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NTDB id 1427 RS RS02760 WP 011000510.1
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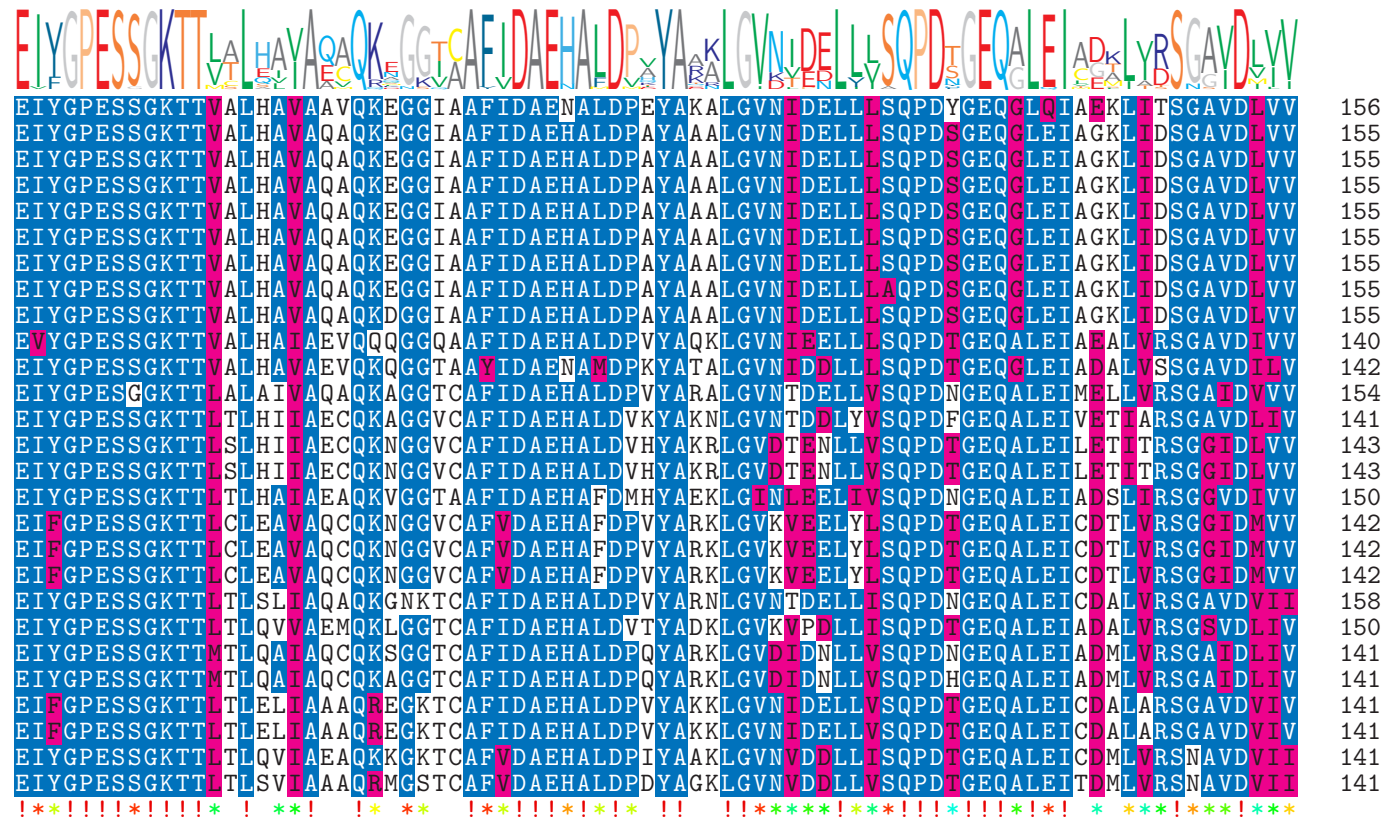
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consensus

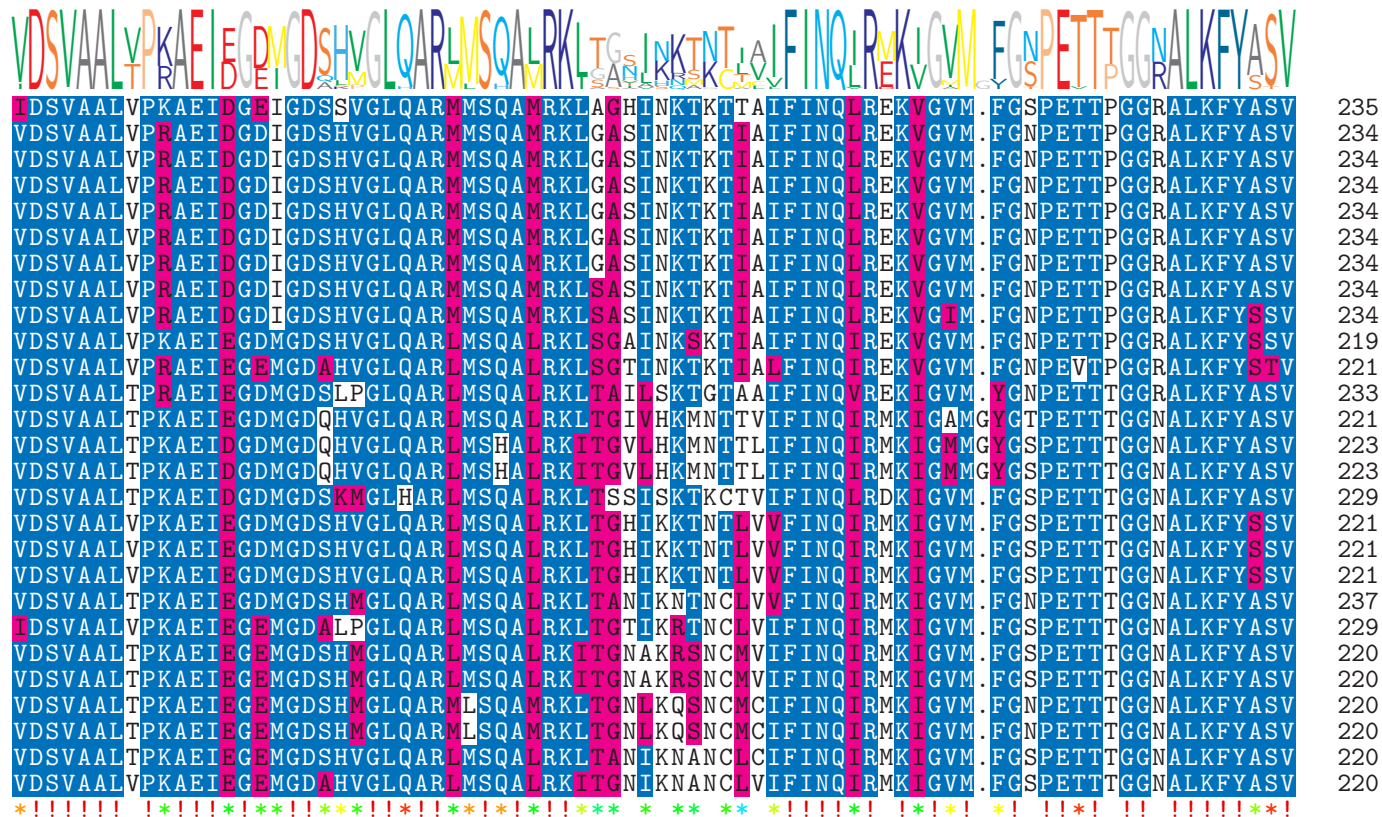
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consensus

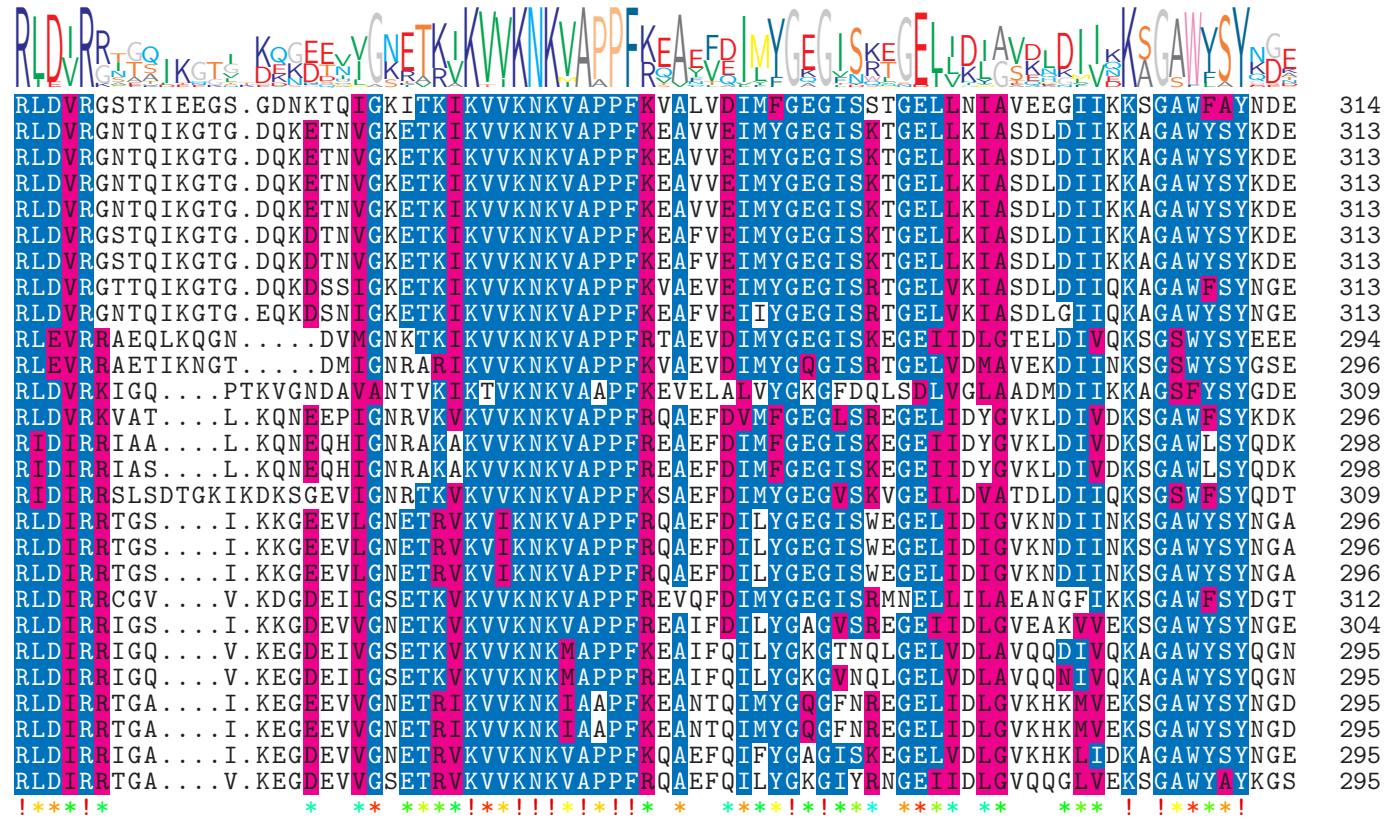
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consensus

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NTDB id 102964 AACH65 RS02930 WP 005348744.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

Sequence	Score
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KIGQGS ENAKKYLAEHPEIFDEIDKQVRSKFG LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEI	386
KIGQGS ENAKKYLAEHPEIFDEIDKQVRSKFG LIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEI	386
KIGQGS ENAKKYLAHDNPEVFAEIDHQVRVQFGLIDGEEAAE....SKKDE....VAQVDS..VNEEVTLDLGDLEIEI	382
KIGQGS ENAKKYLAHDNPEIFDEIDHQVRVQFGLIDGEEIAE....SKKDE....AAQVDS..VNEEVTLDLGDLEIEI	382
KIGQGS ENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEESAMAVASE.....E.....TDDLALDLNGIEIED	378
KIGQGS ENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELED	383
RLGQGR ENAKQFLKENKDIMLMIQEIQIREHYGLDNNGVVQQQAETQEEL.EFE..E.....PDDST.....EE.....	348
RIGQGR ENAKNYLADHEDVEDEVRLKVRMAAIGISDVPEEDLPTTEDE.QI.NIL.PDDST.....EE.....	355
RIGQGEKKTIAIYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAA.....PEAA.....EA.....	363
KLGGRENSKAFILKENPEIAD EITKAIQNSMGT EGMISGSDEDEGE.....E.....	343
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KLGGGRD TVKELIKDNPELAAEELAKIREKLQAK.....	343
KIGQGDKNVRVWLKENPEISDEIDAKIRALNGVEMHI.....TEGTQDET DGERPEE.....	348
KIGQGDKNVRVWLKENPEISDEIDAKIRALNGVEMHI.....TEGTQDET DGERPEE.....	348
KIGQGDKNVRVWLKENPEISDEIDAKIRALNGVEMHI.....TEGTQDET DGERPEE.....	348
KIGQGKNNAVKWLKENPEIADKLEQEIRNTLASNPNSVLKATASDS.NDES...DVEEF....SEEL.....	372
RIGQGRDNCREFLRENAELAREIENKVRHGLVTPMGAVTLAAEV.EED.....	352
KIGQGKNNVIRYFEENNQIAEIEIERNIREQLLTGTGNGAVQI.ED....E...EEDPLL....LES.....	349
KIGQGKNNVIRYLEENPQISTEIEAVIREQLLTKASDQTAAHDET....E...EEDPLL....ES.....	349
KIGQGKANACKY LKENPEIAKTLDKKLREMLLNPNENMQLIAETSSAADDVEFGAVPEEF.....	354
KIGQGKANACKY LKENPEIAKTLDKKLREMLLNPNENMQLIAETSSAADDVEFGAVPEEF.....	354
KIGQGKANVMKLF AENKVMAGEVEEARLRELLLSGAVPVDDKAAPV.EADEFDAESEQEF....E.....	354
KIGQGKANAAKY LLENPEIQGEIEQQIRAKLLV TSA..NTKATPV.AEDLADADI.....	347

logo

		EE	
NTDB id 600 KW2 RS01765 WP 011675413.1	..	387	
NTDB id 269 KZH43 RS08665 WP 001085462.1	EE	388	
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NTDB id 1205 PSJM300 05545 AFN77183.1	..	347	
consensus			

-  non conserved
-  similar
-  ≥ 50% conserved