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NTDB id 1028 TT RS08215 WP 011173991.1	MSXLSIGDRRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
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 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1007783 ABL844 RS25855 WP 126024679.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

NTDB id 1028 TT RS08215 WP 011173991.1	REYBAEELKELRAIPFALBEEAGYRVAHYBPLSIFSYEEYEDSGYEPYQTLREAYMAAHHYPELGLLEYMPSGK	158
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consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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consensus		

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[illegible]

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.....MSVGLLRILVQNQVVTVEQAHHYNE.SQAQKEVLPMLFSDGVISPKSLAALILARVFSYSIDLRLH	65																																																																																																			
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MSALH.....TSPKFTGFFRRLVVEEKHVS AATMTALDAAKRAKQDTVAYLIEEVHLSPSLLAETISAEFAEPYFDLDV	74																																																																																																			
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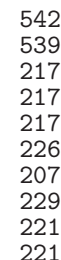
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consensus

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[illegible]

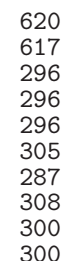
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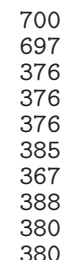
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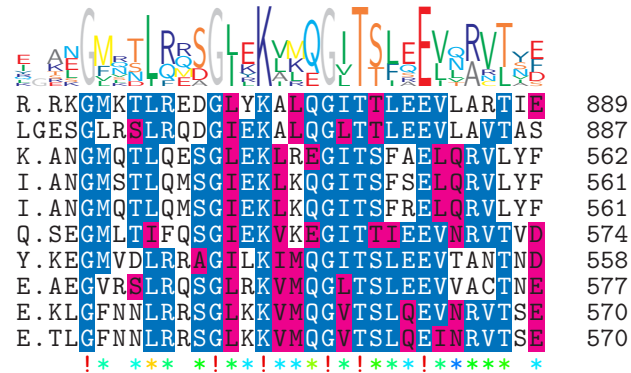
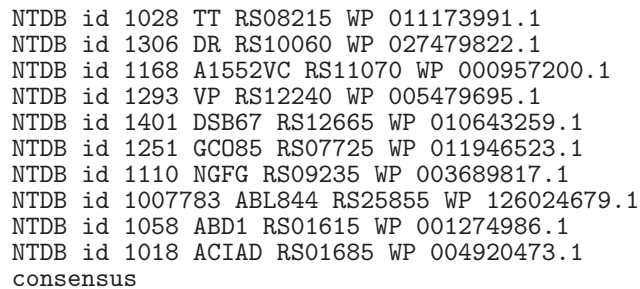
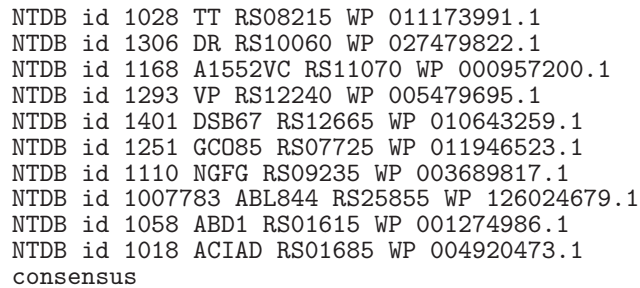
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consensus
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- X non conserved
- X similar
- X > 50% conserved