

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	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 332344 ELB75 RS12220 WP 126984126.1	.....	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
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEIVVPSGASDA	160
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 332344 ELB75 RS12220 WP 126984126.1	.....	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	.....	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
consensus		

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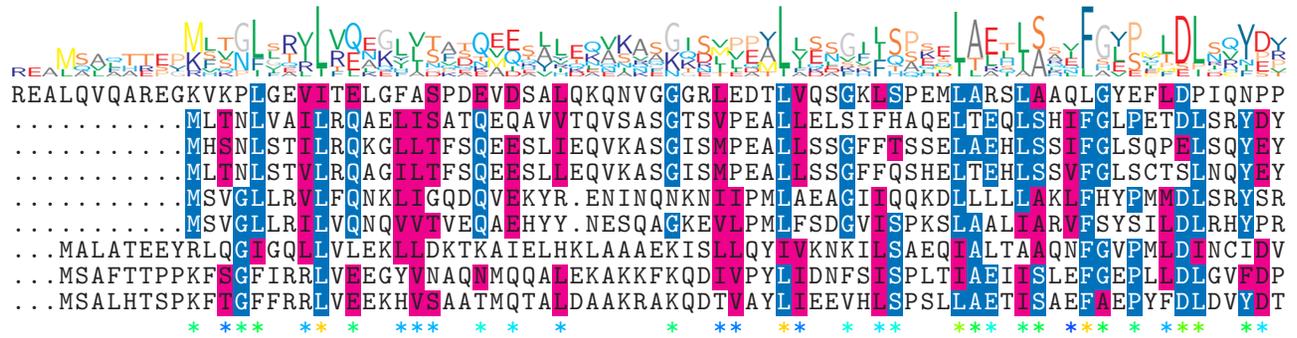
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 332344 ELB75 RS12220 WP 126984126.1	.....	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	320
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
consensus		

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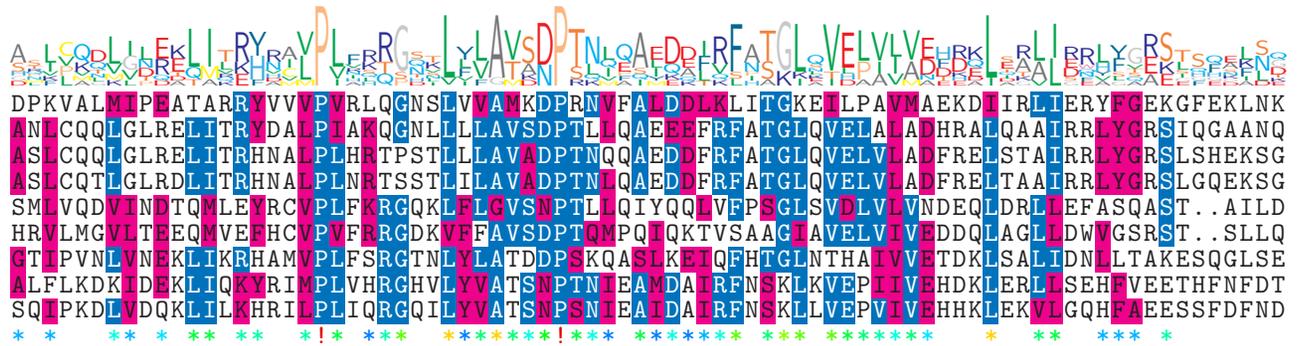
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 NTDB id 332344 ELB75 RS12220 WP 126984126.1  
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



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logo

NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
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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

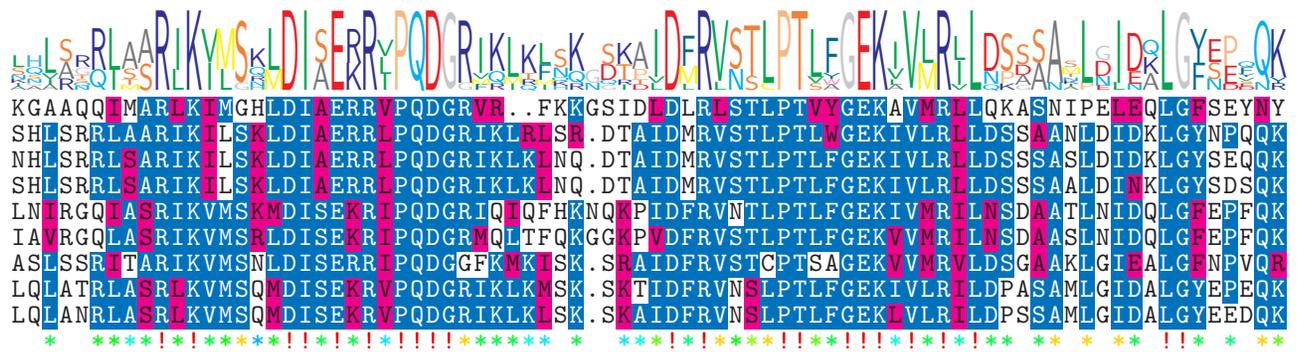
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



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logo

NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



627  
306  
306  
306  
293  
297  
315  
310  
310



 non conserved  
 similar  
  $\geq 50\%$  conserved