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NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
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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
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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGASDA	160
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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	320
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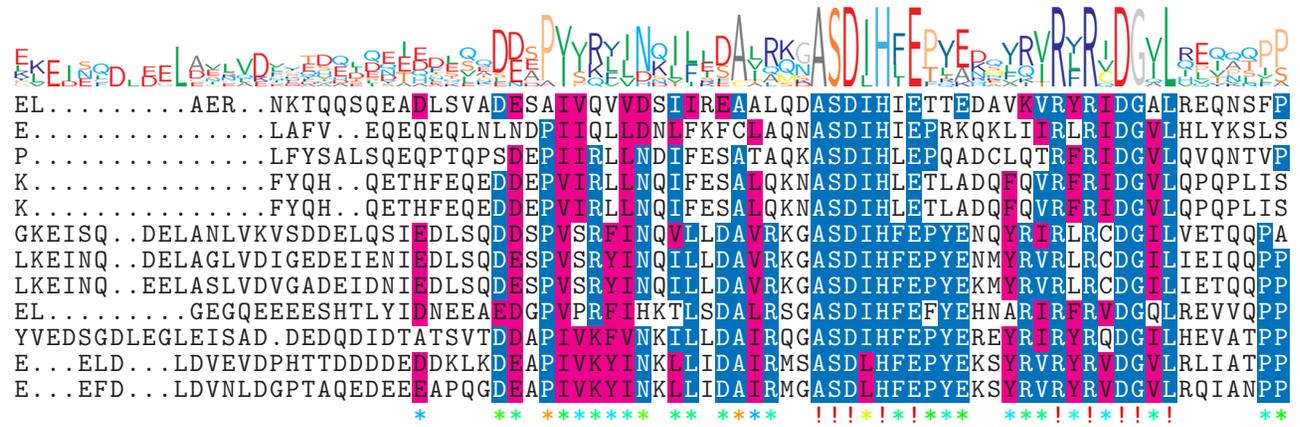
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NTDB id 1110 NGFG RS09235 WP 003689817.1MSVGLLRILLVQNQVVTVEQAEHYY.NESQAGKEVLPMLFSDGVISPKSLAALIAARVFSYSILDLRHY	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQNFVGPMLDINCID	77
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NTDB id 158613 ACT75 RS07360 WP 005540457.1	ELWLKNNQQQQTLLRYLALPLAENGQTLWLGGLDSLNLISACEAFSFLTGKTIIEPVVIESQTLKLTALQDLSR	103
NTDB id 1366 NTHI RS01950 WP 011271972.1	DLWERNQQQQSLLIRYFALPLKEENRNLWLGVDLSNLISACETIAFITGKPVIEPILLESSQLKELLQQLTPHQ	104
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLAADFRELTAAIRRLYGR	149
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NTDB id 1110 NGFG RS09235 WP 003689817.1	HRVLMGVLTEEQMVFEHCVVFRRGDKVFFAVSDPTQMPQIQKTVSAAGIAVELVIVEDDQLAGLLDVGSRST	146
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSKLLKVEPIIVEHDKLERLLSEHFVE	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLLIKHRIPLIQRGQILYVATSNPSNIEAIDAIRFNSKLLVPEVIVEHKKLEKVLGQHF	157
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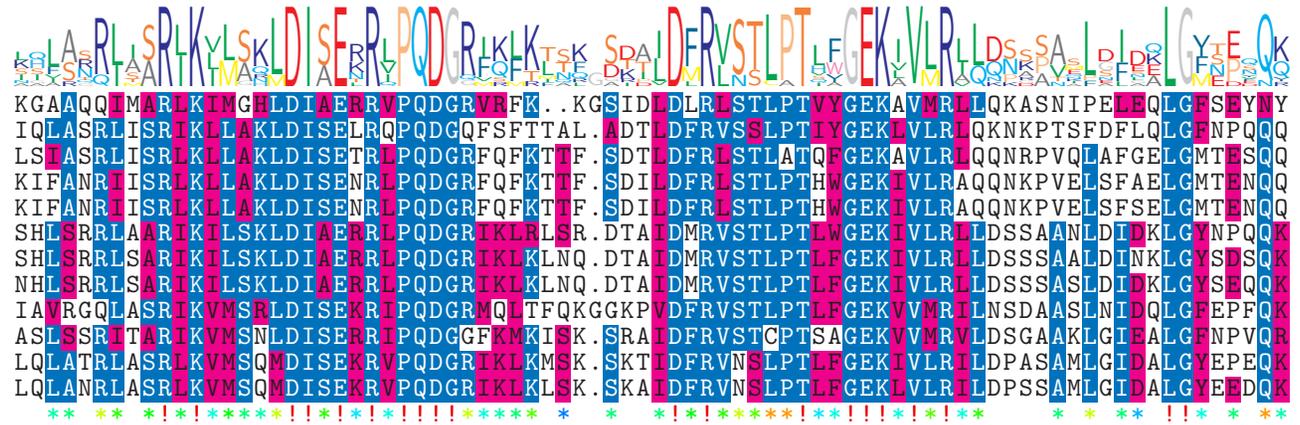
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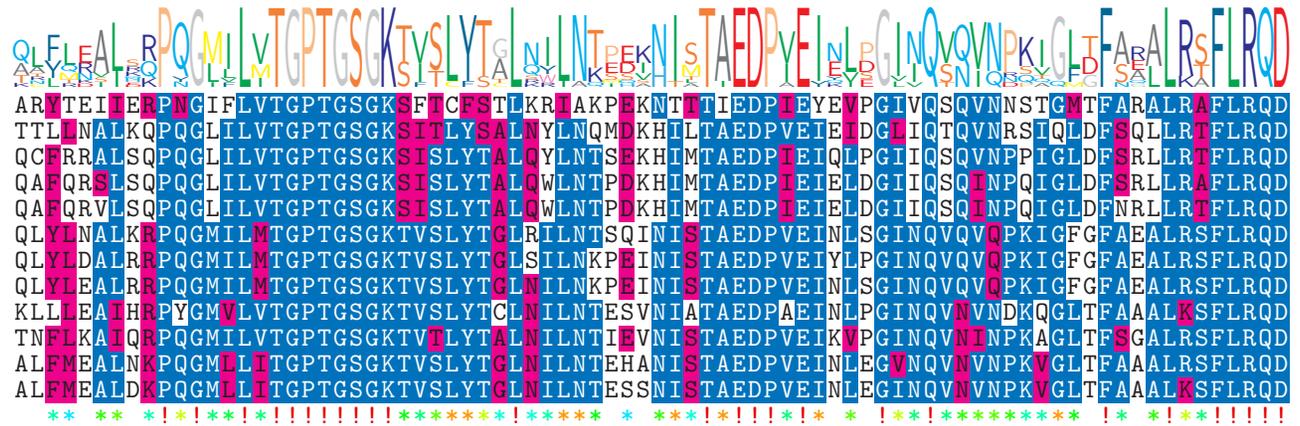
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NTDB id 1306 DR RS10060 WP 027479822.1
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