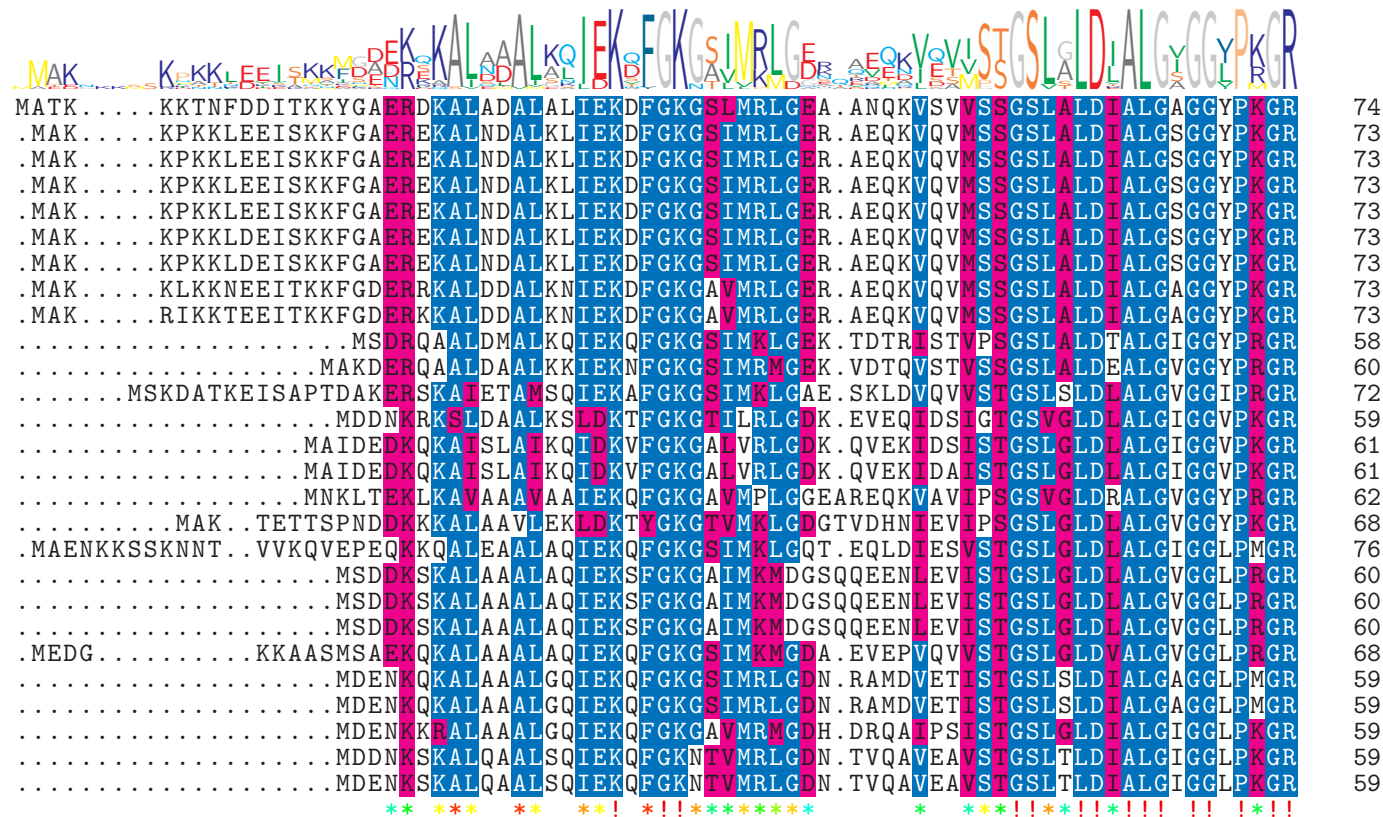


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

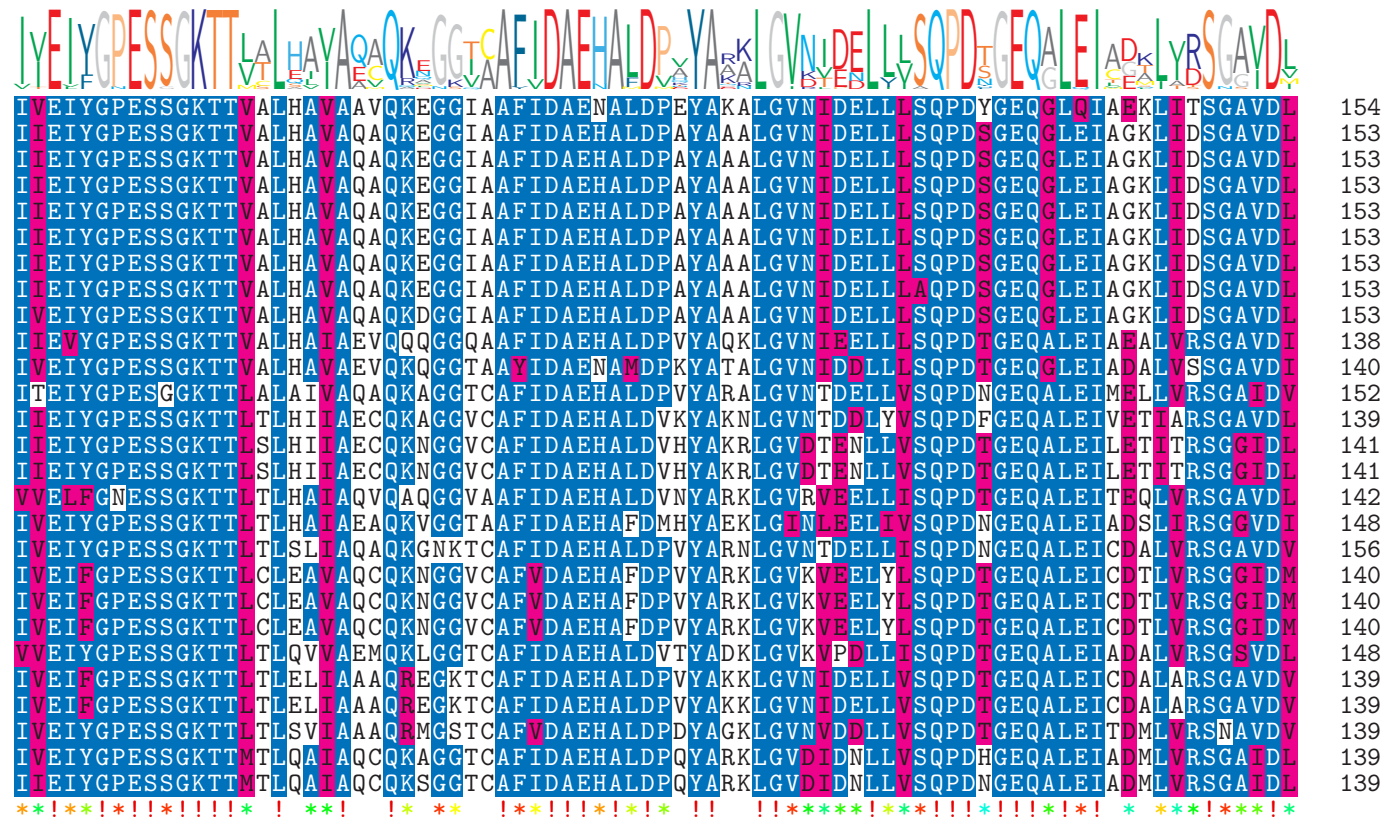
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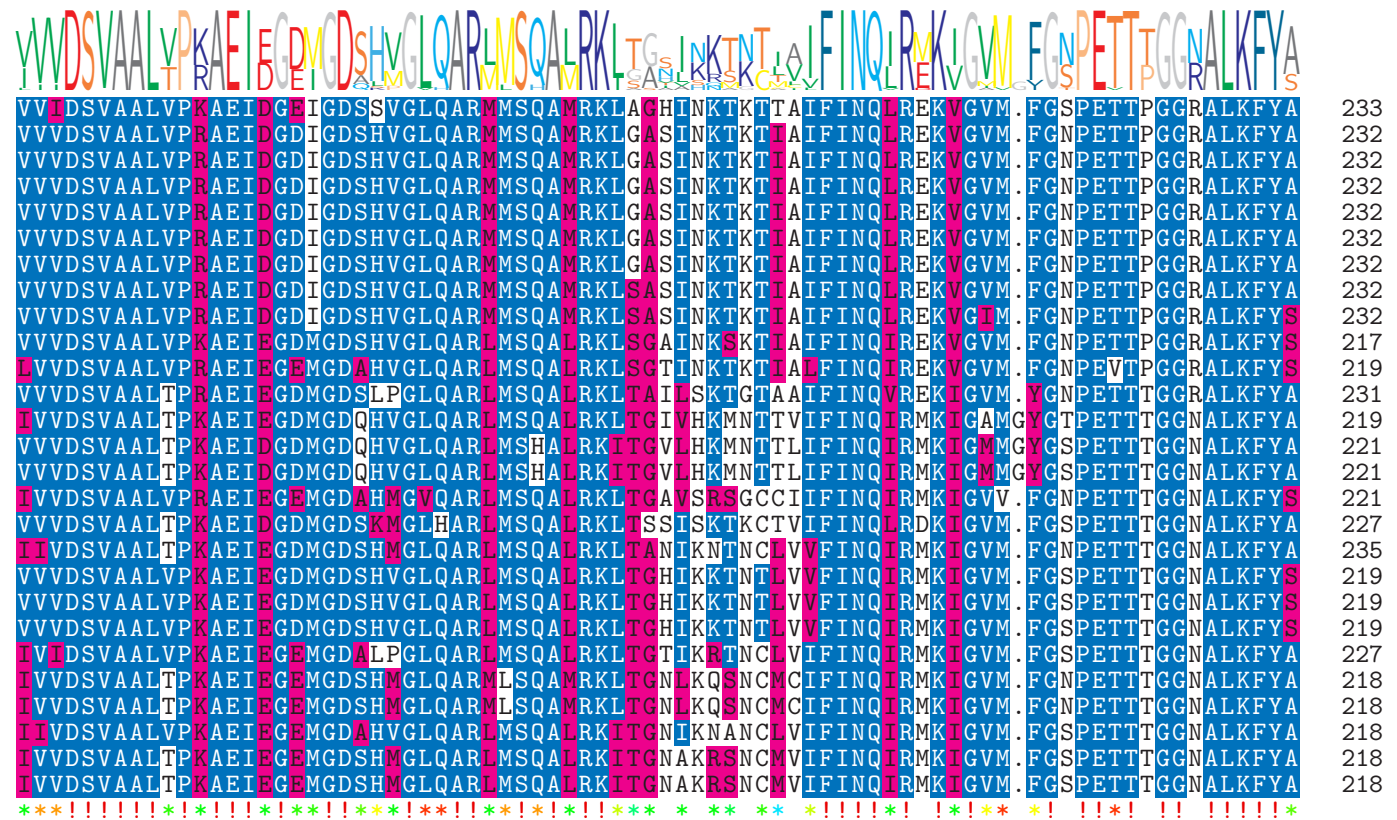
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

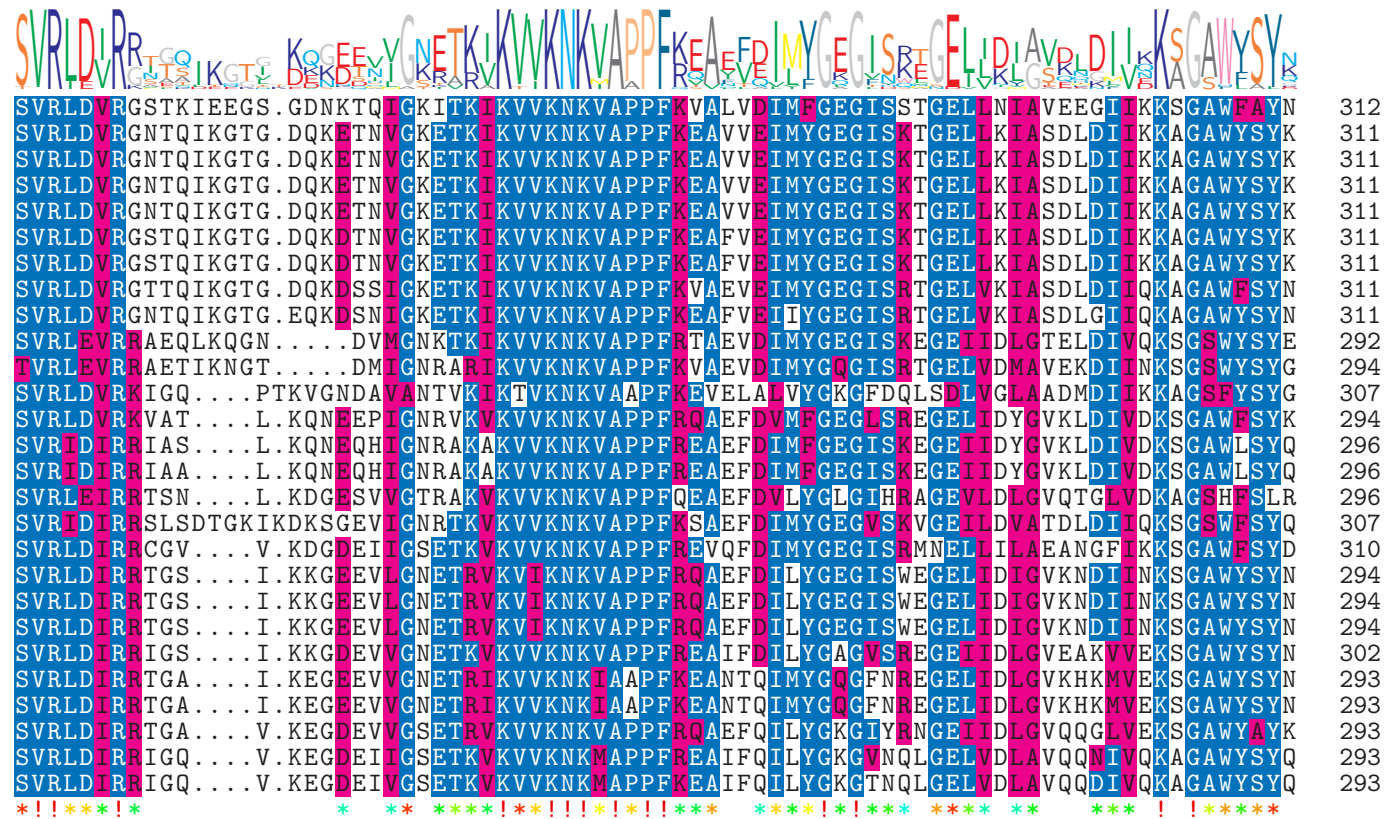
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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consensus

Sequence	Score
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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE . . . PKKEEA . VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE . . . PKKEEA . VNEEVPLDLGDELEIE	385
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DEKIGQGSENAKKYLDADNPEIFDEIDHQRVVFGLIDGEEIAES . . . KKDE . . . AAQVDS . VNEEVTLDLGDLEIE	381
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GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE E TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK ETDEK . . . ADKNLVLELDDTIELE	382
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SERIGGGRENAAKNYLADHEDVEDEVRLKVRVQYGLSDVPEEDLPTTEDE . QI . NILPDDST EE	355
DERIGGQKEKTIATIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAP . EAA EA	363
DKKLGGGRENKSAFLKENPEIADKLEQEIINTLASNPNSVLKATASDSNDES . DVEEF SEEEL	343
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GDKIGGQKANACKYLKENPEIAKTLDDKLREMLLNPNMQLIAETSSAADV . EFGAV PEEF	354
GDKIGGQKANACKYLKENPEIAKTLDDKLREMLLNPNMQLIAETSSAADV . EFGAV PEEF	354
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GNKIGGQKNNVIRYLEENPEIISTEIEAVIREQLLTKASDQTAA . HDETEEEP . DLL ES	349
GNKIGGQKNNVIRYFEENQIAEIERNIREQLLTGTNGAVQ . I . EDEEEP . DLL LES	349

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

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