

logo

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 263013 COL90 RS16990 WP 024743351.1	.....	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 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 1018 ACIAD RS01685 WP 004920473.1	.....	0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRGLGEQLVQEGTISRDL	320
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....MLTNLVAITLRQAELISATQ.	19
NTDB id 1293 VP RS12240 WP 005479695.1	.....MHSNLSTILRQKGLTFSQ.	19
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....MLTNLSTVLRQACILTFSQ.	19
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....MSVGLLRILVQNQVVTVEQ.	19
NTDB id 1251 GC085 RS07725 WP 011946523.1	.....MALATEEYRLQGTGQLLVLEKLLDKTK.	27
NTDB id 263013 COL90 RS16990 WP 024743351.1	.....MNSVSTTNLVGITGIARRLVQDGALEEEA.	29
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....MS..ALHTSPKFTGFFRRLVEEKHVSAAAT.	27
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....MS..AFTTPPKFSGFIRRLVEEGYVNAQN.	27
consensus		

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 consensus



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 consensus



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 consensus



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NTDB id 1306 DR RS10060 WP 027479822.1  
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 consensus



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logo

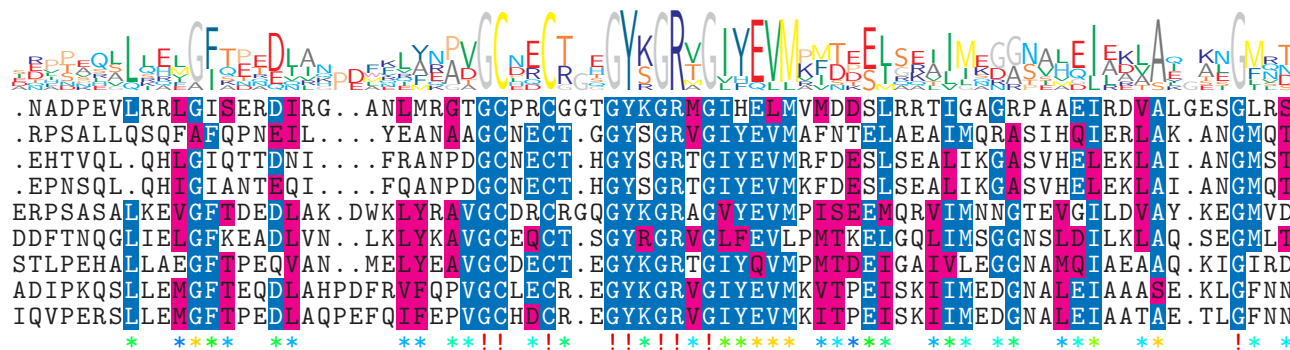
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1



785  
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logo

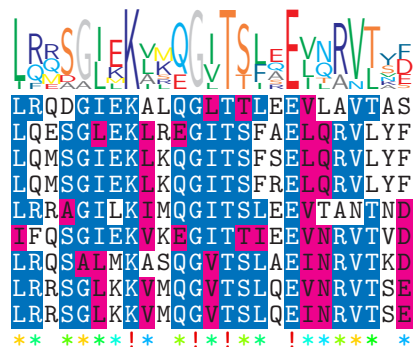
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570

 non conserved  
 similar  
  $\geq 50\%$  conserved