

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1016564 ABXU53 RS23760 WP 003116249.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	REYRAELKELRAIPFALBEEAGYRVAHYBPLSIFSYEEEDSGDYEPYQATREAYMAAATHYPELGLLEYMPSGK	158
NTDB id 1306 DR RS10060 WP 027479822.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	157
NTDB id 1110 NGFG RS09235 WP 003689817.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1016564 ABXU53 RS23760 WP 003116249.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1016564 ABXU53 RS23760 WP 003116249.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1028	TT RS08215 WP 011173991.1		0
NTDB id 1306	DR RS10060 WP 027479822.1		0
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NTDB id 1251	GC085 RS07725 WP 011946523.1		0
NTDB id 1016564	ABXU53 RS23760 WP 003116249.1		0
NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
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consensus			0

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NTDB id 1028	TT RS08215 WP 011173991.1	EALKEA	397
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NTDB id 1168	A1552VC RS11070 WP 000957200.1		66
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NTDB id 1016564	ABXU53 RS23760 WP 003116249.1		71
NTDB id 1058	ABD1 RS01615 WP 001274986.1		74
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consensus			74

logo

NTDB id 1028	TT RS08215 WP 011173991.1	DPPDPGAPL	477
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NTDB id 1293	VP RS12240 WP 005479695.1	YHEYASLCQQ	141
NTDB id 1401	DSB67 RS12665 WP 010643259.1	YHEYASLCQT	141
NTDB id 1251	GC085 RS07725 WP 011946523.1	IDVGTIPVNL	149
NTDB id 1016564	ABXU53 RS23760 WP 003116249.1	LERDSL	146
NTDB id 1058	ABD1 RS01615 WP 001274986.1	YDTSQIPKD	149
NTDB id 1018	ACIAD RS01685 WP 004920473.1	FDPALFLKDK	149
consensus			149

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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1306 DR RS10060 WP 027479822.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

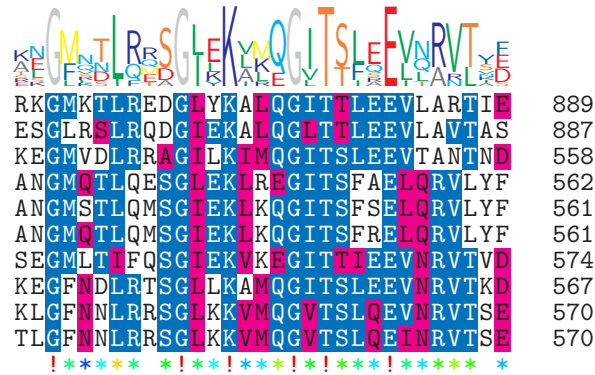
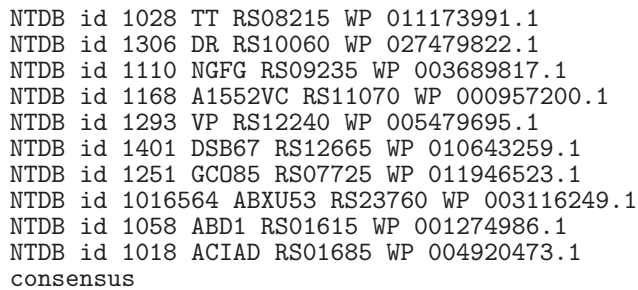
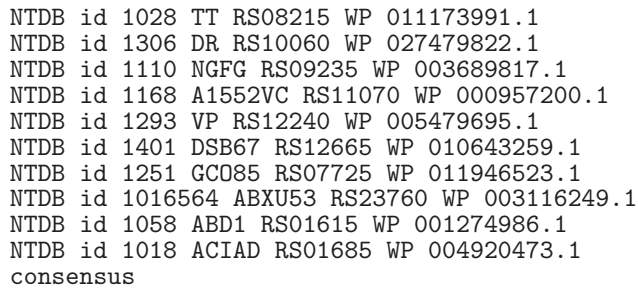
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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