

logo

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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQG YLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNL LLA VQPDP AAL	79
NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
NTDB id 1293	VP RS12240 WP 005479695.1	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
NTDB id 24809	NOC RS01780 WP 011330296.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
consensus		0

logo

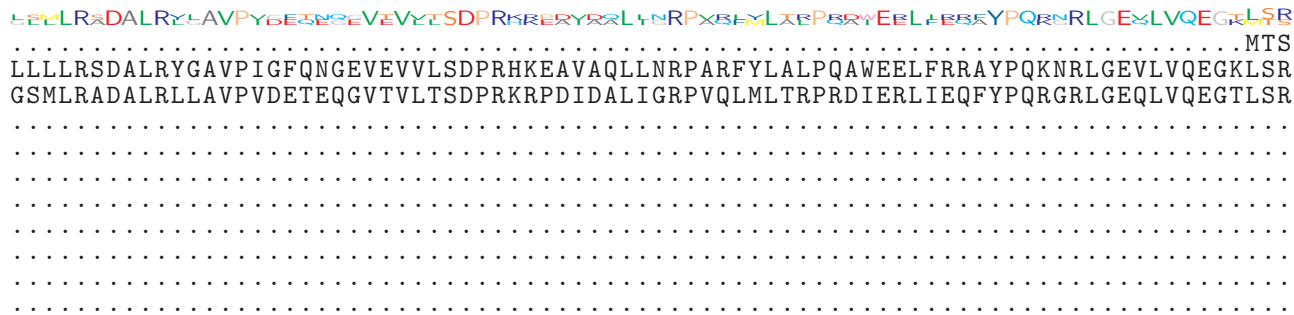
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logo

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NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
consensus		0

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NTDB id 1410 SGL RS13950 WP 010873519.1
 NTDB id 1028 TT RS08215 WP 011173991.1
 NTDB id 1306 DR RS10060 WP 027479822.1
 NTDB id 1168 A1552VC RS11070 WP 000957200.1
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 NTDB id 1251 GC085 RS07725 WP 011946523.1



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consensus

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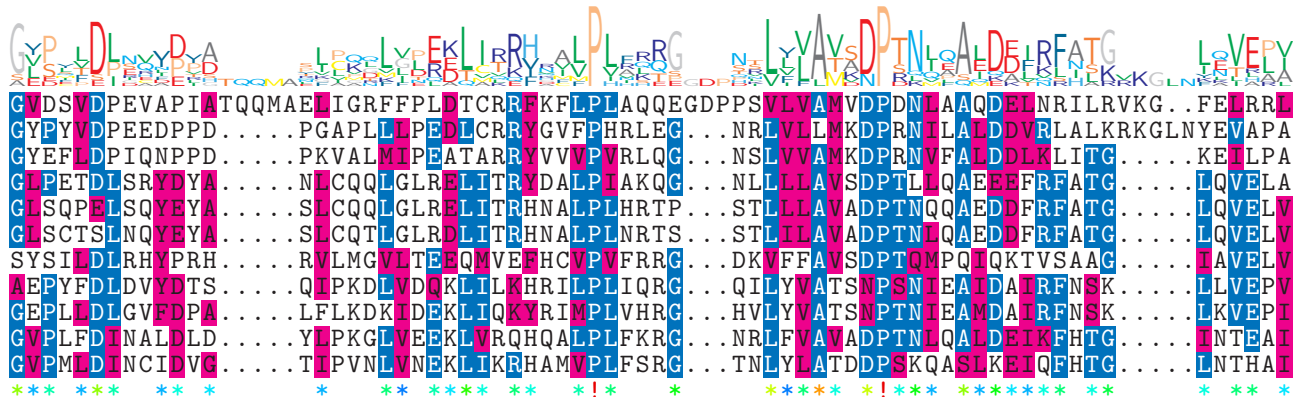


83
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consensus

logo

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 NTDB id 1306 DR RS10060 WP 027479822.1
 NTDB id 1168 A1552VC RS11070 WP 000957200.1
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 NTDB id 24809 NOC RS01780 WP 011330296.1
 NTDB id 1251 GC085 RS07725 WP 011946523.1



161
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consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKKSALTCR <small>X</small> TCAAEL <small>X</small> Q <small>X</small> EWLDCPYCMTSRFS	672
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NTDB id 1110 NGFG RS09235 WP 003689817.1	558
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
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NTDB id 24809 NOC RS01780 WP 011330296.1	571
NTDB id 1251 GC085 RS07725 WP 011946523.1	574
consensus		

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