

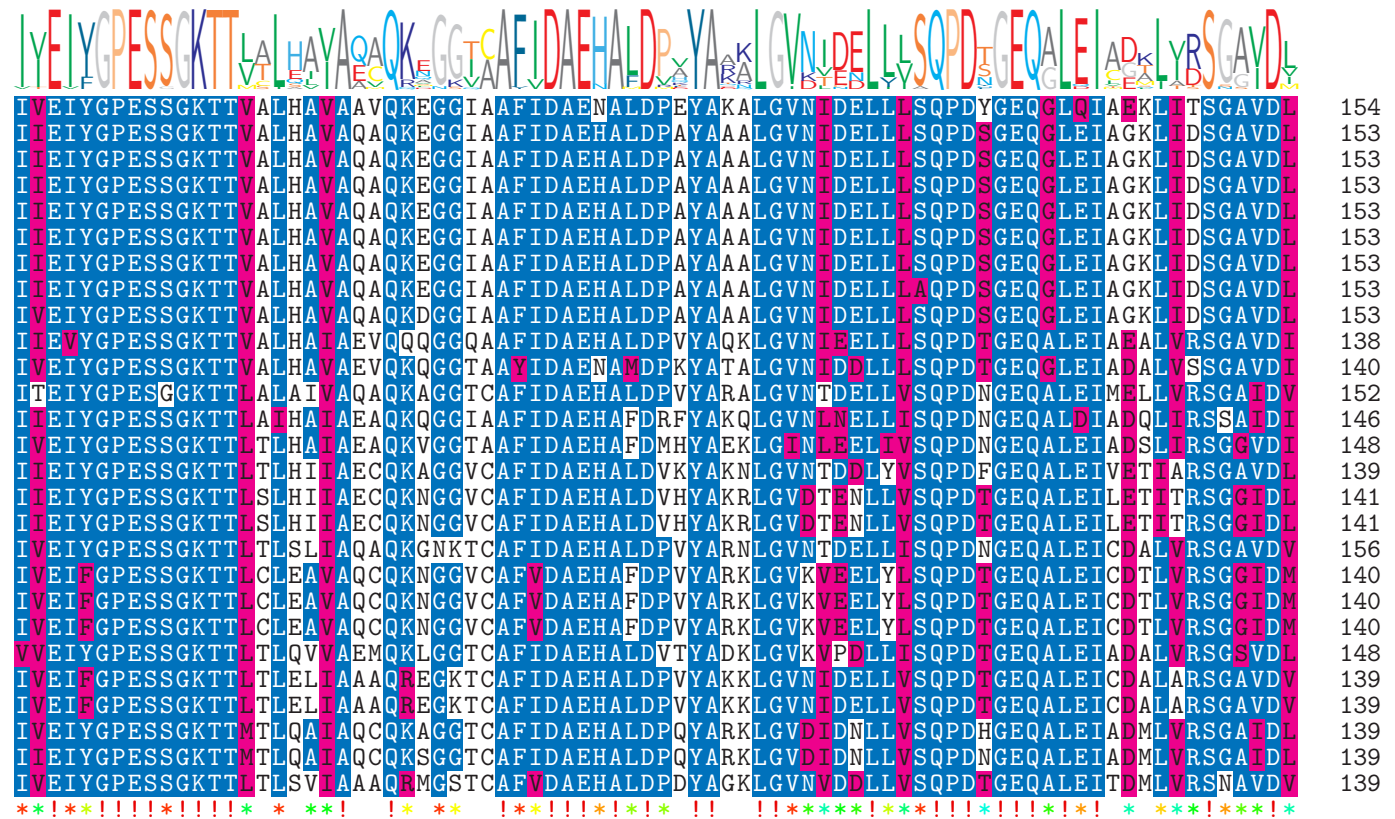
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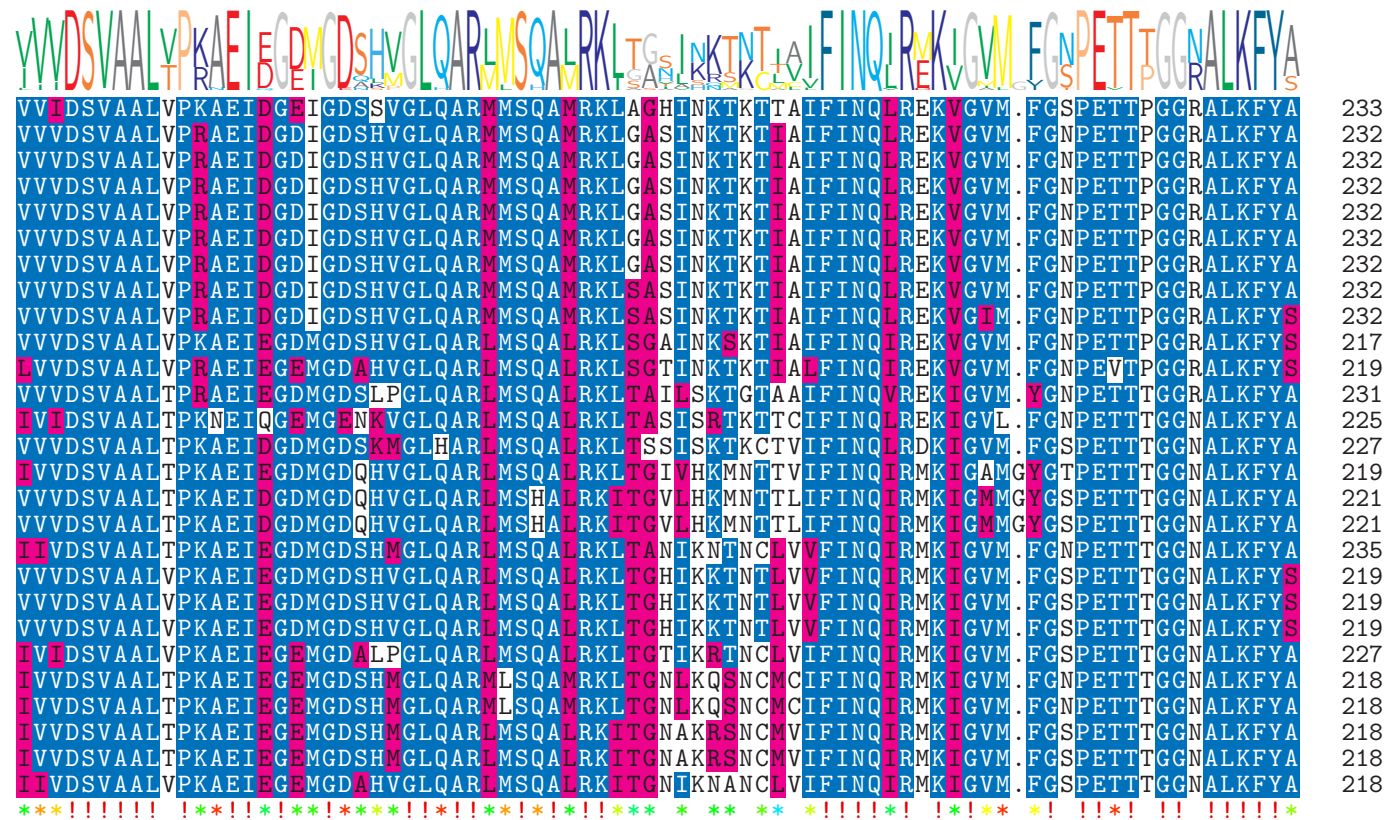
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus



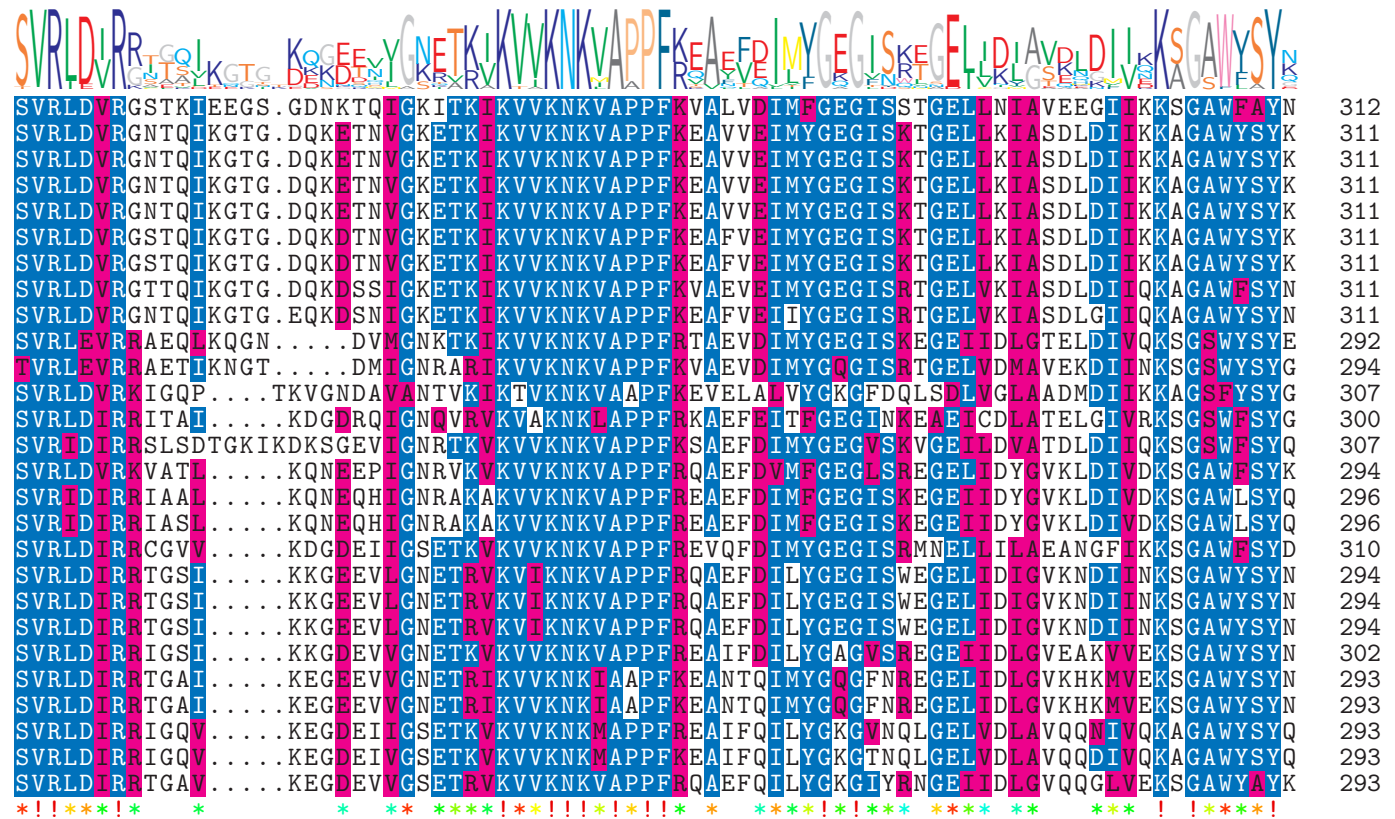
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DEKIGGGSSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQ DTENKKDE PKKEEA . . VNEEVPLDLGD	380
DEKIGGGSSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQ DTENKKDE PKKEEA . . VNEEVPLDLGD	380
DEKIGGGSSENAKKYLAADNPEIFDEIDHQVRVQFLIDGEEIAES KKDE AAQVDS . . VNEEVTLDLGD	376
DEKIGGGSSENAKKYLAADNPEIFDEIDHQVRVQFLIDGEEIAES KKDE VAQVDS . . VNEEVTLDLGD	376
GEKIGGGSSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESA MAVASE E TDDLALDLDN	372
GEKIGGGSSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEK QSSKEK ETDEK . . ADKNLVLEDD	377
EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQQ AEETQEEL . EFE . E	348
SERIGQGRENAKNYLADHEDVEDEVRLKVRVQYGLSDVPEEDLP TTEDE . QI . NILPDDST EE	355
DERIGQGKEKTIAIYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE . A	363
DGKLAQGRESVTKLLKNPELDEIEAKIFAAALRGEDPKDT SDT . PDTDS . DA DFP EEA	357
DTKLGQGRDTPVKELIKDNPELAELEAKIREKLQAK	343
DKKLGQGRNSKAFLENKENPEIADEITKAIQNSMGIIEGMISG SEDDE . . GE . E	343
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEIMP LPDEP . . LE . EME	347
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GERIGQGRDNCREFLENARLARIENKVRHLGVTPMGAV TLA . EEVEE . D	352
GDKIGGQKANACKYLKENPEIAKTLDDKKLREMLLNPNMQL IAETSSAAD . DVEFGAVP EEF	354
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GNKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTAKSDQT AA . HDETEE . EPDLL . . E S	349
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GSKIGGQKANAACKYLENPEIQEIEEQIRAKLLVTSANTK ATPVAEDLA . DADI	347

logo

		ELELEE	
NTDB id 600 KW2 RS01765 WP 011675413.1			387
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consensus			

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