

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1355 HI 0298 AAC21962.1	0
NTDB id 1366 NTHI RS01950 WP 011271972.1	0
NTDB id 311917 D1111 RS04360 WP 005601209.1	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 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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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPFALEGDRLLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGASDA	160
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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 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
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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	240
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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 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
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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	320
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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 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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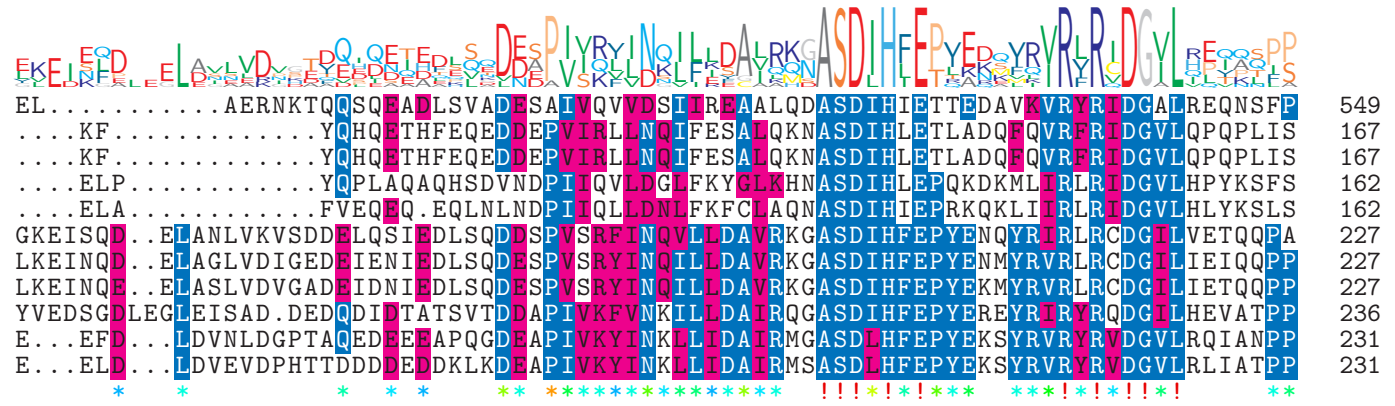
NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQAREGKVKPLGEVITELGFASPDEVDSA LQKQNVGGGRLEDTLVQSGKLSPEMLARSLAAQLGYEFLDPIQNPP	400
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	69
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	77
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMPEATARRVYVVPVRLQGNLVLVAMKDPRNVFALDDLKLTGKEILPAVMAEKDIIRLIERYFGEKGFELKLNK	480
NTDB id 1355 HI 0298 AAC21962.1	DLWERNQQQQSLILRYFALPLKEENRRLWLGVDLSNLSACETIAFITGKPVPELLESSQLKELLQQLTPNQMQVEEQV	104
NTDB id 1366 NTHI RS01950 WP 011271972.1	DLWERNQQQQSLILRYFALPLKEENRRLWLGVDLSNLSACETIAFITGKPVPELLESSQLKELLQQLTPHQMQVEEQV	104
NTDB id 311917 D1111 RS04360 WP 005601209.1	EQWQKNYDERHILRLRYLTVPLREDQTLWLAIDDMNLSACEIFAFVTHKRTEPVLVSSEELKYLGLLQMLPEQQSVYDET	98
NTDB id 1384 A4U84 RS08525 WP 010787006.1	SRWQKNCEEKQLLRLRYLAVPVQETEHKLLAVDDENNLTACEIFAFMTHKQIEPVVIASDELKYLNLALSPEQQPIYEEES	99
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQLGLRELITRYDALPIAKQGNLILLAVSDPTLLQAEEEFRFATGLQVELALADHRAALQAAIRRLYGRSIQGAANQ	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLAADFRELTAAIRRLYGRSLGQEKSG	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQLGLRELITRHNALPLHRTPTSTLLAVADPTNQAEDDFRFATGLQVELVLAADFRELSTAIRRLYGRSLSHEKSG	149
NTDB id 1251 GC085 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLVYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSE	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLILKHRILPLIQRGQILYVATSNSNIEAIDAIRFNSKLLVEPVIVEHHKLEKVLGQHFVAEESSFDND	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNTPTNIEAMDAIRFNSKLLKVEPIIVEHDKLERLLSEHFAVEETHFNFD	157
consensus		

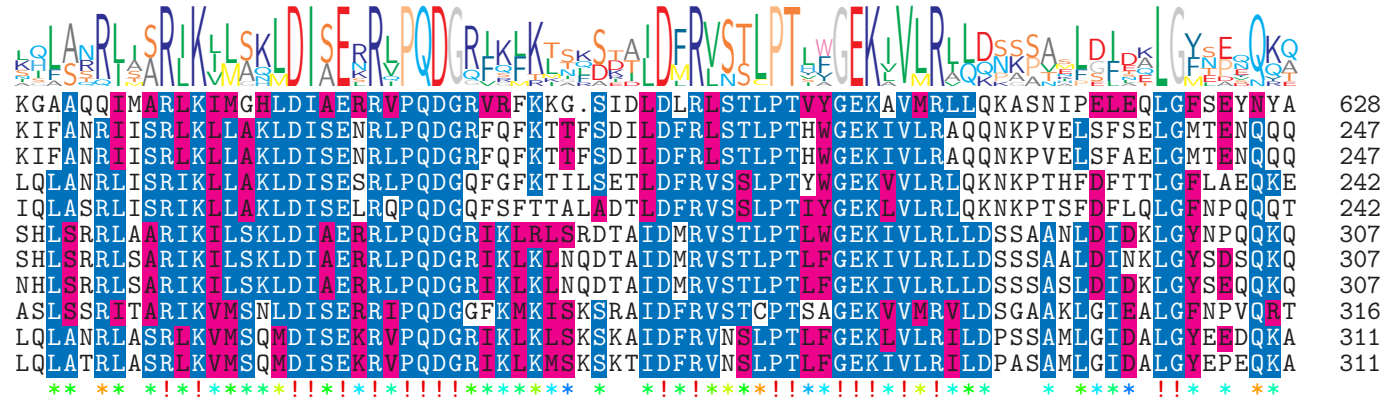
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