

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 410881 BABF1 RS20950 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHLTELSSVFGLSCTSLNQYEYASLCQTLGLRD
MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHLTELSSVFGLSCTSLNQYEYASLCQTLGLRD
.....
.....

80
0
0

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 410881 BABF1 RS20950 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLADFRELTAAIRRLYGRSLGQEKSGLKEINQDEL
LITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLADFRELTAAIRRLYGRSLGQEKSGLKEINQDEL
.....
.....

160
0
0

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 410881 BABF1 RS20950 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LVDIGEDEIENIEDLSQDESPVSRVINQILLDAVRKQASDIHFEPYENMYRVRLRCDGILTEIQPPSHLSRRLSARIKI
LVDIGEDEIENIEDLSQDESPVSRVINQILLDAVRKQASDIHFEPYENMYRVRLRCDGILTEIQPPSHLSRRLSARIKI
.....
.....
.....

240
62
62

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 410881 BABF1 RS20950 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

LSMDIGERRKPNQGSLLYKLNQQNVYLWVSTLPTVYQESLVRLLMPQYSIQSSNKLSFPSTKQTLSELRSHGML
LSMDIGERRKPNQGSLLYKLNQQNVYLWVSTLPTVYQESLVRLLMPQYSIQSSNKLSFPSTKQTLSELRSHGML
.....
.....
.....

320
140
140

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 410881 BABF1 RS20950 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

YFTGPTGSGKTTLLYSYLRKPTNRNVTLEDPEVETRNPDVLQVQNEKAGYTYSAGLKAILRHDPDVIWVGEIRD
YFTGPTGSGKTTLLYSYLRKPTNRNVTLEDPEVETRNPDVLQVQNEKAGYTYSAGLKAILRHDPDVIWVGEIRD
.....
.....
.....

399
220
220

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 410881 BABF1 RS20950 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

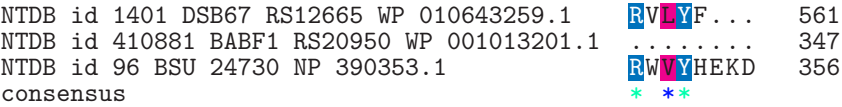
TAEIAIRASLTGHLVLTHTNDARGAIYRLSNMGVTSQIEQSLAIAAQRLLVLLCPFCRRPCCSTYVRSRQVRR
TAEIAIRASLTGHLVLTHTNDARGAIYRLSNMGVTSQIEQSLAIAAQRLLVLLCPFCRRPCCSTYVRSRQVRR
.....
.....
.....

476
299
300

logo



logo



non conserved
 similar
 ≥ 50% conserved