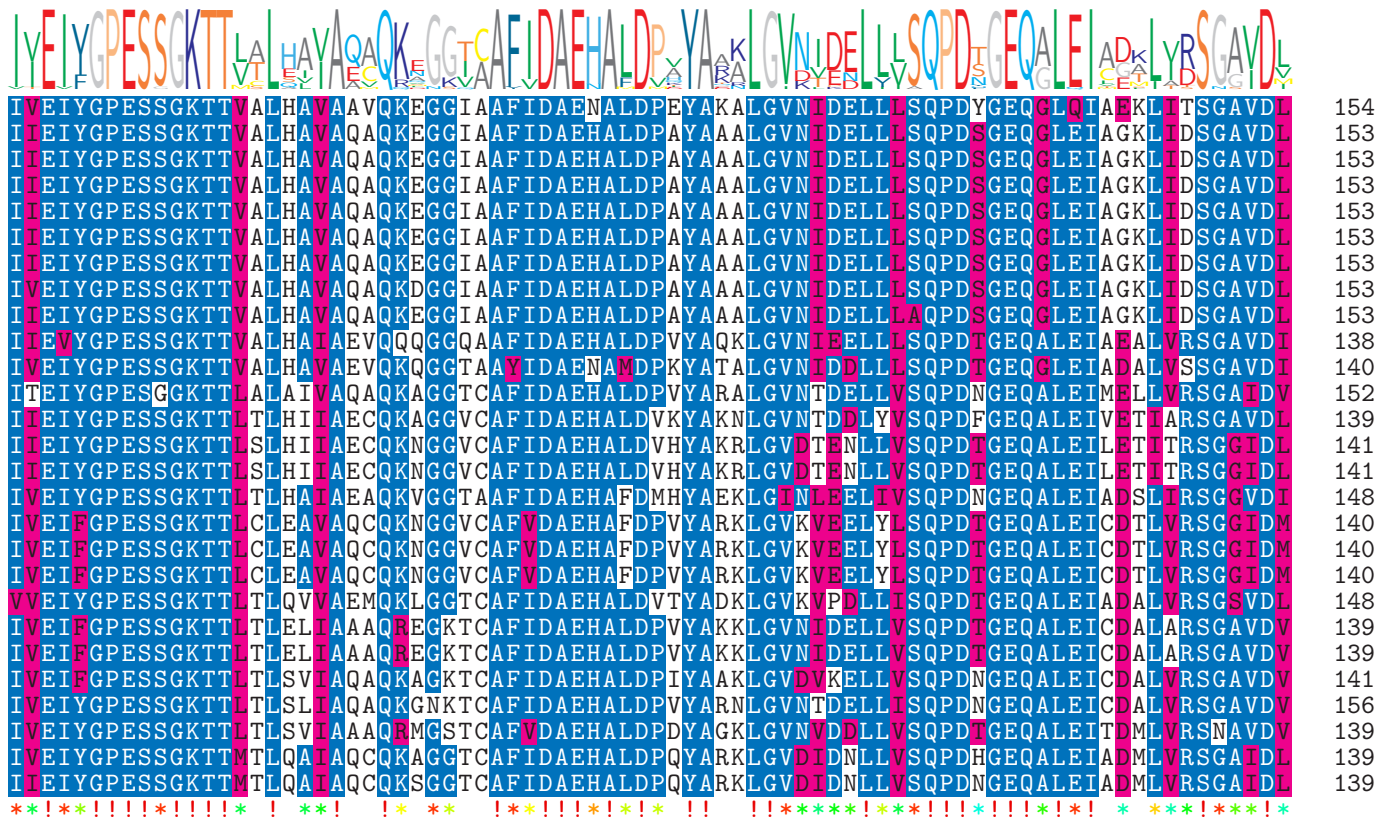
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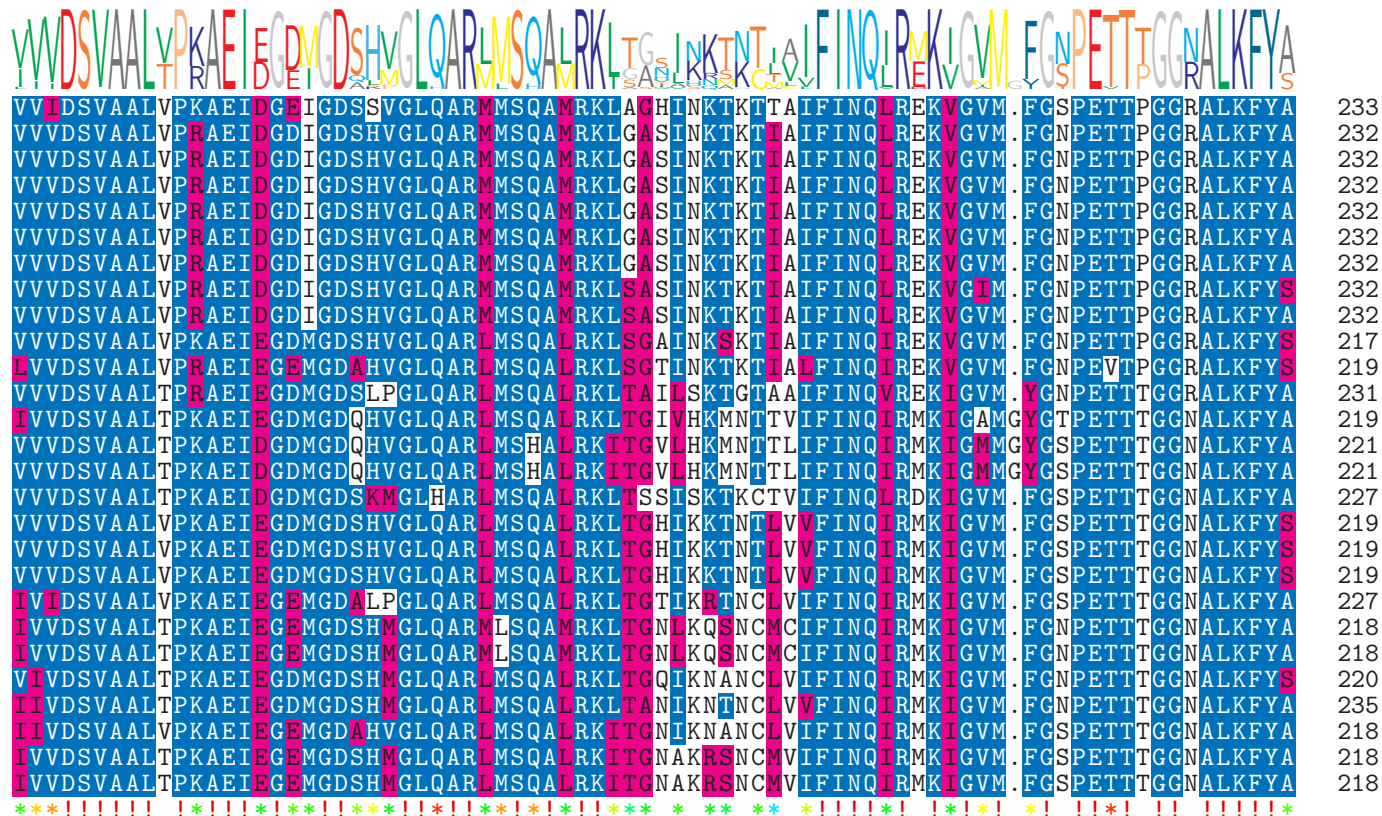
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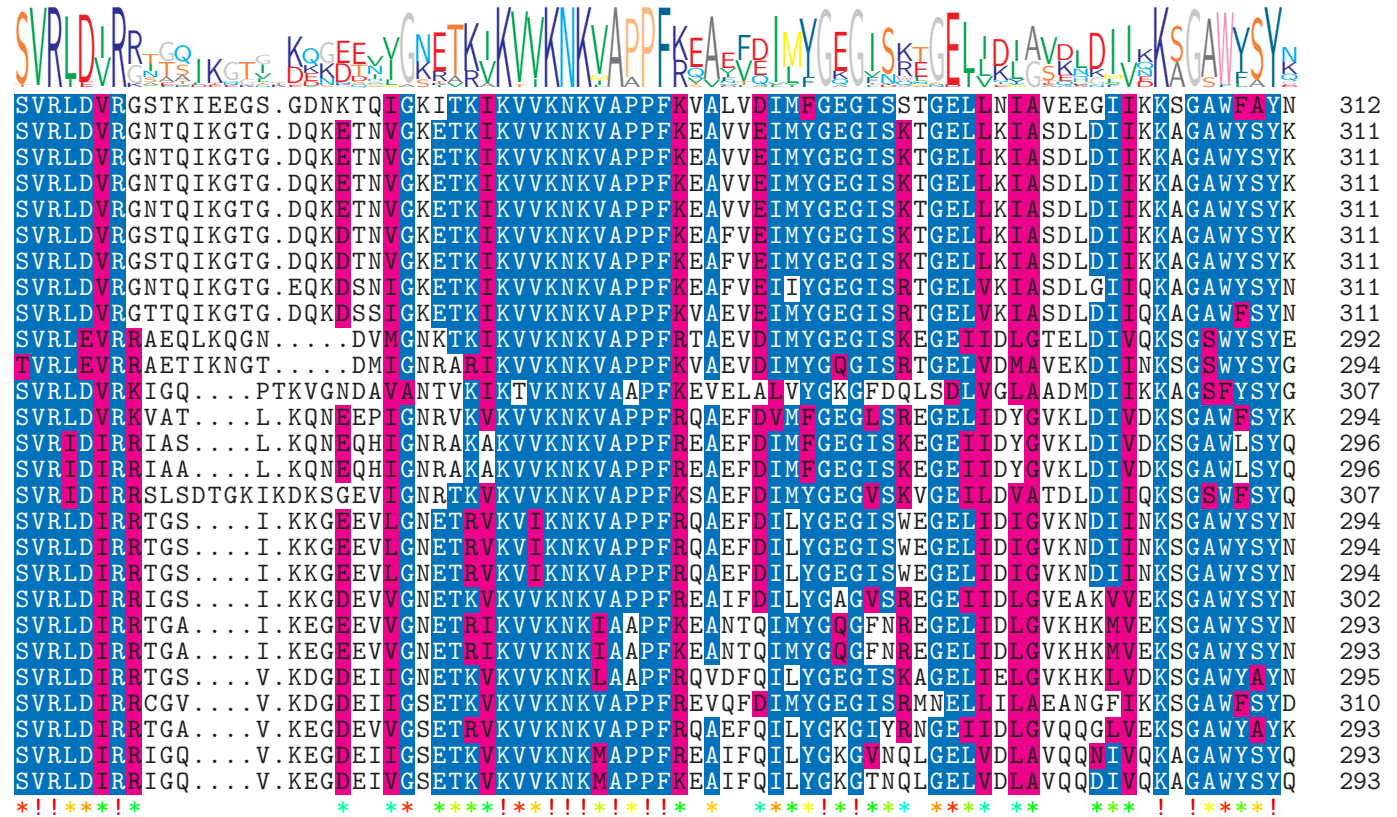
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

Sequence	Score
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DEKIGGQGSENAKKYLAEHPEIFDEIDKQVRSKFG.LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGQGSENAKKYLAADNPEVFAEIDHQVRVQFGLIDGEEAAE....SKKDE....VAQVDS..VNEEVTLDLGDLEIE	381
DEKIGGQGSENAKKYLAADNPEIFDEIDHQVRVQFGLIDGEEAAE....SKKDE....AAQVDS..VNEEVTLDLGDLEIE	381
GEKIGGQGSENAKFLADNPEIFDDIDHKVRVQYGLIEDNTEEEKQSSKEK.....ETDEK..ADKNLVLELDDTTIELE	382
GEKIGGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESSEESAMAVASE.....E.....TDDLALDLNNGIEIE	377
EERLGGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQAETQEEL.EFE.E.....	348
SERIGGQGRENAKNYLADHEDVEDEVRLKVRVQYGLSDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGGQKEKTIAYIAERPEMEQETIDRVMAAIRAGNAGEAPALAPAPAAPEAAE..A.....	363
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logo

		LEE	
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consensus			

- non conserved
- similar
- ≥ 50% conserved