

logo

NTDB id 1306 DR RS10060 WP 027479822.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 320607 D9842 RS12500 WP 121662826.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR
MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR
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.....

80
0
0
0

logo

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NTDB id 320607 D9842 RS12500 WP 121662826.1
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
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.....
.....

240
0
0
0

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consensus

LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL
LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL
.....
.....
.....

320
0
0
0

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consensus

REALQVQAREGKVKPLGEVITELGFASPDEVDSALQKQNVGGGRLLEDTLVQSGKLSPELMSQYITICEQQTYRIFDPIQNSP
REALQVQAREGKVKPLGEVITELGFASPDEVDSALQKQNVGGGRLLEDTLVQSGKLSPELMSQYITICEQQTYRIFDPIQNSP
.....
.....
.....

400
19
0
0

* ** * *

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consensus

SRWYKMLPKOTLRVYVVPVRLTENS L WYAVK DERNYTALEIKMTHKQIPYVMA SKELKYL NRYSP EOPLYNS
DPKVALMIPEATARRVYVVPVRLTENS L WYAVK DERNYTALEIKMTHKQIPYVMA SKELKYL NRYSP EOPLYNS
SRWQKNCEEKQLLLRYLAVPVQETEHKLLWLAVDENNLTACEIFAFMTHKQIEPVVIASDELKYL NALSPEQQPIYEES
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.....

480
99
0
0

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 non conserved
 similar
 $\geq 50\%$ conserved