

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
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NTDB id 1366 NTHI RS01950 WP 011271972.1	0
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NTDB id 1468 ASU2 RS05090 WP 014991734.1	0
NTDB id 39757 IA1 RS00745 WP 000650615.1	0
NTDB id 1436 BW25113 RS00530 WP 001025145.1	0
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 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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consensus		

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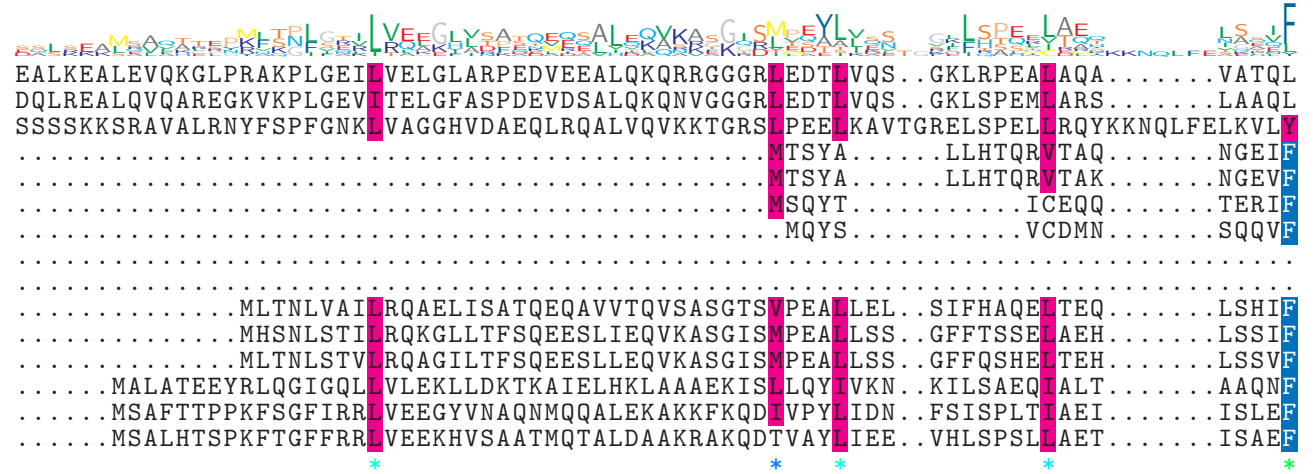
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NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVL																												237										
NTDB id 1410	SGL RS13950 WP 010873519.1																												0										
NTDB id 1366	NTHI RS01950 WP 011271972.1																												0										
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NTDB id 1251	GCO85 RS07725 WP 011946523.1																												0										
NTDB id 1018	ACIAD RS01685 WP 004920473.1																												0										
NTDB id 1058	ABD1 RS01615 WP 001274986.1																												0										
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NTDB id 1028	TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVLSDPRHKEAVAQLLNRPARFYLAALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR																																		317																																											
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NTDB id 1410	SGL RS13950 WP 010873519.1MTS																																		3																																											
NTDB id 1366	NTHI RS01950 WP 011271972.1																																		0																																											
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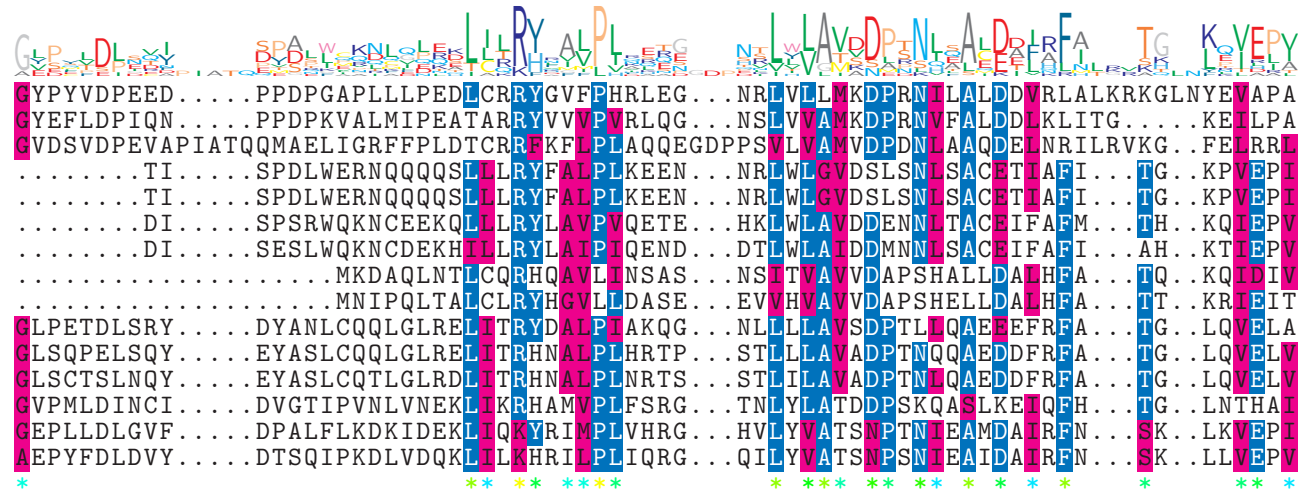
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 consensus



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logo

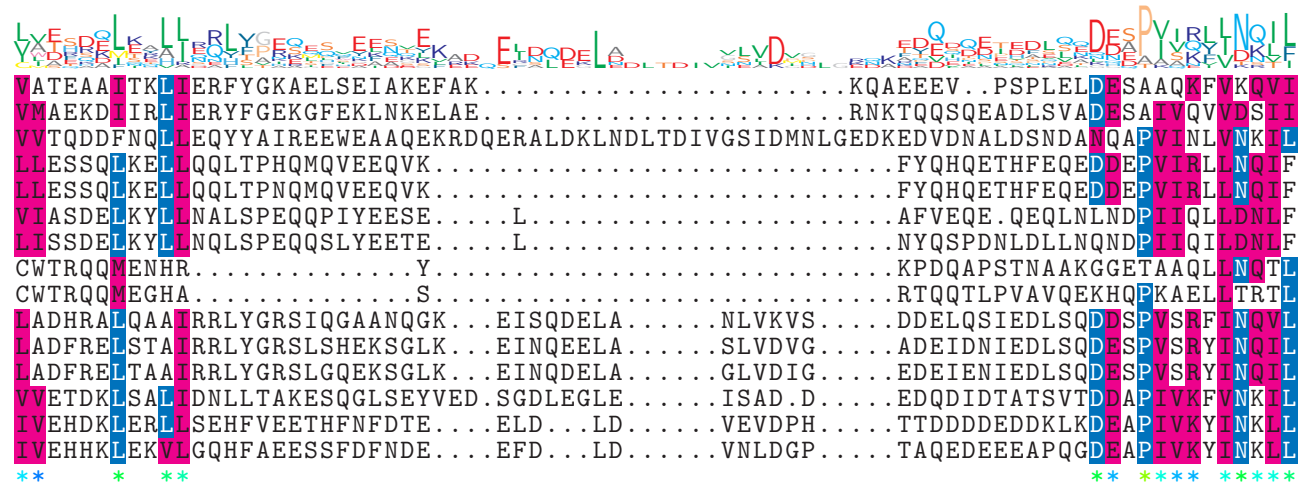
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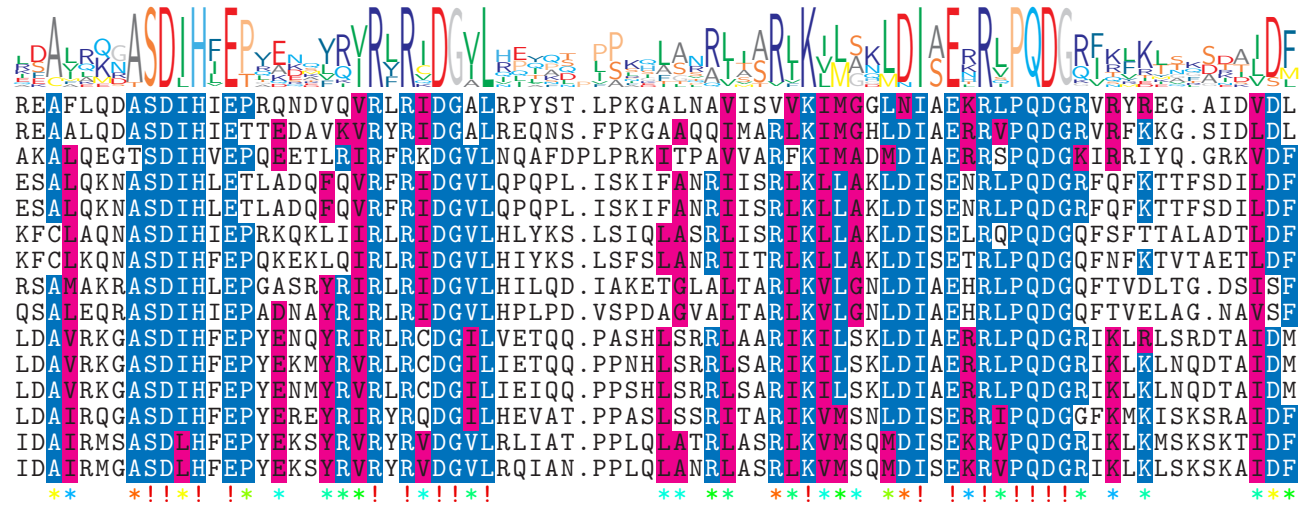
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 consensus



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consensus



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284
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logo

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 NTDB id 1251 GCO85 RS07725 WP 011946523.1
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1

consensus



753
750
480
369
369
364
364
327
327
429
429
429
438
433
433

logo

	MRAKRKSALTCRTCAAELQQEWLDCPYCMTSRFS	
NTDB id 1028 TT RS08215 WP 011173991.1	889
NTDB id 1306 DR RS10060 WP 027479822.1	887
NTDB id 1410 SGL RS13950 WP 010873519.1	MRAKRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
consensus		

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