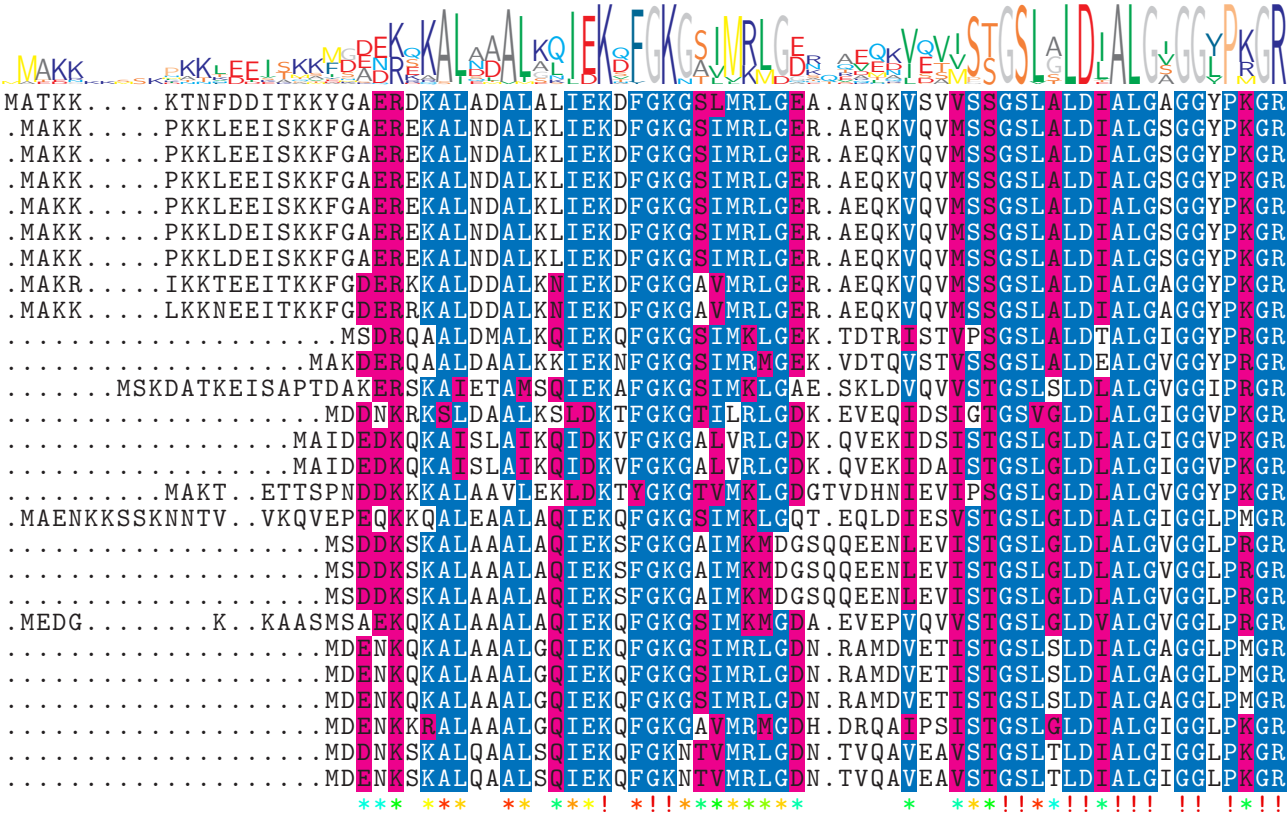


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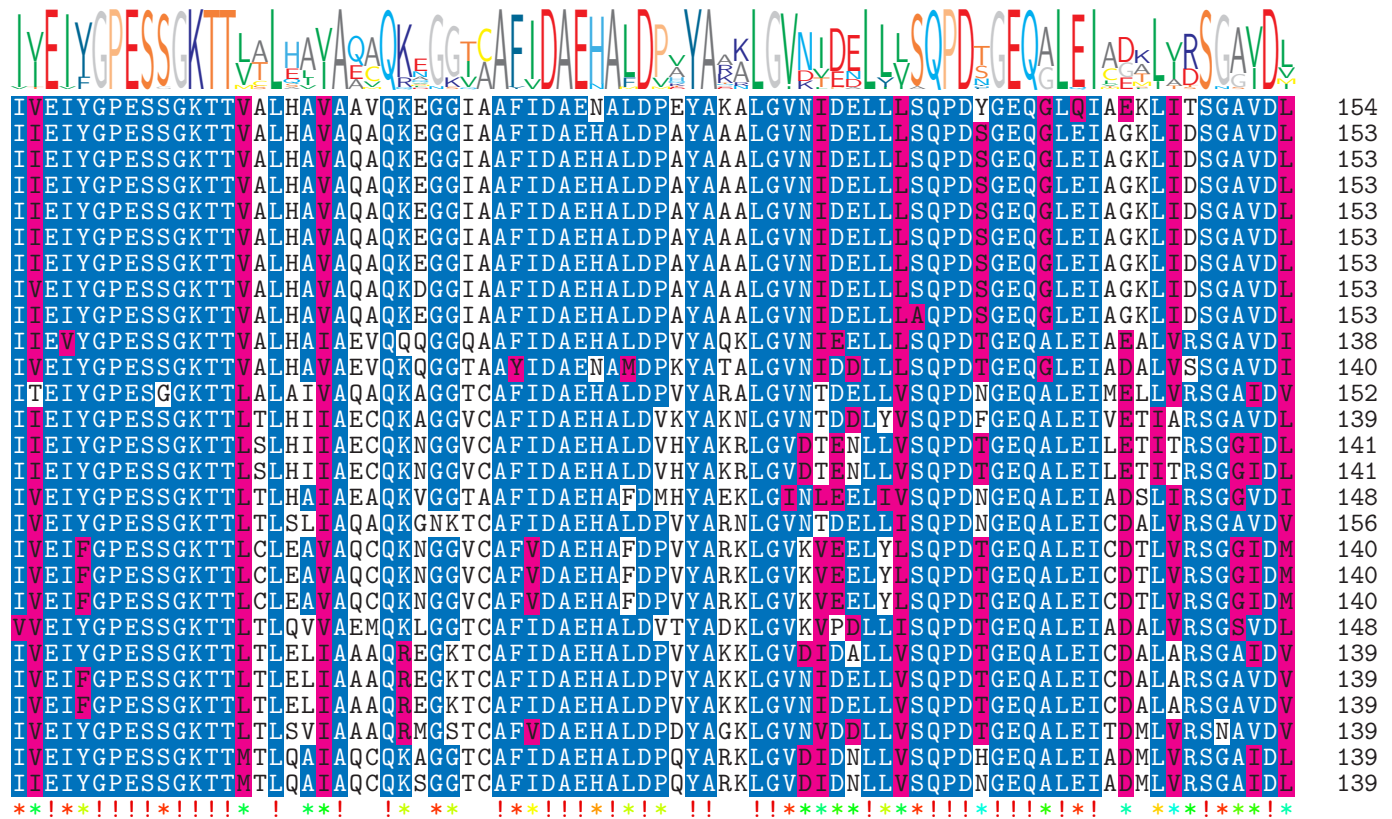


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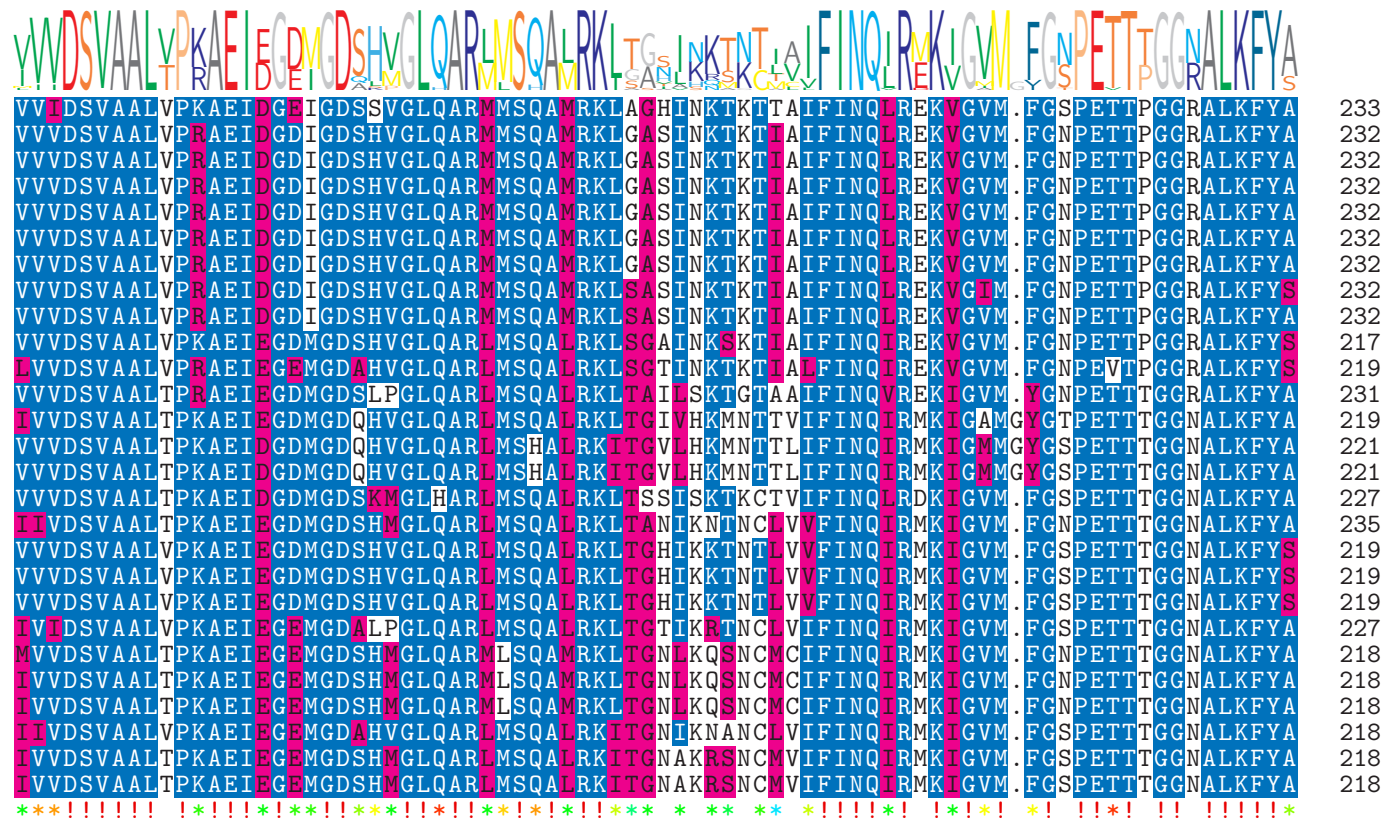
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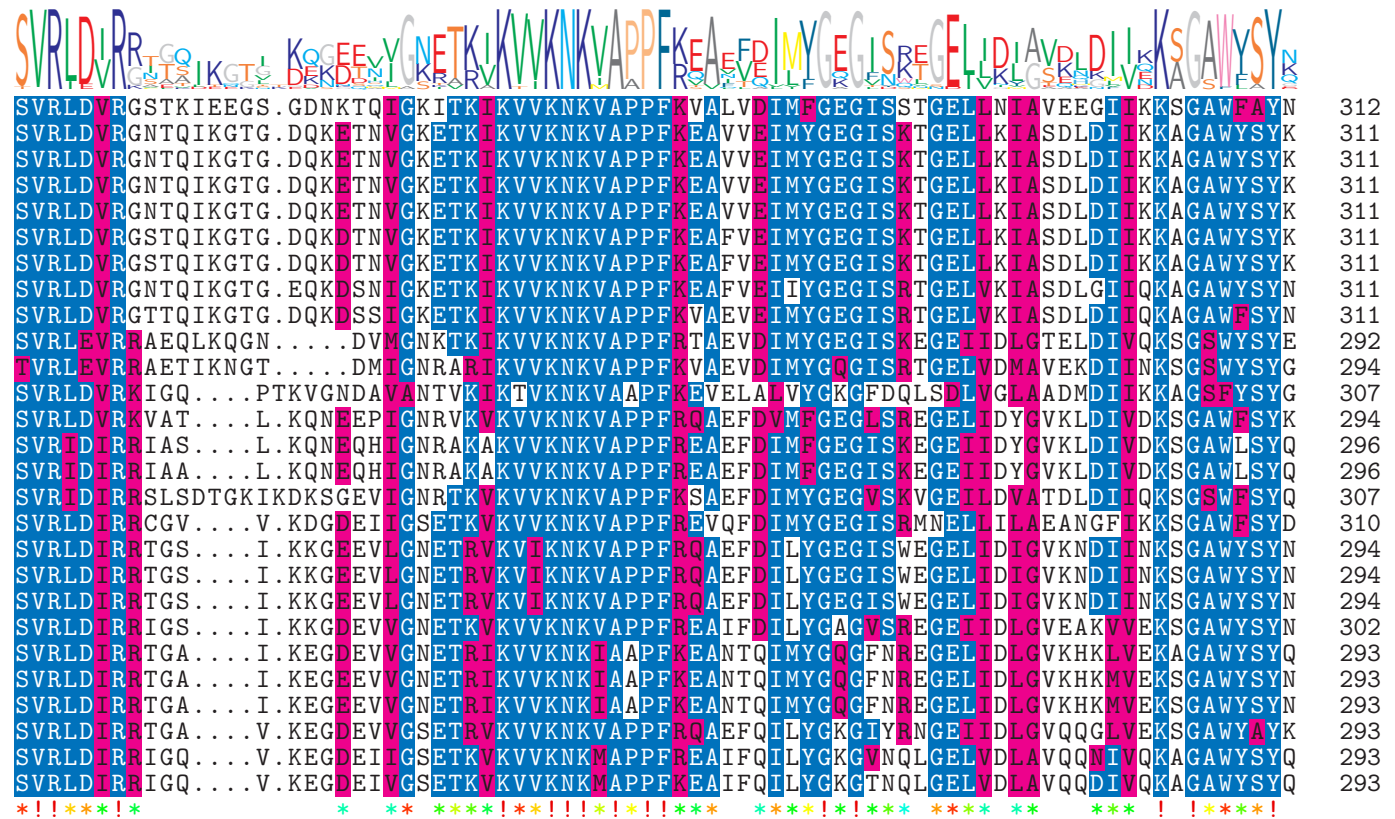
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consensus

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Sequence	Score
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GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE E TDDLALDLNGIEIE	377
EERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNGGVVQQAEETQEEL . EFE . E	348
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logo

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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349	
consensus			

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