

logo

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NTDB id 1028 TT RS08215 WP 011173991.1		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 1110 NGFG RS09235 WP 003689817.1			0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1			0
NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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		SLTPAEKAKELRAIPFALDEEAGVVRVAFLNPLDTSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPP..P	
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NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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		SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	
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consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLED TLVQ..SGKLRPEALAA.....VATQL	388
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NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLS TLRLQKGLLTFSQEESLIEQVKASGISMPEALLS..SGFFTSSSELAEH.....LSSIF	57
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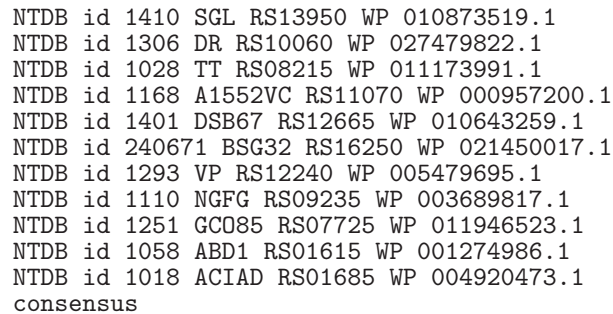
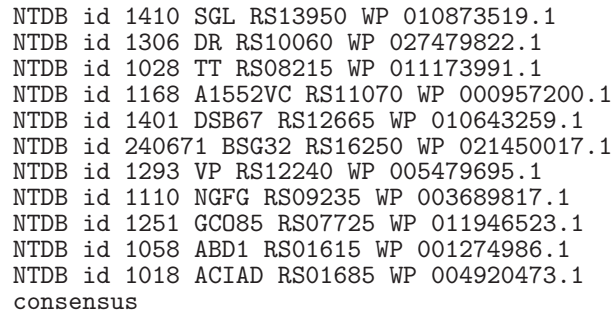
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NTDB id 1251 GC085 RS07725 WP 011946523.1	GVPMLDINCI DVG.....TIPVNLVNEKLIK RHAMVPLFSR...GTNLYLATDDPSKQASLKEIQFHTG.....LNTHAI	132
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLIDLGVFDPA.....LFLKDKLIDEKLIQKYRIMPLVHR...GHVLYVATSNPTNIEAMDAIRFN SK.....LKVEPI	132
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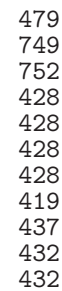
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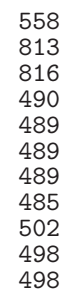

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consensus
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

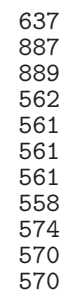
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved