

logo

		MSXLRIGDRRLGAALLDQGYLGDNDLQRALERHSEVGGRLADVLDISGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	
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NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLDISGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
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NTDB id 262409	CU052 RS08830 WP 005446257.1	0
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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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		RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIIVVPSGA	
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	
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NTDB id 1028	TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKQGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVDPDSAA	237
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



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logo

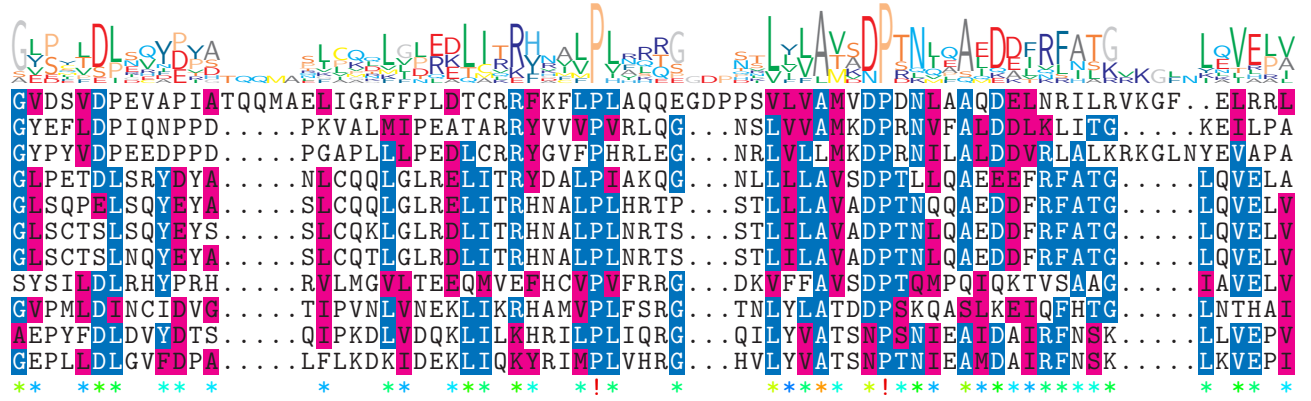
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 consensus



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logo

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 consensus



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logo

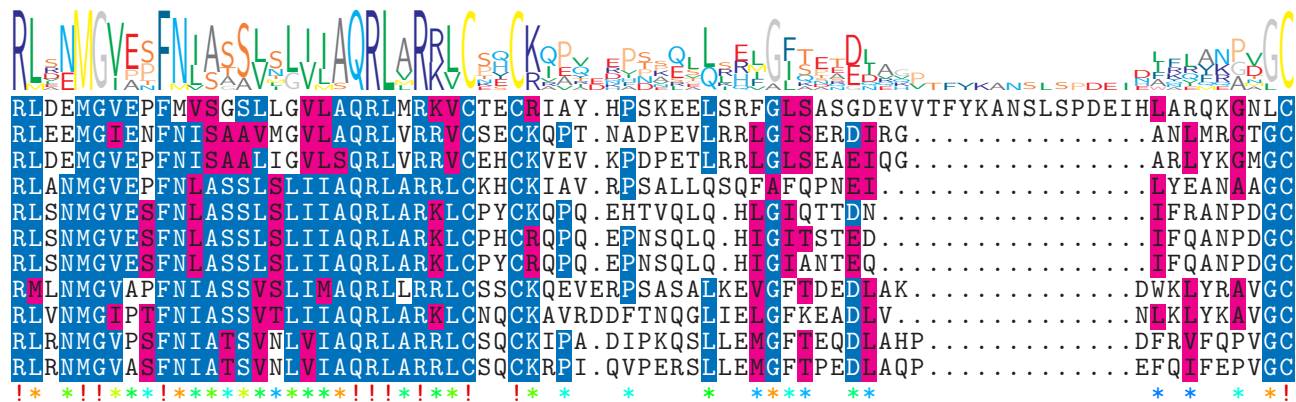
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logo

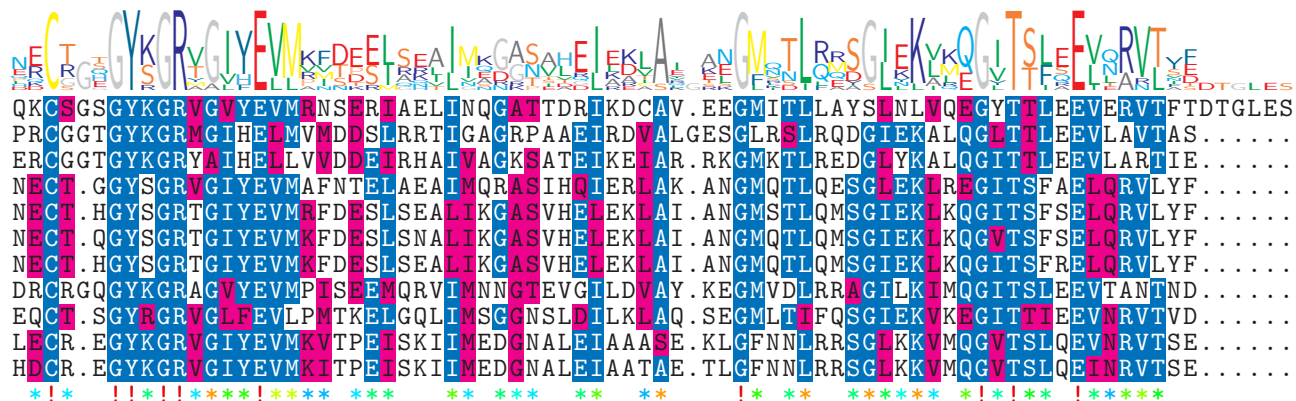
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logo

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 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 ≥ 50% conserved