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NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEIVVPSGASDA	160
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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	240
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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDI DAL I GRPVQLMLTRPRDI ER L I EQFYPPQRGR L GEQLVQEGT LSRDQL	320
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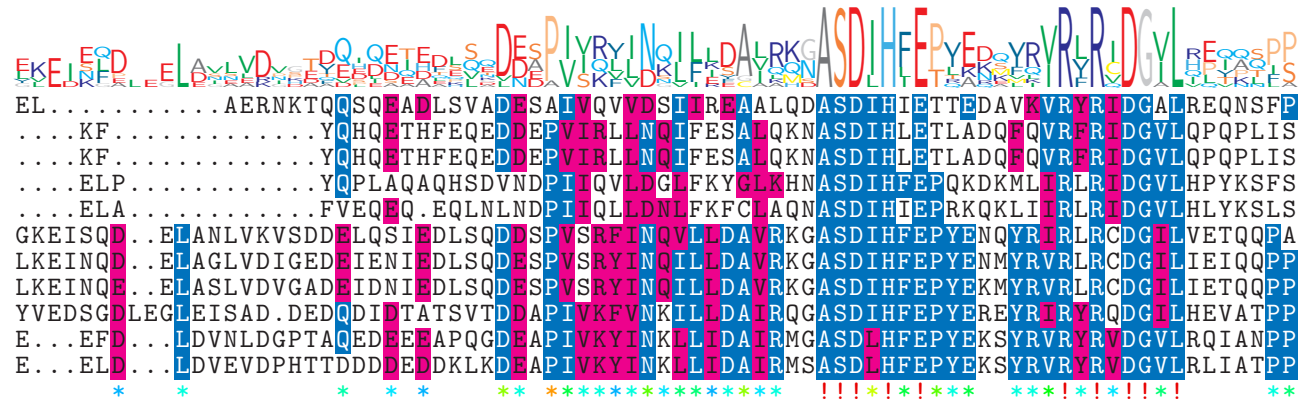
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NTDB id 1355 HI 0298 AAC21962.1	DLWERNQQQQSLILRYFALPLKEENRRLWLGVDLSNLSACETIAFITGKPVPELLESSQLKELLQQLTPNQMQVEEQV	104
NTDB id 1366 NTHI RS01950 WP 011271972.1	DLWERNQQQQSLILRYFALPLKEENRRLWLGVDLSNLSACETIAFITGKPVPELLESSQLKELLQQLTPHQMQVEEQV	104
NTDB id 312232 D1097 RS04745 WP 005597396.1	EQWQKNYDERHILRLRYLTVPLREDQTLWLAIDDMNLSACEIFAFLTTHKRTEPVLVSSEELKYLGLMPEQQSVYDET	98
NTDB id 1384 A4U84 RS08525 WP 010787006.1	SRWQKNCEEKQLLRLRYLAVPVQETEHKLWLAVD DENNLTA CEIFAFMTHKQIEPVVIASDELKYLNLNALSPEQQPIYEE	99
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQLGLRELITRYDALPIAKQGNL LLLAVSDPTLLQAEEBFRFATGLQVELALADHRAALQAAIRRLYGRSIQGAANQ	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLADFRELTAAIRRLYGRSLGQEKSG	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQLGLRELITRHNALPLHRTPTSTLLAVADPTNQAEDDFRFATGLQVELVLADFRELSTAIRRLYGRSLSHEKSG	149
NTDB id 1251 GC085 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDPPSKQASLKEITQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSE	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLILKHRILPLIQRGQILYVATS NPSNIEAIDAIRFNSKLLVEPVIVEHKKLEKVLGQHF AEESFFDND	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATS NPTNIEAMDAIRFNSKLLKVEPIIVEHDKLERLLSEHFAVEETHFNFD	157
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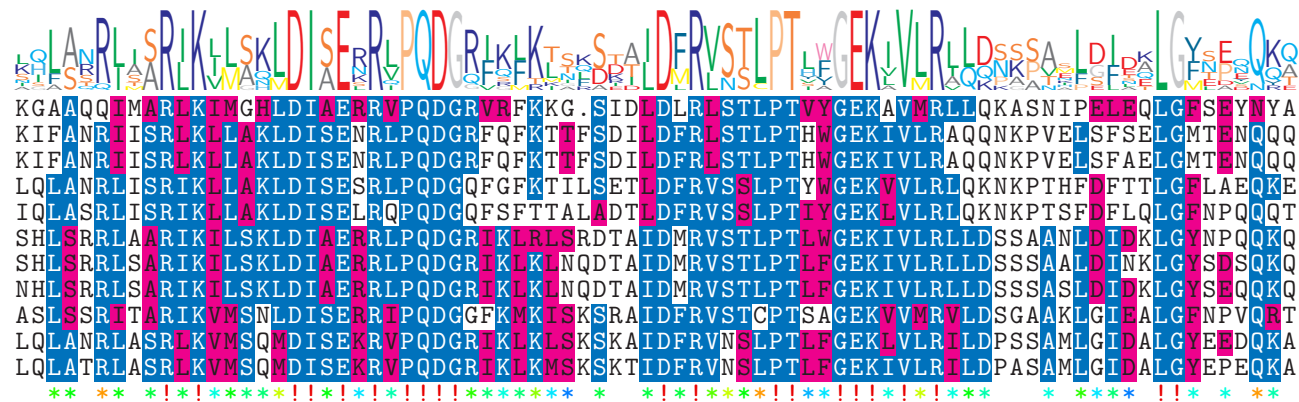
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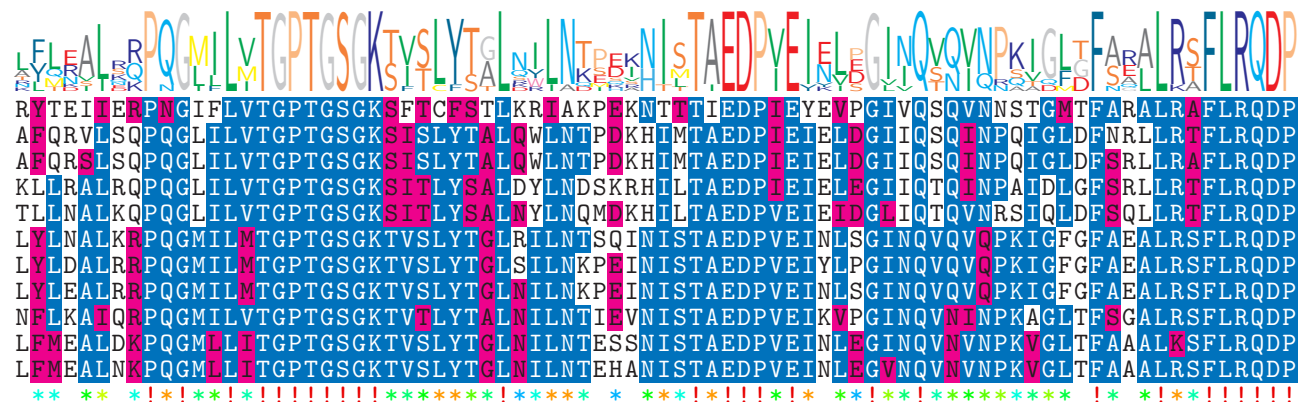
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