

logo

NTDB id 1306 DR RS10060 WP 027479822.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240885 CHR53 RS19715 WP 066385505.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAALR
MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAALR
.....
.....
.....

80
0
0
0

logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 240885 CHR53 RS19715 WP 066385505.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEWPSGASDA
SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEWPSGASDA
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.....
.....

160
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

GRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM
GRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM
.....
.....
.....

240
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTL SRDQL
LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTL SRDQL
.....
.....
.....

320
0
0
0

logo

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consensus

REALQVQAREGKVKPLGEVITELGFASPDEVDSALQKQNVGGGRLLEDTLVQSGKLSPELMSOYTLCEQOTYRLLDPIQNSP
REALQVQAREGKVKPLGEVITELGFASPDEVDSALQKQNVGGGRLLEDTLVQSGKLSPELMSOYTLCEQOTYRLLDPIQNSP
.....
.....
.....

400
19
0
0

* ** * *

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consensus

SRWYKMLPKOTLRVYVVPVRLTENS LWYAVKDRNNTALELHKMTHKQIPYVMA SKELKYLINRYSP EQQP
DPKVALMIPEATARRVYVVPVRLTENS LWYAVKDRNNTALELHKMTHKQIPYVMA SKELKYLINRYSP EQQP
SRWQKNCEEKQLLLRYLAVPVQETEHKLLWLAVDENNLTACEIFAFMTHKQIEPVVIASDELKYLINNALSP EQQP
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.....

480
94
0
0

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 non conserved
 similar
 $\geq 50\%$ conserved