

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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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	SIRPQTALNLQAFPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIIVVPSGASDA	160
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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	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSCGFHPGEEVLGSM	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	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRIGEQLVQECTLSRDQL	320
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MLINLVALRQAEELISATQ.	19
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSQ.	19
NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVGLRLVLQNQVITVEQ.	19
NTDB id 1251 GCO85 RS07725 WP 011946523.1	MALATEEYRLQCTGQLLVLEKLLDKTK.	27
NTDB id 1101908 Q9265 RS07750 WP 011408072.1	MNSVSTTNLVGITCIARRLVQDGALEETA.	29
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MS..ALHTSPKFTGFFRRLVEEKHVSAAAT.	27
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MS..AFTTPPKFSCFIRRLVEEGYVNAQN.	27
consensus	*** *!***** *	

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NTDB id 1018 ACIAD RS01685 WP 004920473.1	MQQALEKAKKFQKDVPVYLIDNFS	77
consensus		

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NTDB id 1101908 Q9265 RS07750 WP 011408072.1	NQNAVKLVSEELIQKYHVLPLFKRCNRL	159
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLIKLRHRLPLLIQRGQIL	157
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consensus		

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consensus		

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



705
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384
384
375
393
397
388
388

logo

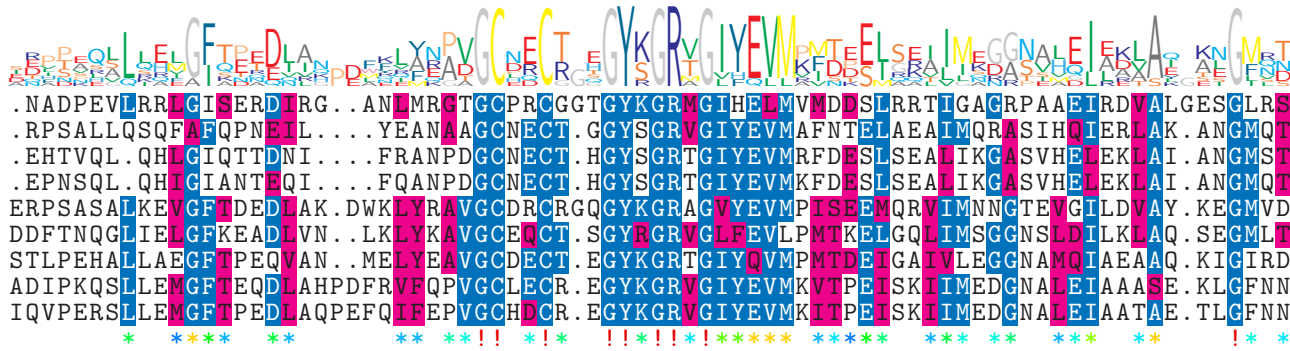
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus



785
464
464
464
455
473
476
467
467

logo

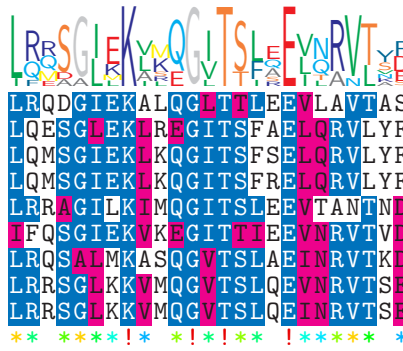
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consensus



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533
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545
545

logo

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consensus



887
562
561
561
558
574
577
570
570

 non conserved
 similar
 $\geq 50\%$ conserved