

logo

		MSXLEIIGDRRLGAALLDAGLLTDEELQRALERHREVEVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVEVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1097939	ACM26V RS24600 WP 416149386.1	0
NTDB id 96	BSU 24730 NP 390353.1	0
NTDB id 1410	SGL RS13950 WP 010873519.1	0
NTDB id 1384	A4U84 RS08525 WP 010787006.1	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 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		REYRKAQAKNLRAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEIVVPSGA	157
NTDB id 1097939	ACM26V RS24600 WP 416149386.1	0
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consensus			

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		SACSEKLEELLYKGYIIRALRYALMQRTEGSLGRTLYSRKIFRERLYEXLAEQEGTEFLESTEGMPEEXK	237
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consensus			

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NTDB id 1028	TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVLS DPRHKEAVAQLLNRPARFYALALPQAW EELFRRAYPQKNRLGEVLVQEGKLSR	317
NTDB id 1306	DR RS10060 WP 027479822.1	GSMLRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSR	317
NTDB id 1097939	ACM26V RS24600 WP 416149386.1	0
NTDB id 96	BSU 24730 NP 390353.1	0
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NTDB id 1384	A4U84 RS08525 WP 010787006.1	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
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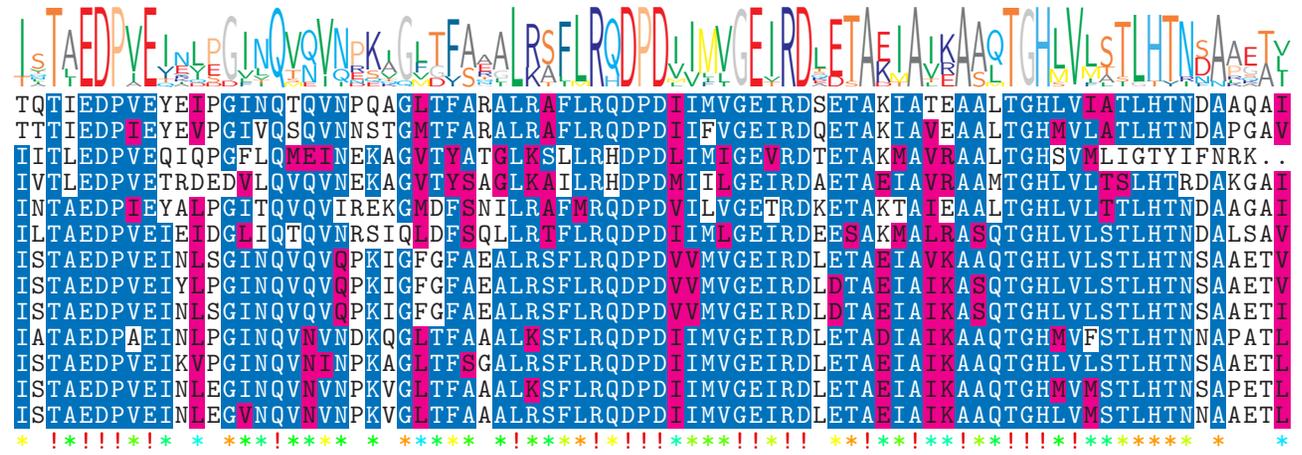
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NTDB id 1306	DR RS10060 WP 027479822.1	DQLREALQVQAREGKVKPLGEV ITELGFASPDEVDSA LKQKNVGGGRLED TLVQS. GKLSPEMLARS..... LAAQL	388
NTDB id 1097939	ACM26V RS24600 WP 416149386.1	0
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NTDB id 1168	A1552VC RS11070 WP 000957200.1	57
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NTDB id 1251	GC085 RS07725 WP 011946523.1	65
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	65
consensus		65

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NTDB id 1028	TT RS08215 WP 011173991.1	GYPYVDPEEDPPDP..... GAPLLPEDLCRRYGVFPHRLE... GNRVLVLMKDPRNILALDDVRLALKRKG LNYEVAPA	460
NTDB id 1306	DR RS10060 WP 027479822.1	GYEFLDPIQNPPDP..... KVALMPEATARRYVVV PVR LQ... GNSLVVAMKDPRNVFALDDLKLITG..... KEILPA	455
NTDB id 1097939	ACM26V RS24600 WP 416149386.1	0
NTDB id 96	BSU 24730 NP 390353.1	0
NTDB id 1410	SGL RS13950 WP 010873519.1	GVDSVDPEVAPIATQMAELIGRFFPLDTCRRFKFLPLAQEGDPPSVLVAMVDPDNLA AQDELNRILRVKG.. FELRRL	161
NTDB id 1384	A4U84 RS08525 WP 010787006.1	TERIFDI... SPSR... WQKNCEEKQLLRYLAVPVQET... EHKLWLAVD DENNLTA CEIFA... FMTH... KQIEPV	74
NTDB id 1168	A1552VC RS11070 WP 000957200.1	GLPETDLSRYDYAN... LCQQLGLRELITRYDALPIAKQ... GNLLLLAVSDPTLIQAEEFR... FATG.. LQVELA	124
NTDB id 1401	DSB67 RS12665 WP 010643259.1	GLSCTSLNQY EYAS... LCQTLGLRDLITRHNA LPLNRT... SSTLILAVADPTNLQAEDDFR... FATG.. LQVELV	124
NTDB id 1293	VP RS12240 WP 005479695.1	GLSQPELSQY EYAS... LCQQGLGLRELITRHNA LPLHRT... PSTLLAVADPTNQAEDDFR... FATG.. LQVELV	124
NTDB id 1110	NGFG RS09235 WP 003689817.1	SYSILDLRHYPRHR... VLMGVLTEE QMVEFHCVPVFRR... GDKVFFAVSDPTQMPQIQKT V... SAAG.. IAVELV	123
NTDB id 1251	GC085 RS07725 WP 011946523.1	GVPMLDINCIDVGT... IPVNLVNEKLIKRRHAMVPLFSR... GTNLVYATDTPSKQASLKEIQ... FHTG.. LNTHAI	132
NTDB id 1058	ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTSQ... IPKDLVDNKLILKHRILPLIQR... GQILYVATSNPSNIEAIDAIR... FNSK.. LLVEPV	132
NTDB id 1018	ACIAD RS01685 WP 004920473.1	GEPLLDLVGFDPAL... FLKDKIDEKLIQKYRIMPLVHR... GHVLYVATSNPTNIEAMD AIR... FNSK.. LKVEPI	132
consensus		132

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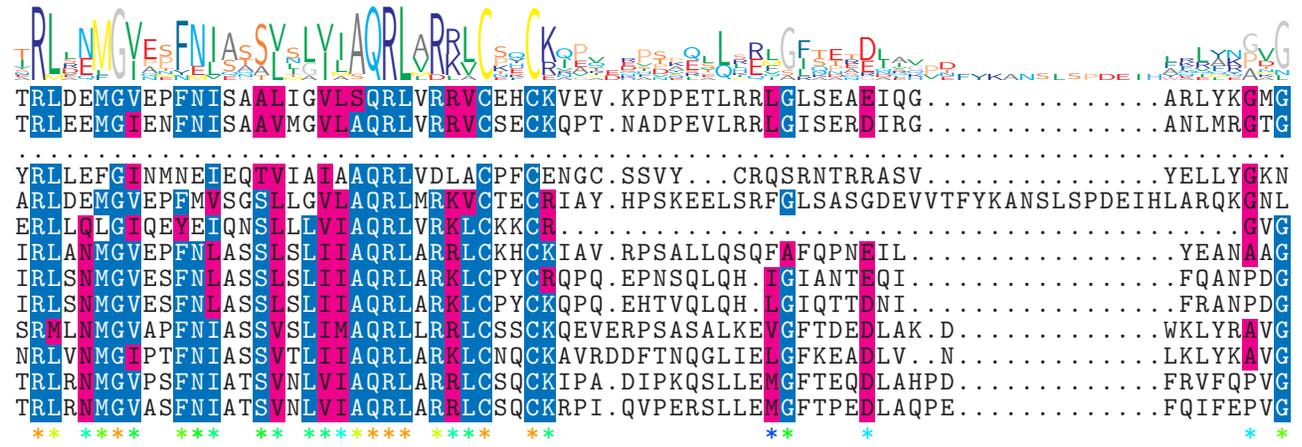
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



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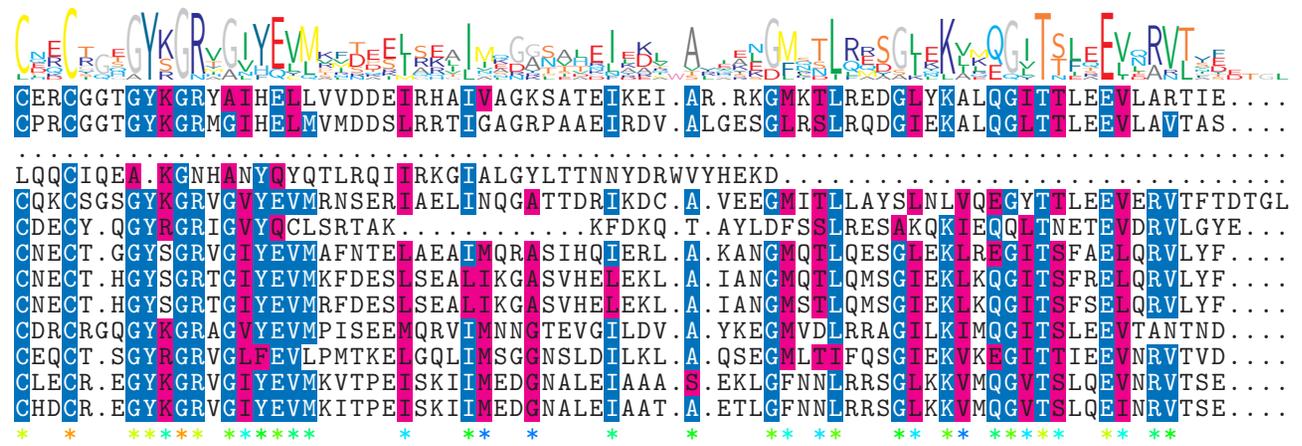
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 NTDB id 1251 GC085 RS07725 WP 011946523.1
 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



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NTDB id 1028 TT RS08215 WP 011173991.1
 NTDB id 1306 DR RS10060 WP 027479822.1
 NTDB id 1097939 ACM26V RS24600 WP 416149386.1
 NTDB id 96 BSU 24730 NP 390353.1
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 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



889
887
247
356
635
461
562
561
561
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574
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570

logo

NTDB id 1028 TT RS08215 WP 011173991.1	ESEMRAKRKSALTCRTCAAELQQEWLDCPYCMTSRFS	889
NTDB id 1306 DR RS10060 WP 027479822.1	887
NTDB id 1097939 ACM26V RS24600 WP 416149386.1	247
NTDB id 96 BSU 24730 NP 390353.1	356
NTDB id 1410 SGL RS13950 WP 010873519.1	ESEMRAKRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
NTDB id 1384 A4U84 RS08525 WP 010787006.1	461
NTDB id 1168 A1552VC RS11070 WP 000957200.1	562
NTDB id 1401 DSB67 RS12665 WP 010643259.1	561
NTDB id 1293 VP RS12240 WP 005479695.1	561
NTDB id 1110 NGFG RS09235 WP 003689817.1	558
NTDB id 1251 GC085 RS07725 WP 011946523.1	574
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
consensus		

- non conserved
- similar
- ≥ 50% conserved