

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	80
NTDB id 1110 NGFG RS09235 WP 003689817.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1293 VP RS12240 WP 005479695.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1006212 ABHN08 RS19460 WP 073475466.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0
consensus	MSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLI	0

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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
NTDB id 1110 NGFG RS09235 WP 003689817.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1293 VP RS12240 WP 005479695.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1006212 ABHN08 RS19460 WP 073475466.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
consensus	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0

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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	240
NTDB id 1110 NGFG RS09235 WP 003689817.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0
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NTDB id 1293 VP RS12240 WP 005479695.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0
NTDB id 1006212 ABHN08 RS19460 WP 073475466.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0
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consensus	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVLGSM	0

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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	320
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	19
NTDB id 1293 VP RS12240 WP 005479695.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	19
NTDB id 1401 DSB67 RS12665 WP 010643259.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	19
NTDB id 1251 GC085 RS07725 WP 011946523.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	27
NTDB id 1006212 ABHN08 RS19460 WP 073475466.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	23
NTDB id 1058 ABD1 RS01615 WP 001274986.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	27
NTDB id 1018 ACIAD RS01685 WP 004920473.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	27
consensus	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	27

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consensus



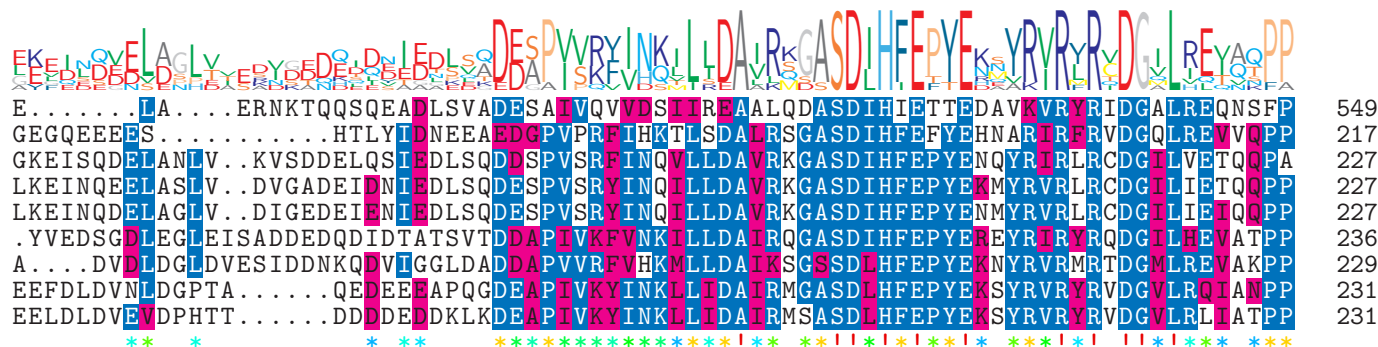
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



 non conserved
 similar
 $\geq 50\%$ conserved