

logo

	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	
NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 328060 EGH15 RS09995 WP 003689817.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

logo

	SIRPQTALNLQAFPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	
NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
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NTDB id 328060 EGH15 RS09995 WP 003689817.1	0
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consensus		

logo

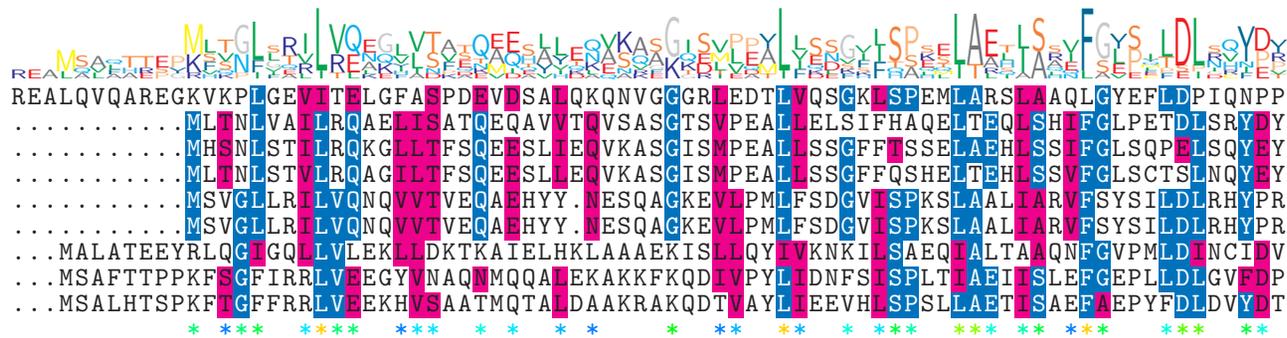
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

logo

	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	
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consensus		

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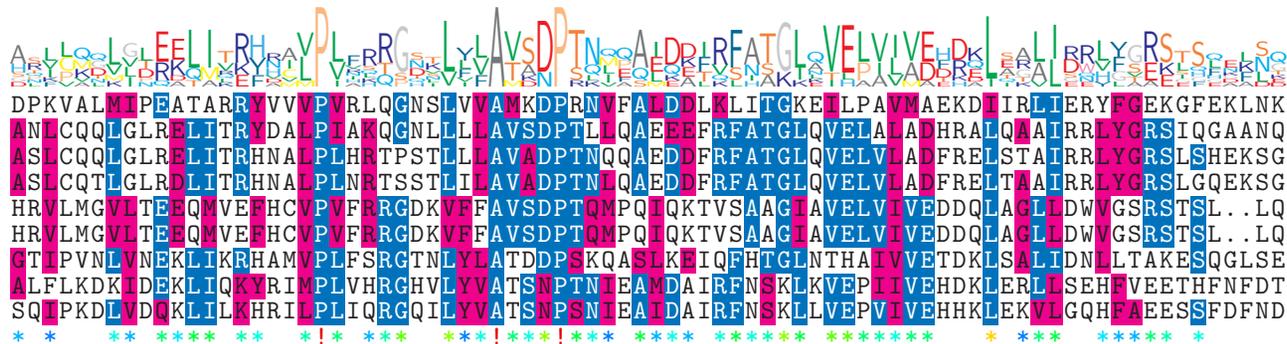
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



400
69
69
69
68
68
77
77
77

logo

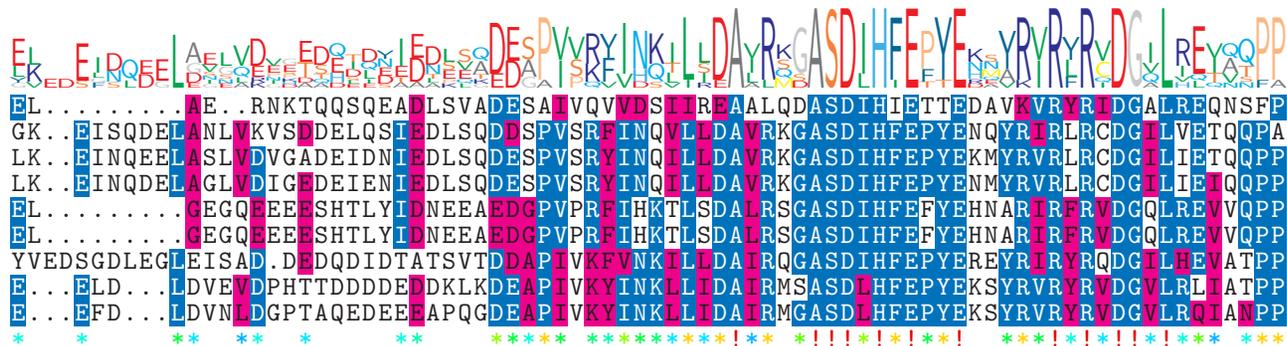
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
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 consensus



480
149
149
149
146
146
157
157
157

logo

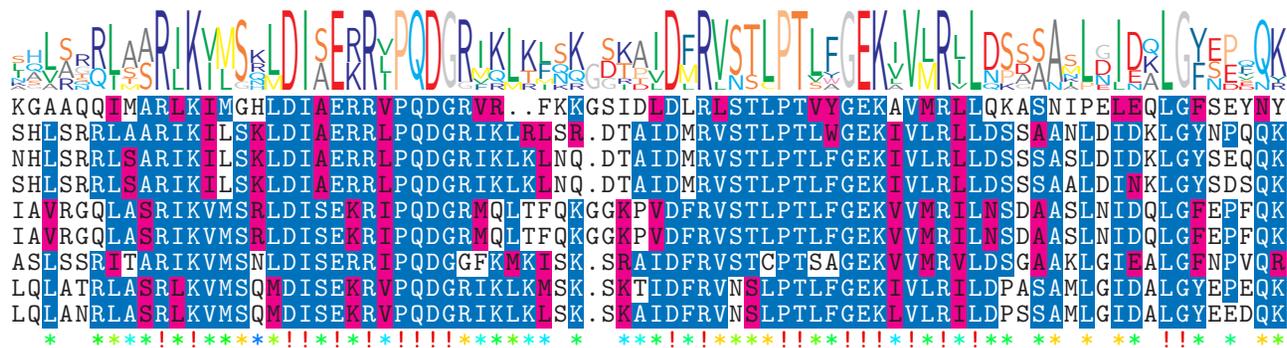
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



549
227
227
227
217
217
236
231
231

logo

NTDB id 1306 DR RS10060 WP 027479822.1
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



627
306
306
306
297
297
315
310
310

 non conserved
 similar
 $\geq 50\%$ conserved