

logo

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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 1401	DSB67 RS12665 WP 010643259.1		0
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NTDB id 1293	VP RS12240 WP 005479695.1		0
NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GCO85 RS07725 WP 011946523.1		0
NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

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		SLTPAEKAKELRAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPP..P	
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NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPVPP..P	158
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NTDB id 1293	VP RS12240 WP 005479695.1		0
NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GCO85 RS07725 WP 011946523.1		0
NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

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NTDB id 1293	VP RS12240 WP 005479695.1		0
NTDB id 1110	NGFG RS09235 WP 003689817.1		0
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NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

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NTDB id	1410	SGL RS13950 WP 010873519.1
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NTDB id	1028	TT RS08215 WP 011173991.1
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NTDB id	1058	ABD1 RS01615 WP 001274986.1
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consensus		

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NTDB id	1410	SGL RS13950 WP 010873519.1
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consensus		

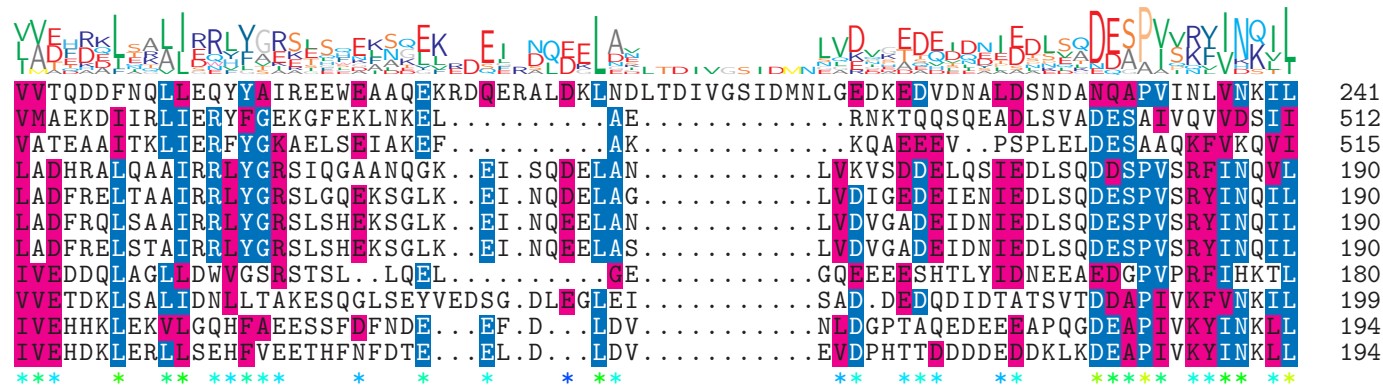
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.....MLTNLST	57
.....MQTNLST	57
.....MHSNLST	57
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.....MALATEEYRLQ	65
.....MSALHTSPKF	65
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logo

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		

GVDSVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQEGDPPSVLVAMVDPDNLAAQDELNRILRVKGF..ELRRL	161
GYEFLDPIQNPPD.....PKVALMPEATARRVVVPVRLQ...GNSLVVAMKDPRNVFALDDDLKLITG.....KEILPA	455
GYPYVDPEEDPPD.....PGAPLLLPEDLCRRYGVPFHRLE...GNRLVLMLKDPNRLALDDVRLALKRKGLNVEVAPA	460
GLPETDLSRYDYA.....NLCQQLGLRELITRYDALPIAKQ...GNLLLLAVSDPTLLQAEEDFRFATG.....LQVELA	124
GLSCTSLNQYEYA.....SLCQTLGLRDLITRHNALPLNRT...SSTLILAVADPTNLQAEDDFRFATG.....LQVELV	124
GLNQPELSQYEYA.....SLCQQLGLRELITRHNALPLQRT...PSTLLAVADPTNQQAEDDFRFATG.....LQIELV	124
GLSQPELSQYEYA.....SLCQQLGLRELITRHNALPLHRT...PSTLLAVADPTNQQAEDDFRFATG.....LQVELV	124
SYSILDLRHYPRH.....RVLMGVITEEQMVEFHCVPVFR...GDKVFFAVSDPTQMPQIQKTVSAAG.....IAVELV	123
GVPMLDINCDVIG.....TIPVNLVNEKLIKRRHAMVPLFSR...GTNLYLATDDPSKQASLKEIQFHTG.....LNTHAI	132
AEPYFDLDVYDTS.....QIPKDLVDQKLILKHRILPLIQR...GQILYVATSNPSNIEAIDAIRFNISK.....LLVEPV	132
GEPLIDLGVFDPA.....LFLKDKIDELIKQYRIMPLVHR...GHVLYVATSNPTNIEAMDIAIRFNISK.....LKVEPI	132

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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus
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consensus

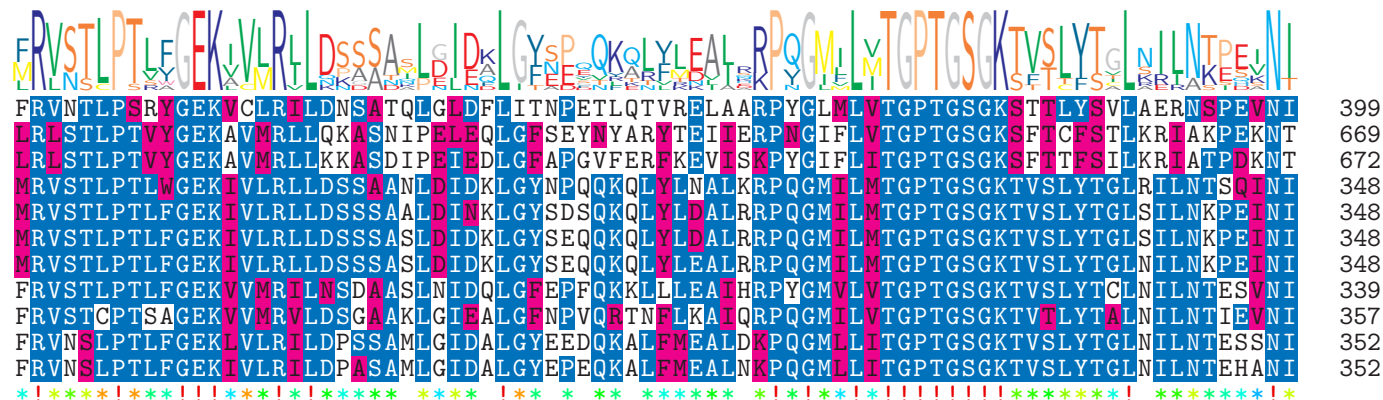
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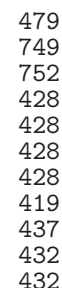
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consensus

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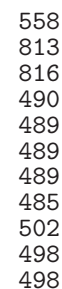
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consensus
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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

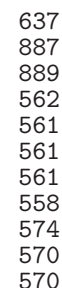
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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logo

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NTDB id 1168 A1552VC RS11070 WP 000957200.1		562
NTDB id 1401 DSB67 RS12665 WP 010643259.1		561
NTDB id 203710 K04M1 RS13545 WP 054730502.1		561
NTDB id 1293 VP RS12240 WP 005479695.1		561
NTDB id 1110 NGFG RS09235 WP 003689817.1		558
NTDB id 1251 GC085 RS07725 WP 011946523.1		574
NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved