

logo

		MSXLRIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLLGAAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384	A4U84 RS08525 WP 010787006.1	0
NTDB id 1355	HI 0298 AAC21962.1	0
NTDB id 119783	NTHI723 RS06930 WP 044364868.1	0
NTDB id 1366	NTHI RS01950 WP 011271972.1	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1293	VP RS12240 WP 005479695.1	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	0
NTDB id 1251	GC085 RS07725 WP 011946523.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
	consensus	0

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		RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
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		SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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NTDB id 1306 DR RS10060 WP 027479822.1	GSMLRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGR LGEQLVQEGT LSR	317
NTDB id 1028 TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEV LSDPRHKEAVAQLLNRPARFYLALPQAW EELFRRAYPQKNRLGEVLVQEGK LSR	317
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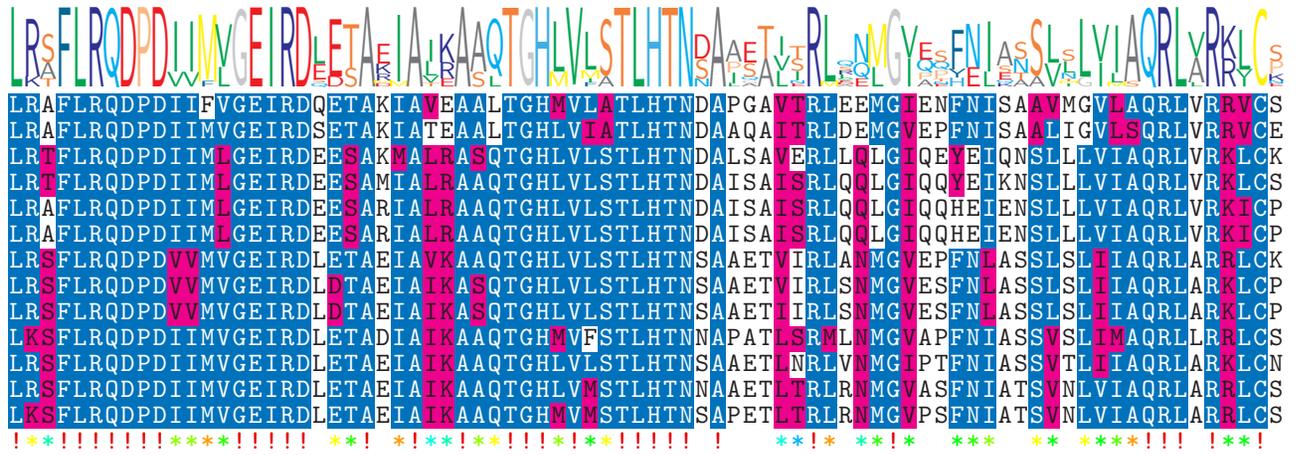
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NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEI LVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAAQAVATQLGYPYVDPEE	397
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NTDB id 1110 NGFG RS09235 WP 003689817.1	65
NTDB id 1251 GC085 RS07725 WP 011946523.1	74
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	74
consensus	74

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NTDB id 1306 DR RS10060 WP 027479822.1	NPPDPKVALM IPEATARRVYVVPVRLQGN SLVVMKDP RNVFALDD LKLTG.....	472
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NTDB id 1366 NTHI RS01950 WP 011271972.1	ISPDLWERNQQQQL LRLRYFALPLKEENRNLWLGVD SLSNLSACETIAFITG.....	96
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQ LGLRELITRYDALPIAKQGN LLLAVSDPTLLQAE EEFRFATG.....	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YEYASLCQT LGLRLITRHNA LPLNRTSSTLILAVADPTNLQAE DDFRFATG.....	141
NTDB id 1293 VP RS12240 WP 005479695.1	YEYASLCQQ LGLRELITRHNA LPLHRT PSTLLAVADPTNQQAE DDFRFATG.....	141
NTDB id 1110 NGFG RS09235 WP 003689817.1	YPRHRVLMG VLT EEQMV EHCVPVFRRGDK VFFAVSDPTQMPQIQKT VSAAG.....	140
NTDB id 1251 GC085 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIK RHAMVPLFVRG TNLYLATDPSKQASLKEI QFHTG.....	149
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQK LILKHRI LPLIQRGQI LYVATSNPSNIEAIDAIRFNSK.....	149
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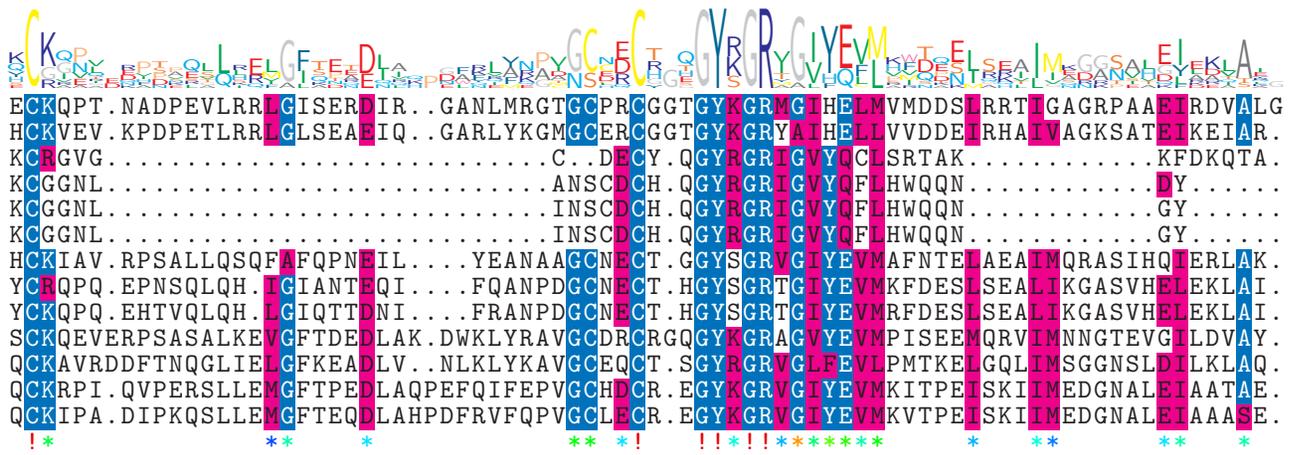


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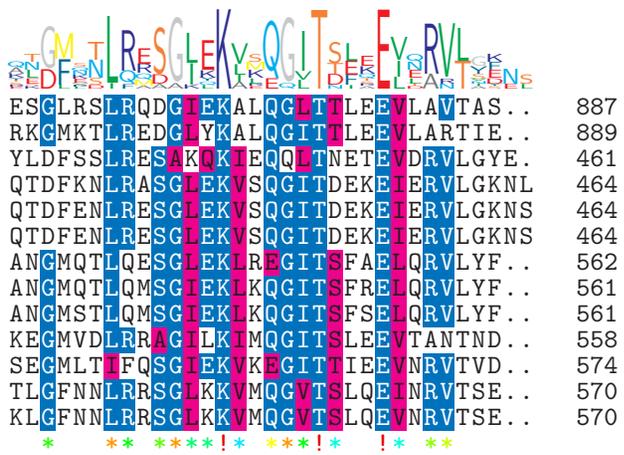


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consensus

 non conserved
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 $\geq 50\%$ conserved