

logo

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NTDB id 1028 TT RS08215 WP 011173991.1		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 1293 VP RS12240 WP 005479695.1			0
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NTDB id 1251 GC085 RS07725 WP 011946523.1			0
NTDB id 402404 FOB21 RS08290 WP 005265509.1			0
NTDB id 1058 ABD1 RS01615 WP 001274986.1			0
NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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		SLTPAEKAKELRAIPFALDEEAGVVRVAFLNPLDTSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPP..P	
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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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		SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFGHPGEEVL	
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consensus			

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NTDB id 1401	DSB67 RS12665 WP 010643259.1		0
NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GCO85 RS07725 WP 011946523.1		0
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consensus			

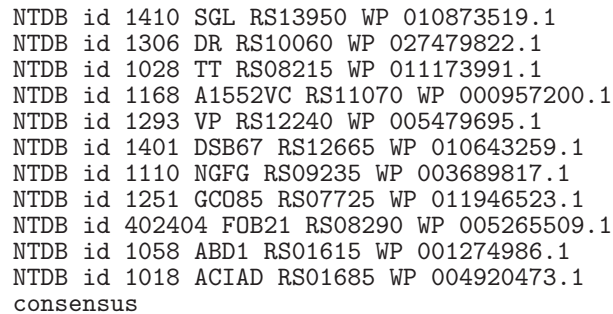
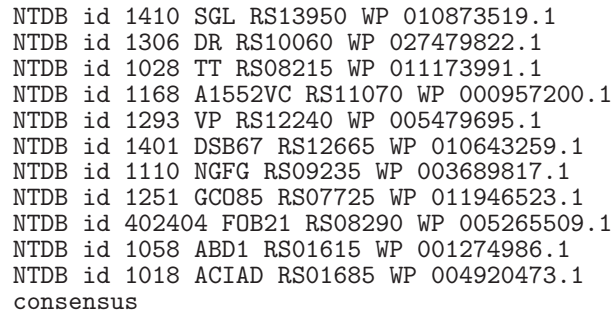
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NTDB id 1293	VP RS12240 WP 005479695.1	MHSNLSTILRQKGLTFSQEESLIEQVKASGISMPEALLS..SGFFTSSLEAHE.....LSSIF	57
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NTDB id 1251	GC085 RS07725 WP 011946523.1	MALATEEYRLQGLLQGLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVK..NKILSAEQIALT.....AAQNF	65
NTDB id 402404	FOB21 RS08290 WP 005265509.1	MTALQGSPRFTGFIRRLVEEGVISAEDMRSALAHAKQEKIDIVAELIN..QQQLSPTVIAET.....ISVEF	65
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	MSAFTTPPKFTSGFIRRLVEEGYVNAQNMQQALEKAKFKFKQDITVPYLID..NFSISPLTIAEI.....ISLEF	65
consensus		* * * * *	

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NTDB id 1251	GC085 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKLIKRRHAMVPLFSRG...TNLYLATDPSKQASLKEIQFHTG.....LNTHAI	132
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	GEPLDLIGVEDPA....LFLKDKDIEKLQIKYRIMPLVHRG...HVLVYVATSNPTNIEAMDIAIRFNSK.....LKVEPI	132
consensus		* * * * *	

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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

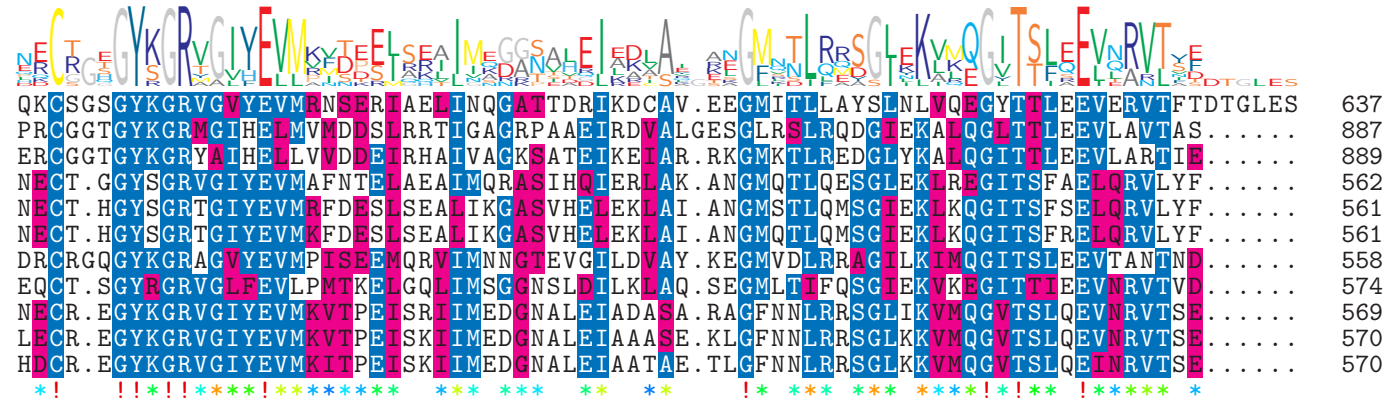
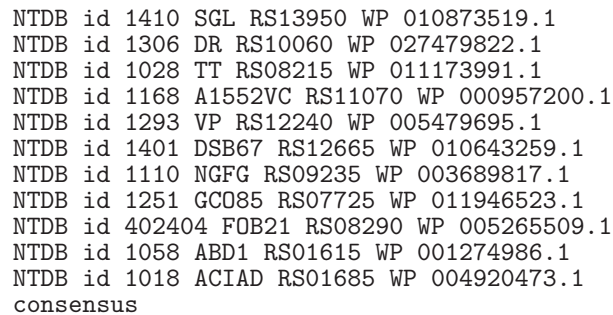
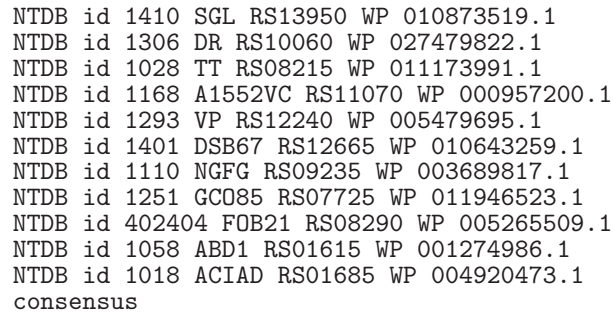
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consensus

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved