

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1169735 ACH8EK RS10295 WP 003689817.1	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	0

logo

NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
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consensus	0

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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEGTTFLPNPSGFHPGEEVLGSM	240
NTDB id 1168 A1552VC RS11070 WP 000957200.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEGTTFLPNPSGFHPGEEVLGSM	0
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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	0

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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDI DALIGRPVQLMLTRPRDIERLIEQFYFQRGRLGEQLVQEGTLSRDQL	320
NTDB id 1168 A1552VC RS11070 WP 000957200.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDI DALIGRPVQLMLTRPRDIERLIEQFYFQRGRLGEQLVQEGTLSRDQL	0
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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	0

logo

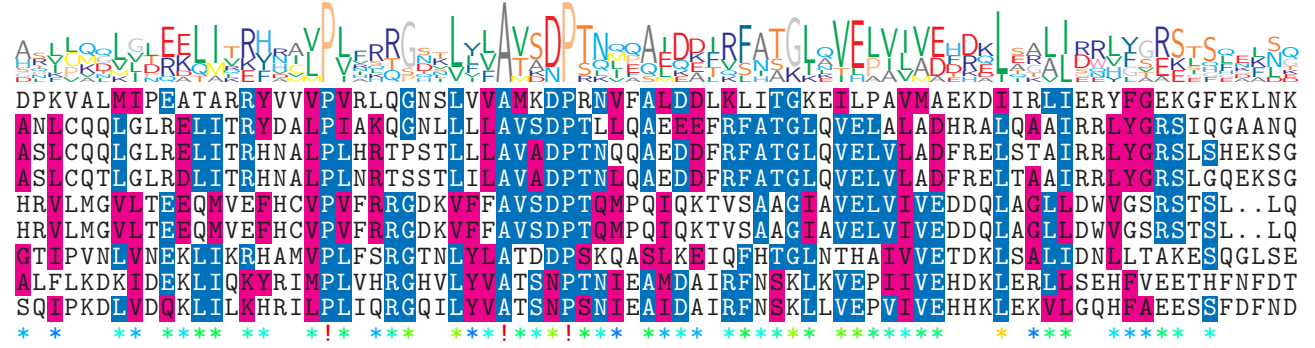
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 consensus



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logo

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
 consensus



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logo

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 consensus



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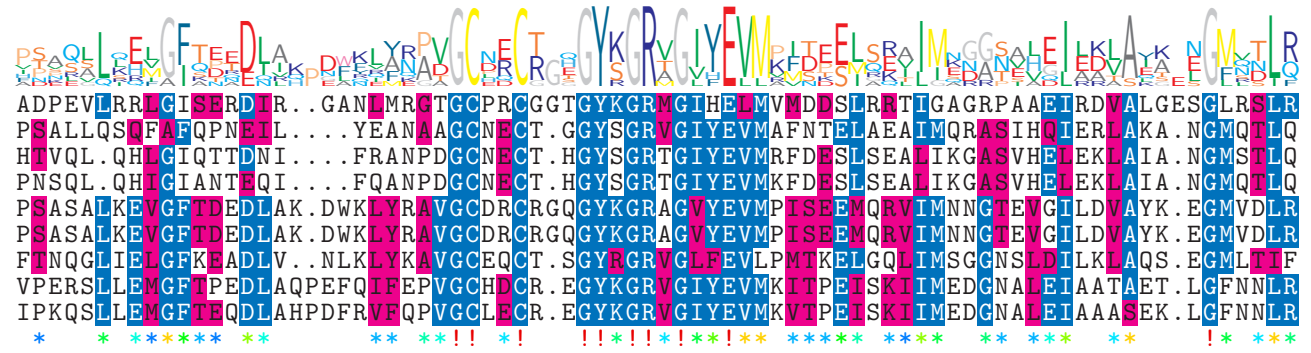
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
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786
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469

logo

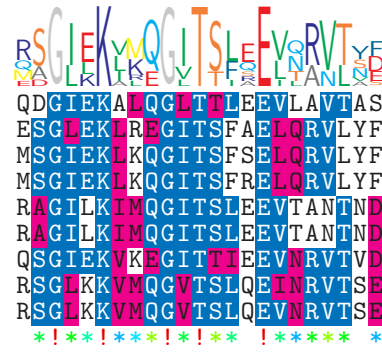
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
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558
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574
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570

 non conserved
 similar
 $\geq 50\%$ conserved