

logo

		MSXLRIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1384	A4U84 RS08525 WP 010787006.1	0
NTDB id 1355	HI 0298 AAC21962.1	0
NTDB id 1065049	ACHWYH RS05740 WP 112103477.1	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 1058	ABD1 RS01615 WP 001274986.1	0
consensus		0

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		SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1028	TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
consensus		0

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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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NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus	0

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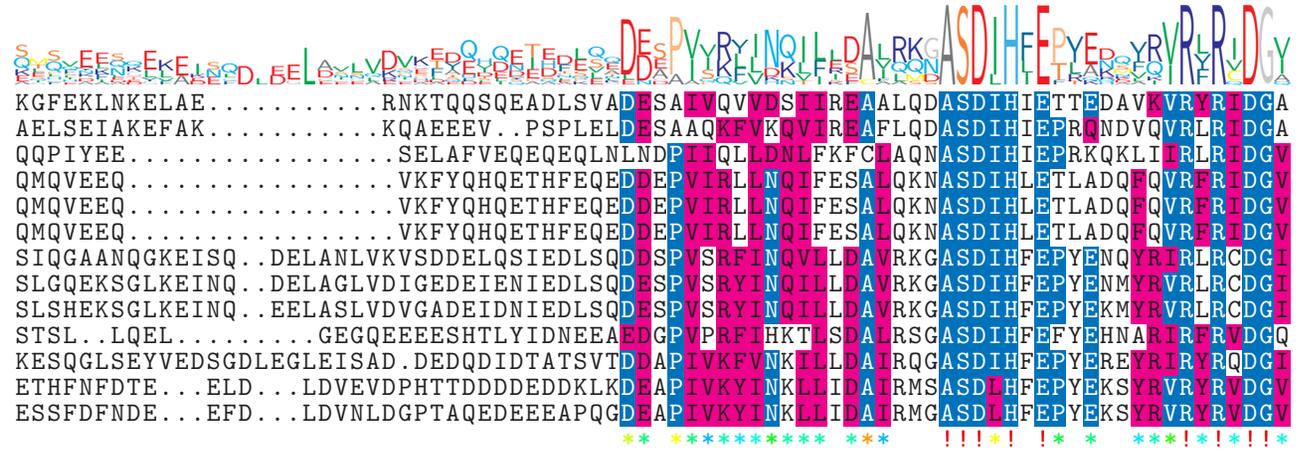
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NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEI LVELGLARPEDVEEALQKQRGGGRLEDTLVQSGKLRPEALAAQAVATQLGYPYVDPEE	397
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NTDB id 1293 VP RS12240 WP 005479695.1	66
NTDB id 1110 NGFG RS09235 WP 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 IPEATARRYVVV PURLQGNSLVVAMKDP RNVFALDD LKLTG.....	472
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NTDB id 1366 NTHI RS01950 WP 011271972.1	ISPDLWERNQQQQS L LRYFALPLKEENR L WLGVD SLSNLSACETIAFITG.....	96
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQ L GLRELITRYDALPIAKQGN L LLLAVSDPTLLQAE EEFRFATG.....	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YEYASLCQT L GLRLITRHNA L PLNRTSSTL L LAVADPTNLQAE DDFRFATG.....	141
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDL V DQK L L LKHRI L PLIQRGQV L YVATSNPSNIEAIDAIRFNSK.....	149
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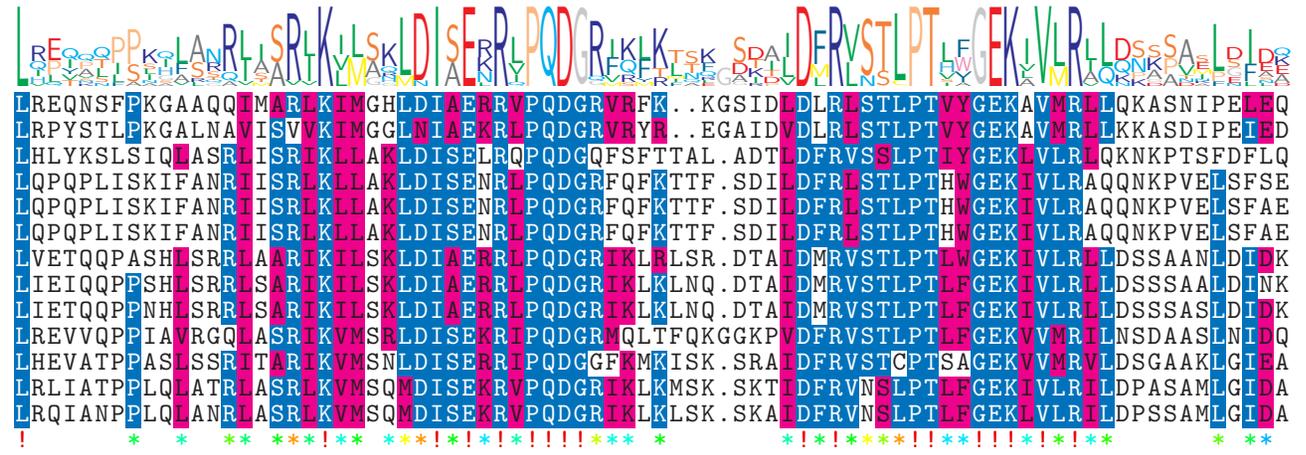
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 consensus



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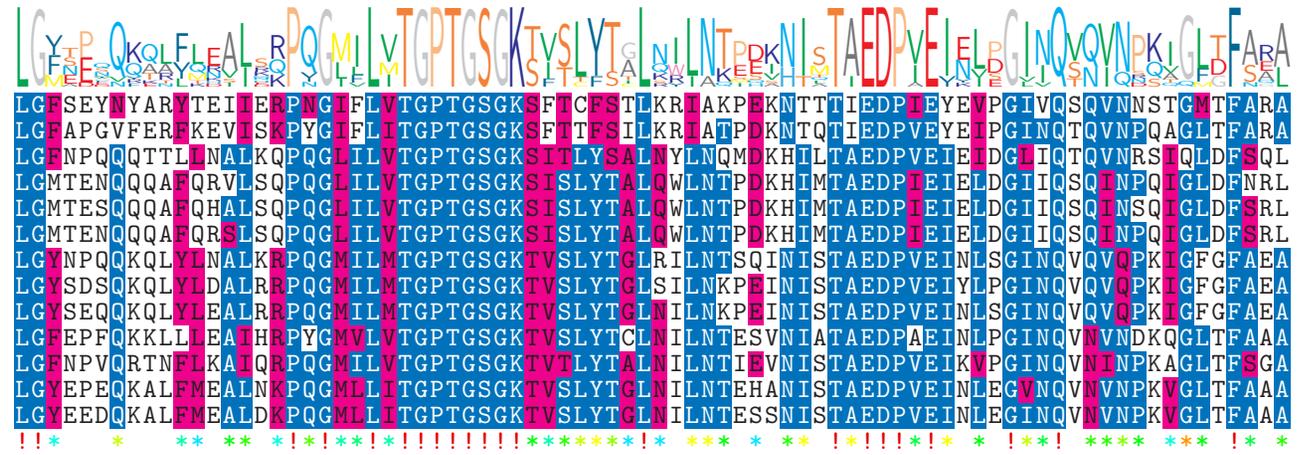
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



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NTDB id 1306 DR RS10060 WP 027479822.1
 NTDB id 1028 TT RS08215 WP 011173991.1
 NTDB id 1384 A4U84 RS08525 WP 010787006.1
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 consensus



699
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 non conserved
 similar
 $\geq 50\%$ conserved