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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus		

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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consensus		

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consensus

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[illegible]

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NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus

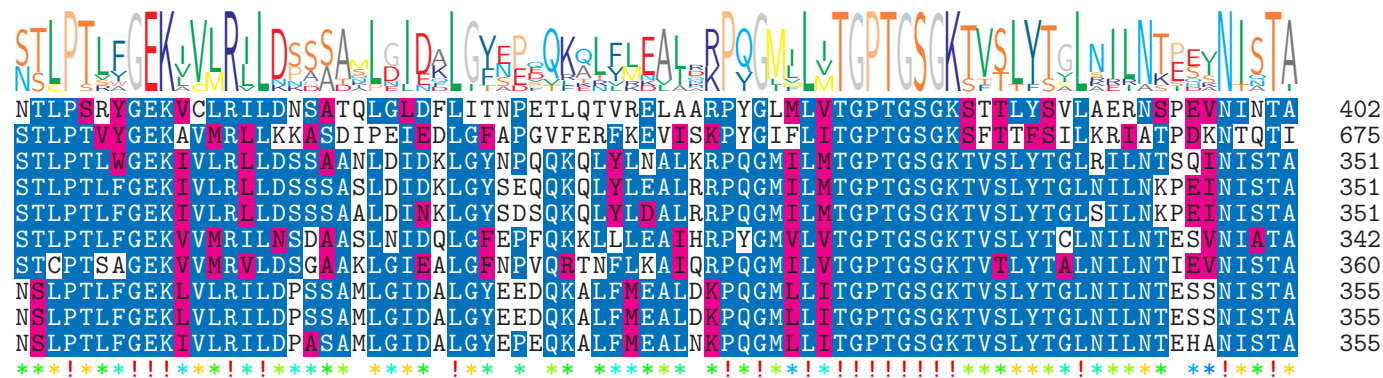
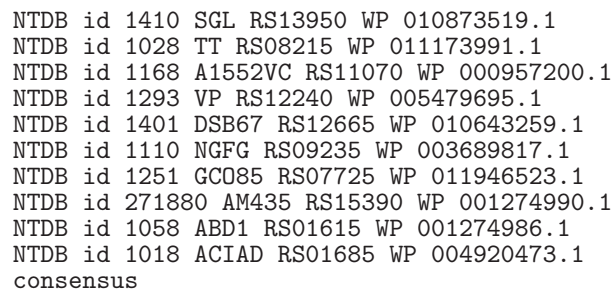
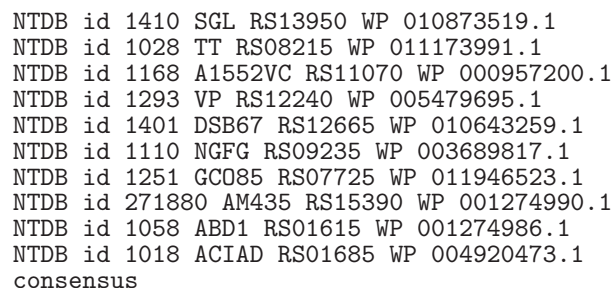
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4	GEILVELGLARPDVEEAL	60
5	QKRRGGRLDTLVQ	60
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7	LSIFHAQE	60
8	LTETQ	60
9	LSHIFGLP	60
10	MHSNLS	60
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13	KASGIS	60
14	MP	60
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38	MF	59
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42	IARV	59
43	SYS	59
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50	IALT	68
51	AAQNF	68
52	FGVP	68
53	MSALHTSP	68
54	KFTGFFRR	68
55	LVEEKH	68
56	VSAAT	68
57	MTALDAAK	68
58	RAKQDT	68
59	VAYLIE	68
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88	GEP	68

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consensus

[illegible]

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consensus



logo

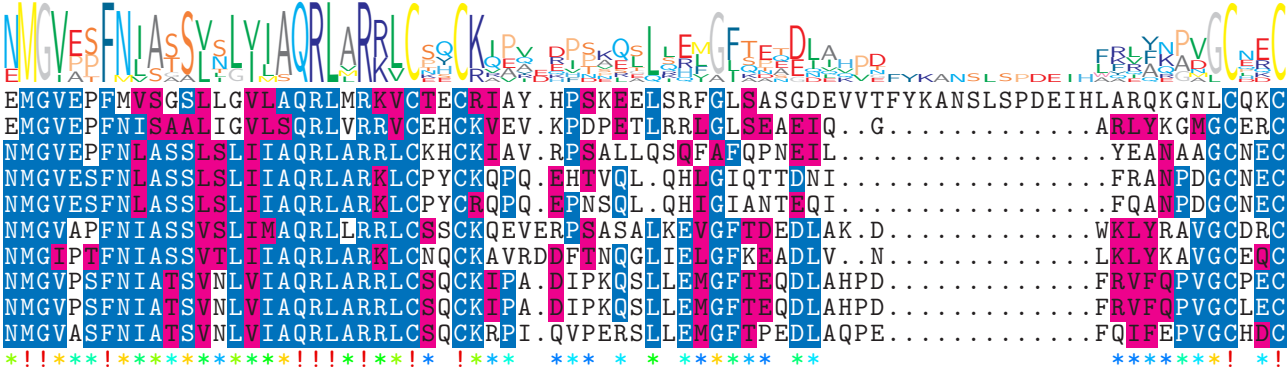
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consensus		



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431
431
422
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435
435
435

logo

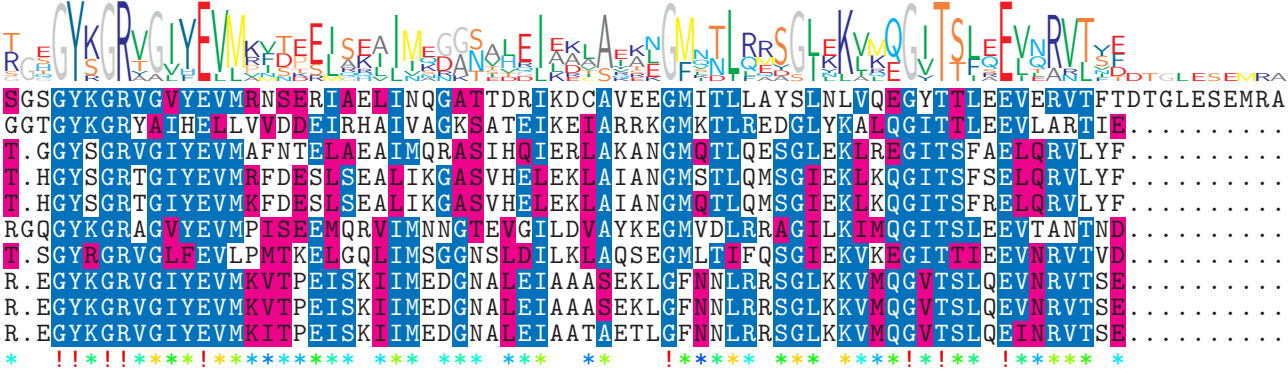
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NTDB id	271880	AM435 RS15390 WP 001274990.1
NTDB id	1058	ABD1 RS01615 WP 001274986.1
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consensus		



561
819
493
492
492
488
505
501
501
501

logo

NTDB id	1410	SGL RS13950 WP 010873519.1
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consensus		



641
889
562
561
561
558
574
570
570
570

logo

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NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved