

logo

		MSXLRIGDRRLGAALLDQGYLGDNDLQRALERHSEVGGRLADVLDISGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	
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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 81494	YZOS03 RS02850 WP 005379975.1	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

logo

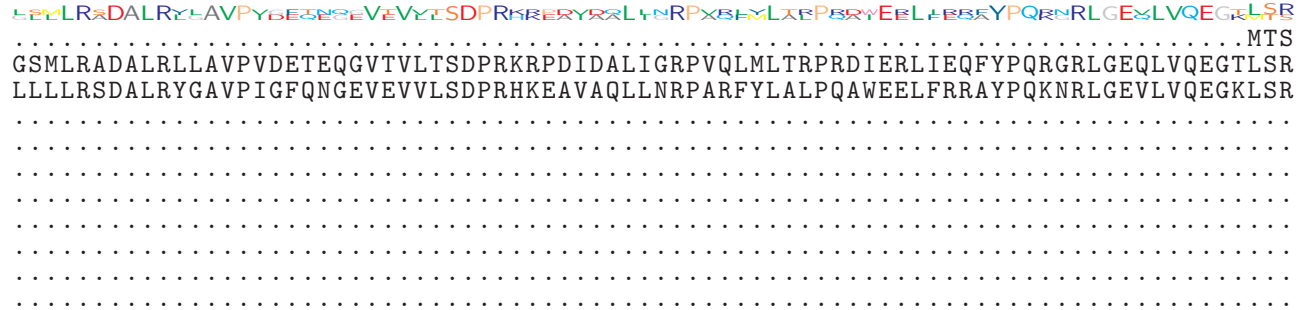
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consensus			

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		SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVL	
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NTDB id 1028	TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAA	237
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consensus			

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 NTDB id 81494 YZOS03 RS02850 WP 005379975.1
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 NTDB id 1110 NGFG RS09235 WP 003689817.1
 NTDB id 1251 GC085 RS07725 WP 011946523.1
 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



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logo

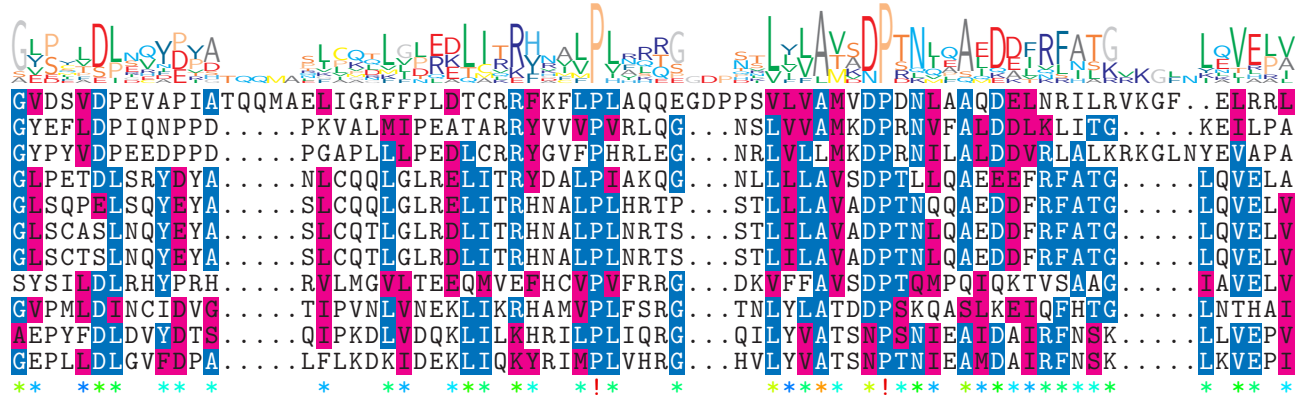
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 NTDB id 1401 DSB67 RS12665 WP 010643259.1
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 consensus



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logo

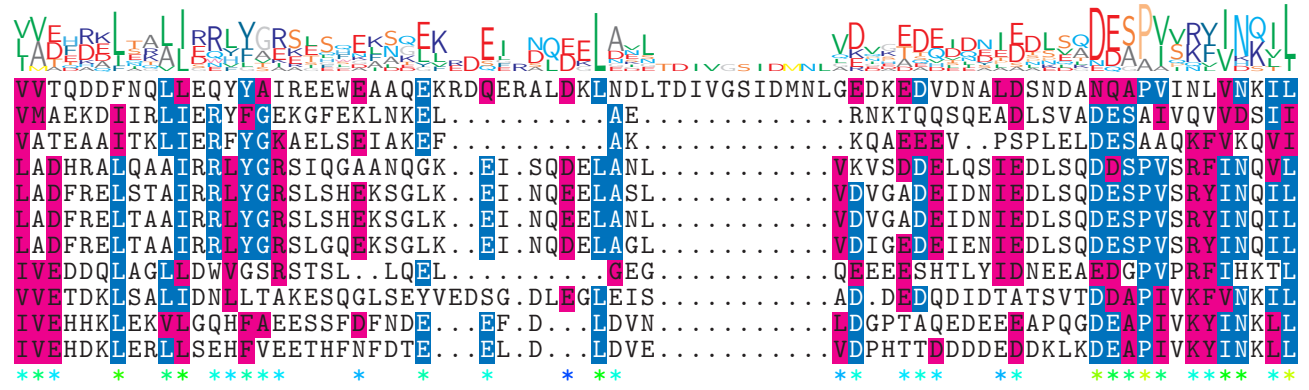
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 NTDB id 1251 GC085 RS07725 WP 011946523.1
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 consensus



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logo

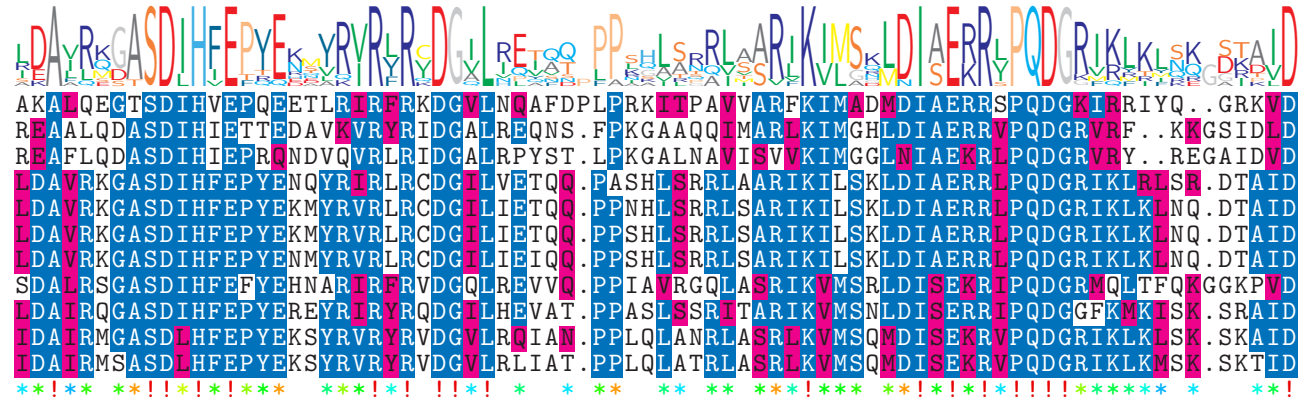
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



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512
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logo

NTDB id 1410 SGL RS13950 WP 010873519.1
 NTDB id 1306 DR RS10060 WP 027479822.1
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 NTDB id 1110 NGFG RS09235 WP 003689817.1
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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

NTDB id 1410 SGL RS13950 WP 010873519.1
 NTDB id 1306 DR RS10060 WP 027479822.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKRSALTCR <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 1251 GCO85 RS07725 WP 011946523.1	574
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
consensus		

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