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NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....	0
NTDB id 1293 VP RS12240 WP 005479695.1	.....	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 154229 WUE2121 RS09920 WP 002221967.1	.....	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		

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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIYVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSEEEVLDLGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....	0
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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTDLLGRILV.RKGLPEEALYRALAEQKGLFLESTEGIVPDPSAA	237
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NTDB id 1306 DR RS10060 WP 027479822.1	LSMLRSDALRYLAVPYBESNREVEVYVSDPRKREERYRRLYLRPXSHMLTSPBRYEELHESYYPQERRLGEXLVQEGKLSR	317
NTDB id 1028 TT RS08215 WP 011173991.1	GSMLRADALRLLAVPVDETEQQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGELVQEGTLSR	317
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NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
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NTDB id 1306 DR RS10060 WP 027479822.1	DQLREALLQVQAREGKVKPLGEVITELGFASPEVDSALQKQNVGGGRLDRTLQVSGKLSPEMLARSLAAQLGYEFLDPIQ	397
NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRGGRLDRTLQVSGKLRPEALAQAVATQLGYPYVDPEE	397
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....	66
NTDB id 1293 VP RS12240 WP 005479695.1	.....	66
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	66
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	74
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consensus	.....	74

logo

NTDB id 1306 DR RS10060 WP 027479822.1	NPPDPKVALMIP EATARRVVPVRLQGNLSLVAMKDPN VFALDDLKLITG.....KEILPAVMAEKDIIRLIERYFGE	472
NTDB id 1028 TT RS08215 WP 011173991.1	DPPDPGAPLLIPEDLCRRYGVFPHRLEGNRLVLLMKDPRNILALDDVRLALKRKGGLNYEVAPAVATEAAITKLIERYFGK	477
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQQLGLRELITRYDALPIAKQGNLLLLAVSDPTLLQAE EEFRFATG.....LQVELALADHRALQA AIRRLYGR	141
NTDB id 1293 VP RS12240 WP 005479695.1	YEYASLCQQQLGLRELITRHNA LPHRTPSTLLAVADPTNQQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YEYASLCQTLGLRDLITRHNA LPLNRTSSTLILAVADPTNIQAEDDFRFATG.....LQVELVLADFRELTA AIRRLYGR	141
NTDB id 154229 WUE2121 RS09920 WP 002221967.1	YPRHRVLMGVLT EEQMVEFHCVPVFRRGDKVFFAVSDPTQMPQIQKTVSAAG.....IEVELVIVEDDQLAGLLDWVGSR	140
NTDB id 1110 NGFG RS09235 WP 003689817.1	YPRHRVLMGVLT EEQMVEFHCVPVFRRGDKVFFAVSDPTQMPQIQKTVSAAG.....I AVELVIVEDDQLAGLLDWVGSR	140
NTDB id 1251 GC085 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVVETDKLSALIDNLLTA	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLKDKID EKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMD AIRFNSK.....LKVEPIIVEHDKLERLLSEHFVE	149
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQK LILKHRILPLIQRGQILYVATSNPSNIEAID AIRFNSK.....LLVEPVIVEHKKLEKVLGQHFAE	149
consensus	.....	149



logo

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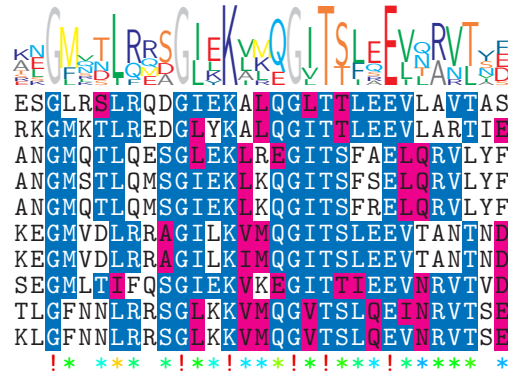
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 consensus



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- non conserved
- similar
- ≥ 50% conserved