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NTDB id 1142 GTF74 RS11550 WP 000344154.1
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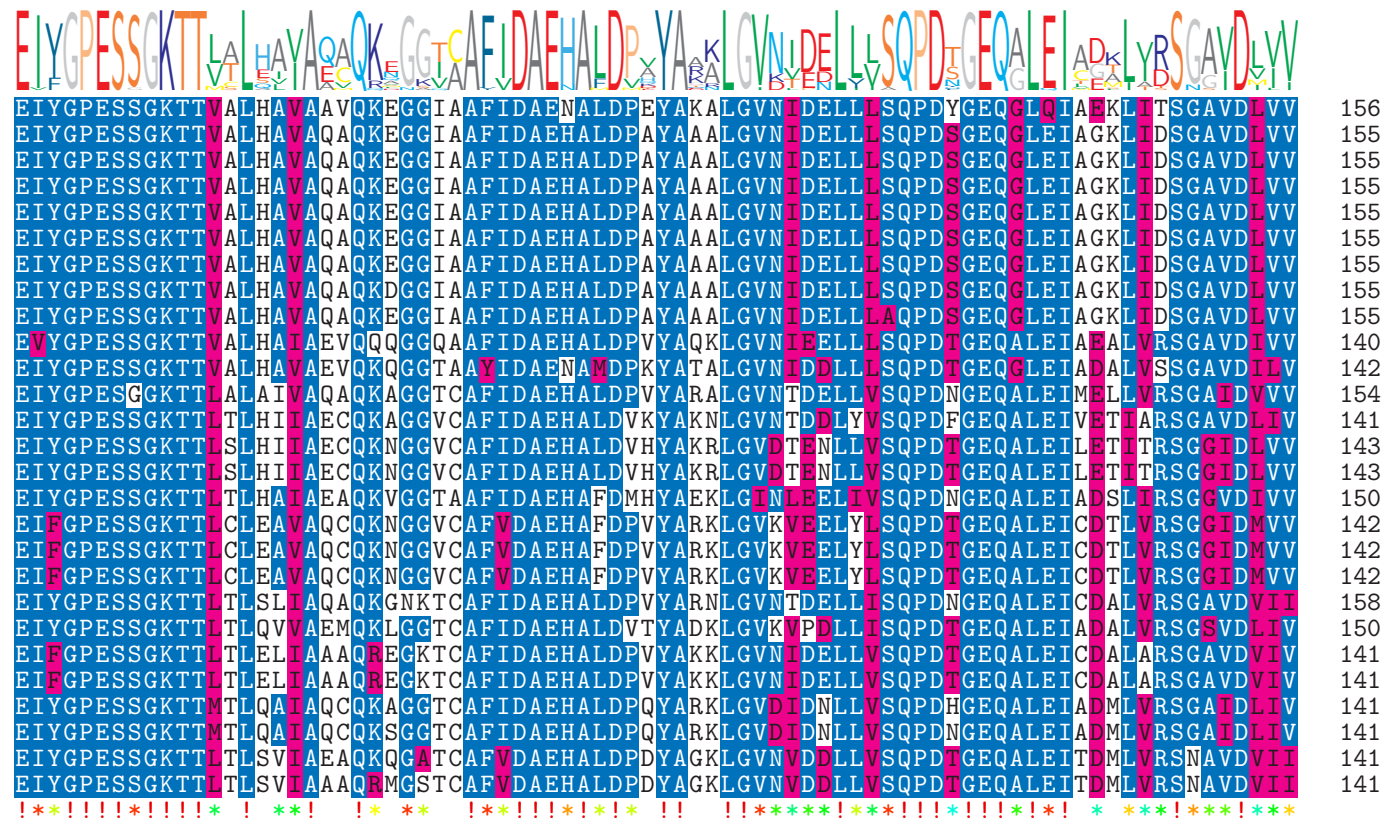
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consensus

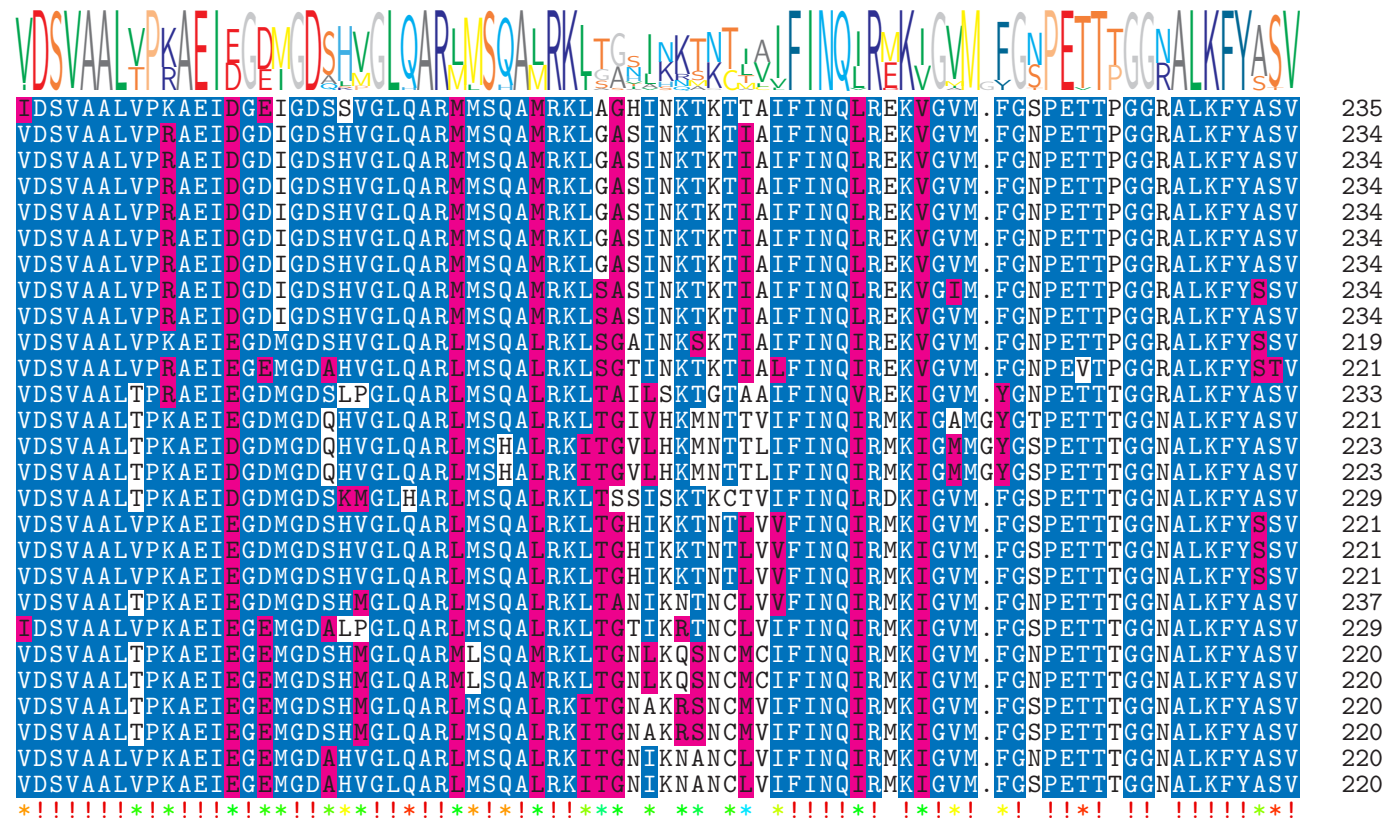
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consensus

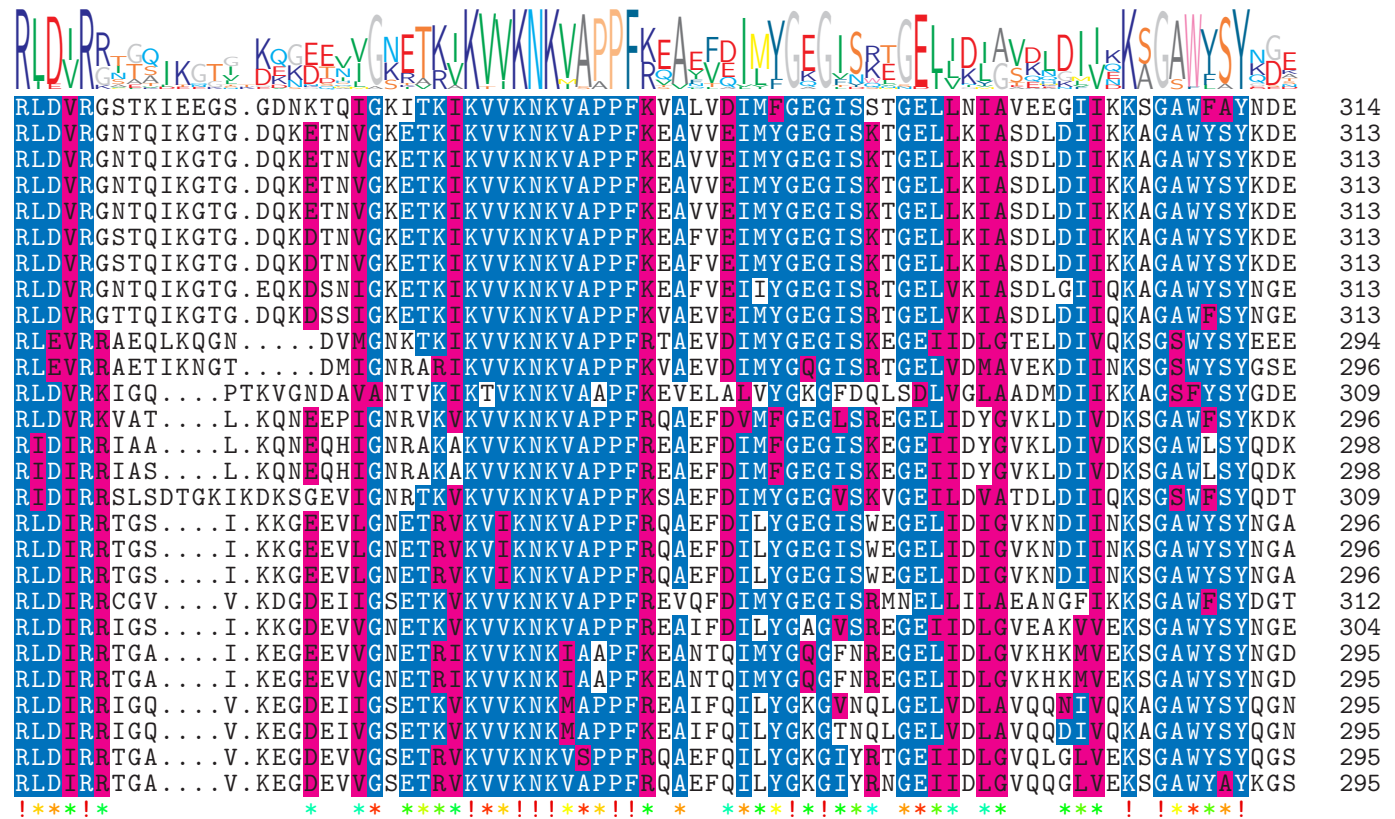
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consensus

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NTDB id 1205 PSJM300 Q5545 AFN77183.1

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Sequence	Score
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KIGGQSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK...KDE...PKKEEA...VNEEVPLDLGDELE	383
KIGGQSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK...KDE...PKKEEA...VNEEVPLDLGDELE	383
KIGGQSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENK...KDE...PKKEEA...VNEEVPLDLGDELE	383
KIGGQSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAES...K...KDE...AAQVDS...VNEEVPLDLGDELE	379
KIGGQSENAKKYLAHNPEIFDEIDHQVRVQFGLIDGEEIAES...K...KDE...VAQVDS...VNEEVPLDLGDELE	379
KIGGQSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEEKSSKE...K...ETDEK...ADKNLVLELDDTIE	380
KIGGQSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVAS...E...E...TDDLALDLNGIE	375
RLGGQRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQ...EEL.EFE.E...	348
RIGGQRENAKKNYLADHEDVEDEVRLKVRVQYGLSDVPEEDLPTTEDE...QI.NILPDDST...EE...	355
RIGGQKEKTIAYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAEA...	363
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KIGGQKANACKYLKENPEIAKTLKKLREMLLNPNMQ...LIAETSSAADVEFGAVP...EEF...	354
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KIGGQKANAACKYLEDNPEIGSVLEKTRDQLLAKSGPV...KADAEVADAEAD...	346
KIGGQKANAACKYLENPEIGQIEEQIRAKLLVTSANT...KATPVAEDLADADI...	347

logo

NTDB id 600 KW2 RS01765 WP 011675413.1		387
NTDB id 269 KZH43 RS08665 WP 001085462.1	IEIEE	388
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved