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				MSXLSIGCDRLGALLLDXGGLDNLQRALERHSEVGGRLAEVLYDSQYERERIASXIEERFGIPLVQLHRVENSPPXXK	
NTDB id 1306	DR	RS10060	WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028	TT	RS08215	WP 011173991.1	MSVLTIGDKRLGAILLDAGLLTDEELQRALERHREVGGS�AEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1366	NTHI	RS01950	WP 011271972.1	.....	0
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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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				RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	
NTDB id 1306	DR	RS10060	WP 027479822.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVPEYQTTKSAFLYALAKHYPELGLPVPPP..P	157
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				SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	
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NTDB id 1306 DR RS10060 WP 027479822.1	GSMLRADALRLLAVPVDETEGGVTVLTSDPRKRPDIDALIGRPVQMLLTPRPDIERLIEQFYPQGRGLGELVQEGTL	317
NTDB id 1028 TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEIVELGLARPEDVEEALQKQRRGGGRLEDTLVQSCKIRPEALAAQAVATQLGYPVVDPEE	397
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....MLTNLSTVLRQAGILTFSSQEESSLEQVKASGISMPEALLSSGFFQSHETHEHSSVFGLSCTSLNQ	66
NTDB id 1293 VP RS12240 WP 005479695.1	.....MHSNLSTILRQKGLLTFSSQEESSLEQVKASGISMPEALLSSGFFTSSELAEHSSVFGLSQPFLSQ	66
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....MSVGLLRILVQNQVVTVQAEHYYNESQAGKEVLPMLFSDGVISPKSLAALIAVRFYSISILDLRH	65
NTDB id 1251 GCO85 RS07725 WP 011946523.1	.....MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYLVKNKILSAEQIALTAQNFQVPMLDINC	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDTPVYLDNFSISPLTIAEISLEFGEPLLDLGV	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....MSALHTSPKFTGFFRRILVEEKHVSAATMQTALDAAKRAKQDTPVAYLIEEVHISPSLLAETISAEFAEPYFDLDV	74
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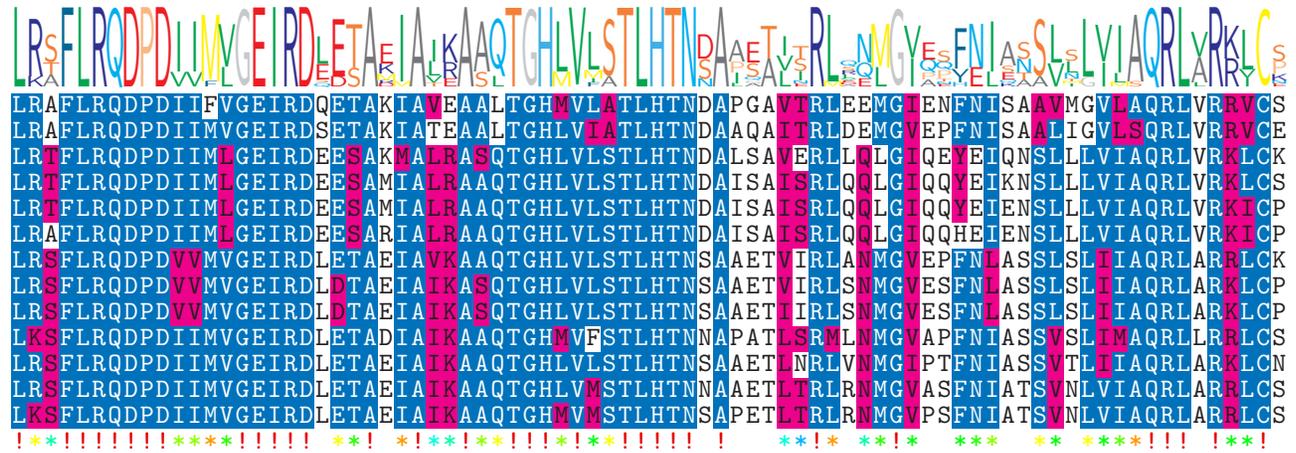
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NTDB id 1028 TT RS08215 WP 011173991.1	DPPDPGAPLILPEDTCRRYGVFPHRLEGNRLVLLMKDPRNLLALDDVRLALKRKGLNVEVAPAVATEAAITKLIERFYGK	477
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NTDB id 1355 HI 0298 AAC21962.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPN	96
NTDB id 1140664 DQN24 RS03295 WP 111695393.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPR	96
NTDB id 1366 NTHI RS01950 WP 011271972.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPH	96
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQILGLRELITRYDALPIAKQGNLLAVSDPTLLQAEFEFRFATG.....LQVELALADHRAALQAAIRRLYGR	141
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDPSKQASLKEIQFHTG.....LNTHAIVVETDKLSALIDNLT	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLKDKIDEKLIQKVRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSK.....LKVEPIVEHDKLERLSEHFVE	149
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLKRRILPLIQRGQILYVATSNPSNIEATDAIRFNSK.....LLVEPVIVEHKKLEKVLGQHFAE	149
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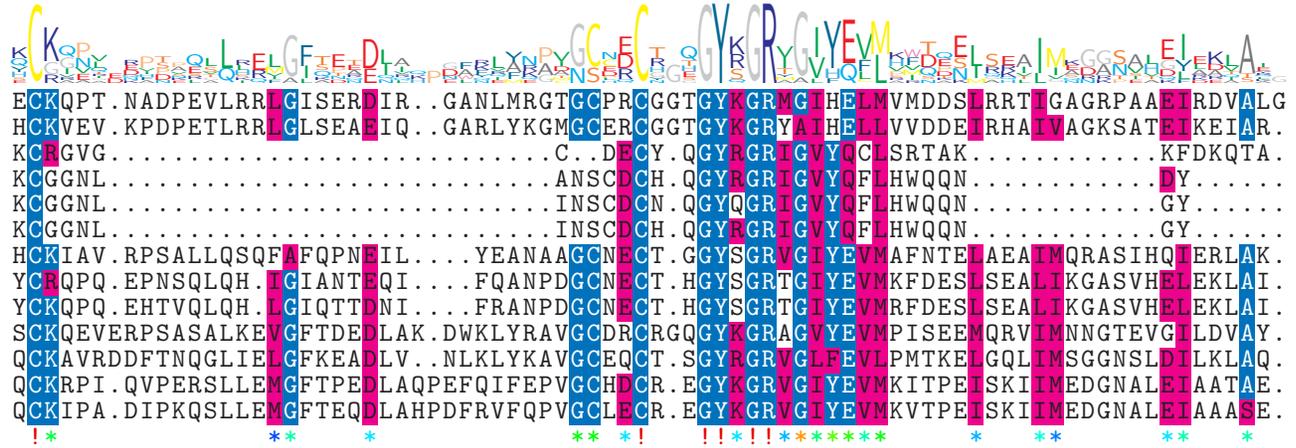
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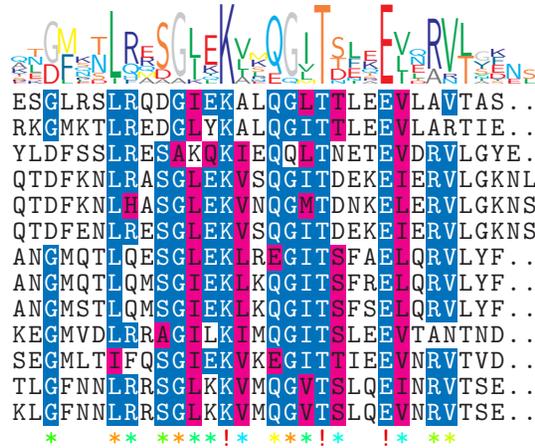
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