

logo

	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKK	
NTDB id 1410 SGL RS13950 WP 010873519.1	0
NTDB id 153855 MOTHE RS07525 WP 011393063.1	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKK	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

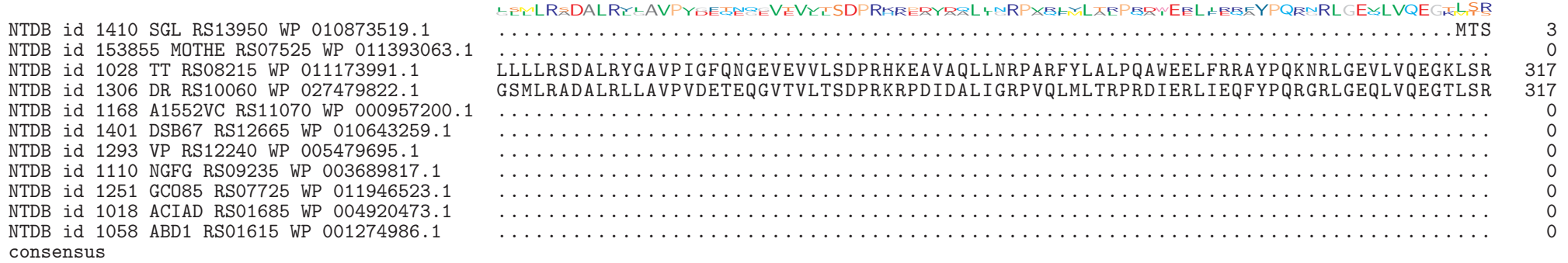
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	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTL SLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

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	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVDPDSAA	
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consensus		

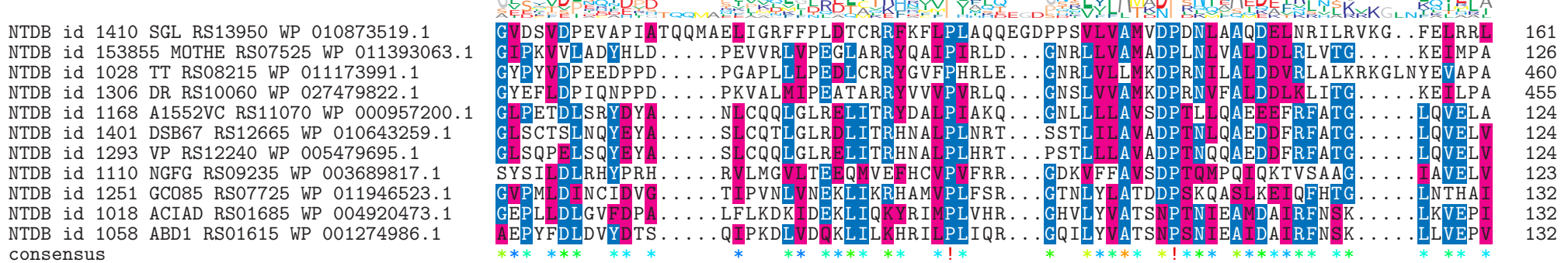
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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	574
NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
consensus		

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