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NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSIGCDRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLDISGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS�AEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
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consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKQWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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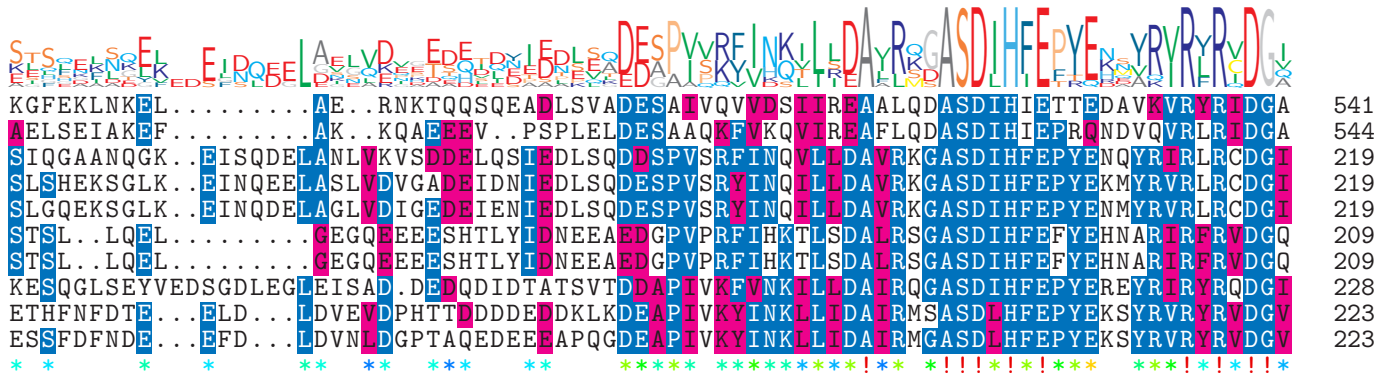
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NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVGLLRILVQNVVTVVEQAHHYY.NESQACKEVLPMLFSDCVISPKSLAAL IARVFSYSILDIRH	65
NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQCTGQLLVLEKLLDKTKATELHKLAAAEKISLLQYIVKNKILSAEQIALTAQNFVGPMLDINC	74
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFFRRLVBEKHVSAATMTALDAAKRAKQDTWAYLIEEVHISPSLSLAETISAEFAEPYFDLDV	74
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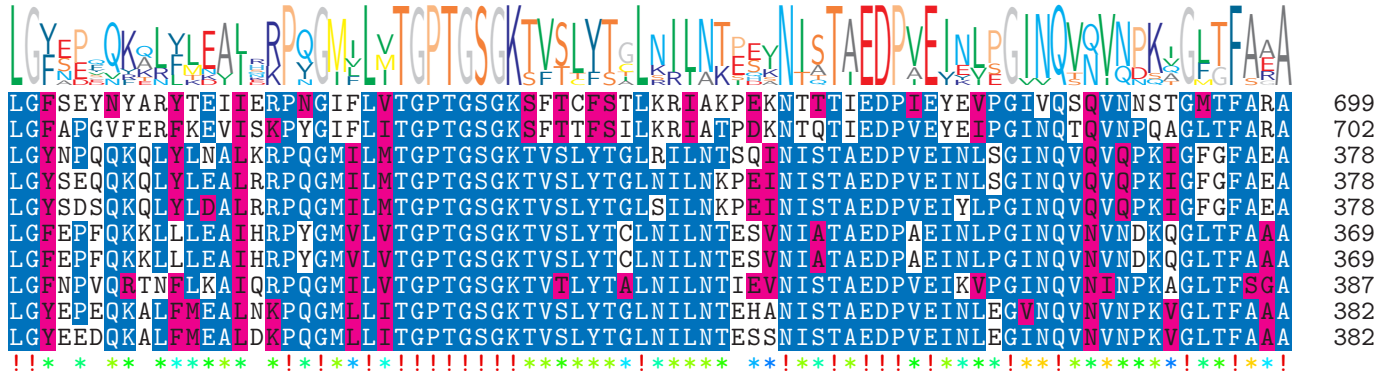
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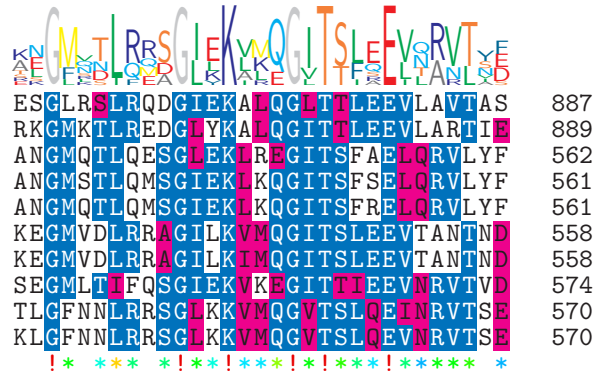
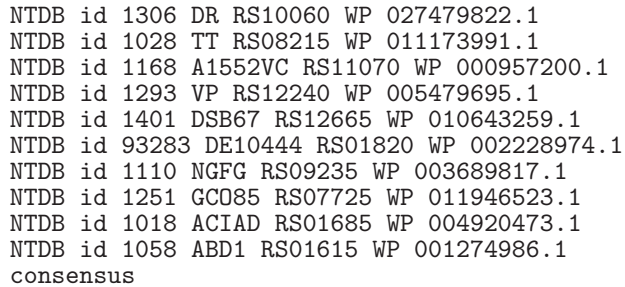
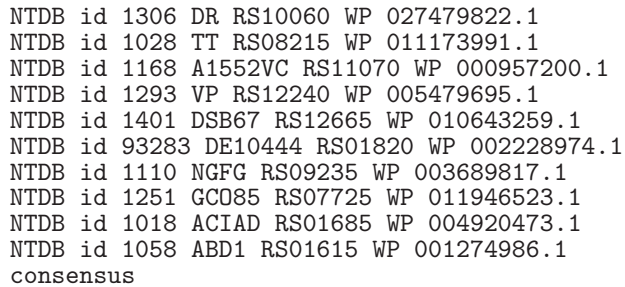
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consensus

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- X non conserved
- X similar
- X $\geq 50\%$ conserved