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NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAALR	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 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	0
NTDB id 166729 ATY43 RS03455 WP 011408072.1	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	SIRPQTALNLQAFPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEIVVPSGASDA	160
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NTDB id 166729 ATY43 RS03455 WP 011408072.1	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	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPNSGFHPGEEVLGSM	240
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus		

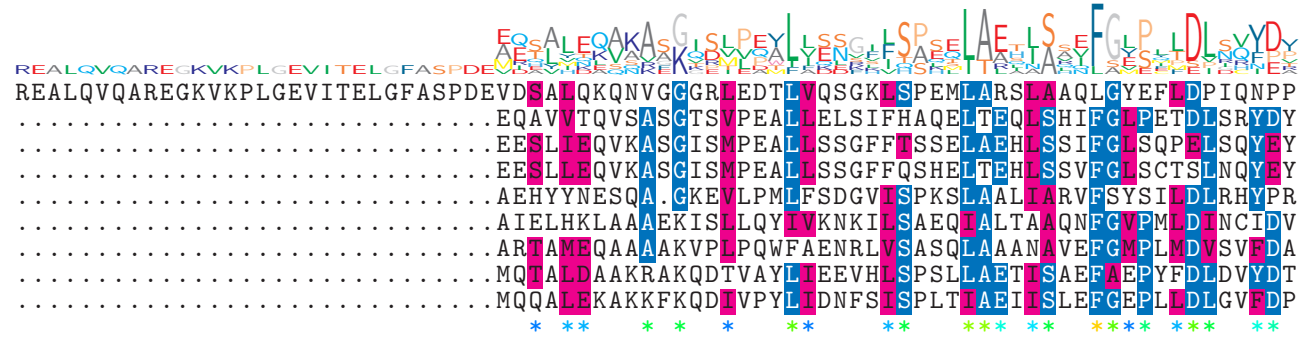
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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRDPIDALIGRPVQLMLTRPRDIERLIEQFYPQGRGLGEQLVQEGTLSRDQL	320
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NTDB id 1110 NGFG RS09235 WP 003689817.1MSVGLLRITLVQNQVVTVEQ.	19
NTDB id 1251 GC085 RS07725 WP 011946523.1MALATEEYRLQGTGQLLVLEKLLDKTK.	27
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NTDB id 1018 ACIAD RS01685 WP 004920473.1MS..AFTTPPKFSGFIRRLVEEGYVNAQN.	27
consensus		

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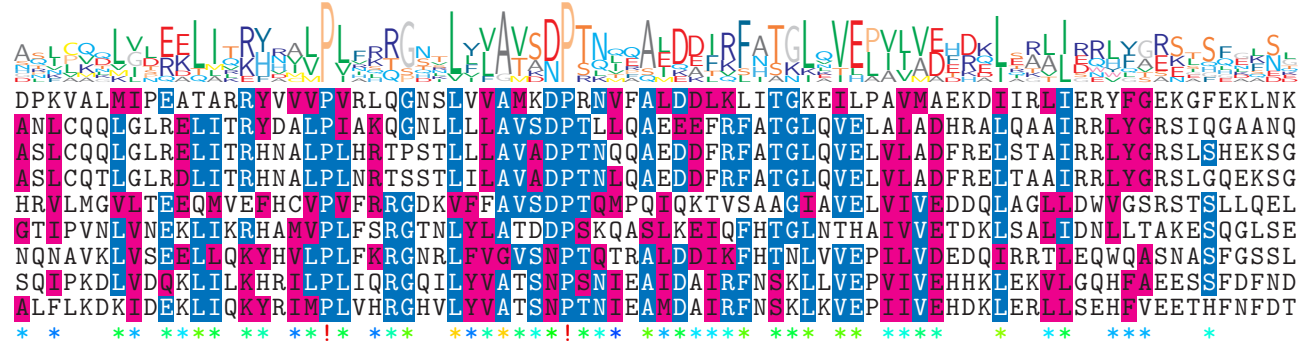
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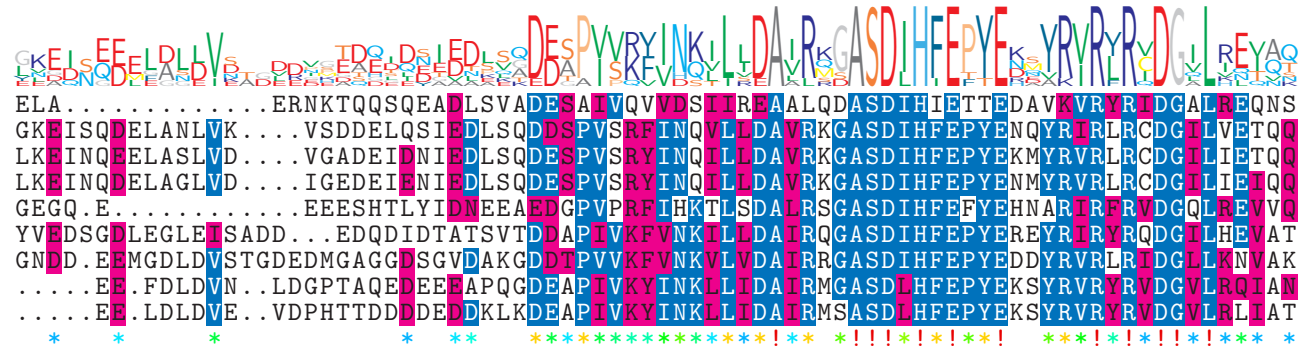
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NTDB id 1306 DR RS10060 WP 027479822.1
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625
304
304
304
295
313
317
308
308

 non conserved
 similar
 $\geq 50\%$ conserved