

logo

	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKK	
NTDB id 1410 SGL RS13950 WP 010873519.1	0
NTDB id 153876 MOTH RS08015 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
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

logo

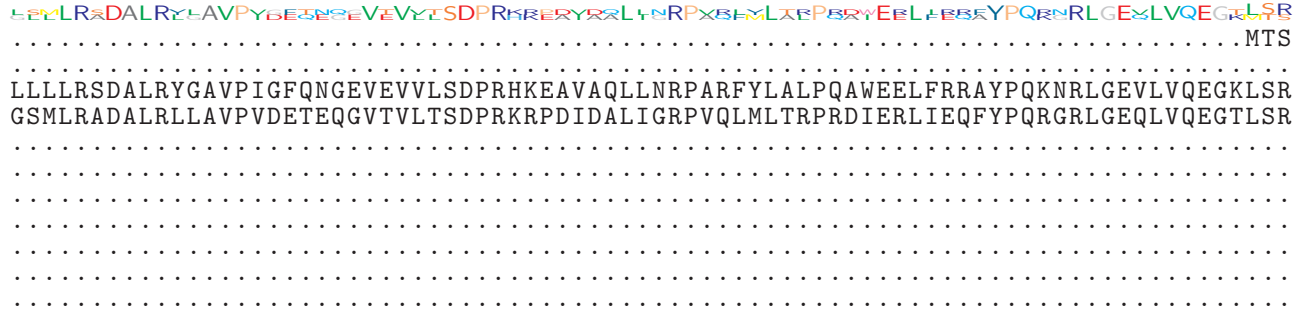
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

logo

	SGEGQKDLKLGELLQKQWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAA	
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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



3
0
317
317
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consensus

logo

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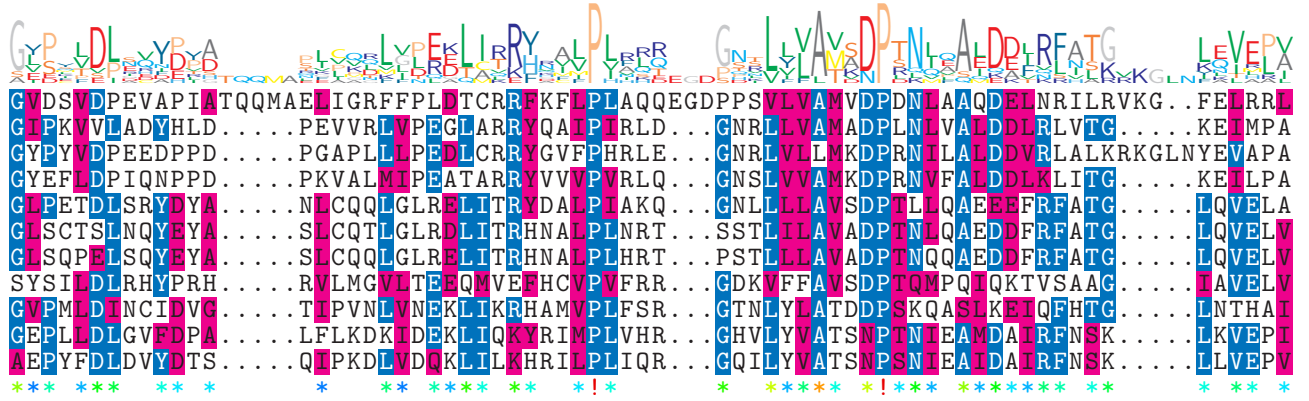


83
59
388
388
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57
56
65
65
65

consensus

logo

NTDB id 1410 SGL RS13950 WP 010873519.1
 NTDB id 153876 MOTHAS RS08015 WP 011393063.1
 NTDB id 1028 TT RS08215 WP 011173991.1
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 NTDB id 1293 VP RS12240 WP 005479695.1
 NTDB id 1110 NGFG RS09235 WP 003689817.1
 NTDB id 1251 GC085 RS07725 WP 011946523.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 NTDB id 1058 ABD1 RS01615 WP 001274986.1



161
126
460
455
124
124
124
123
132
132
132

consensus

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKSALTCR <small>X</small> TCAAEL <small>X</small> Q <small>X</small> EWLDCPYCMTSRFS	672
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NTDB id 1293 VP RS12240 WP 005479695.1	561
NTDB id 1110 NGFG RS09235 WP 003689817.1	558
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 ≥ 50% conserved