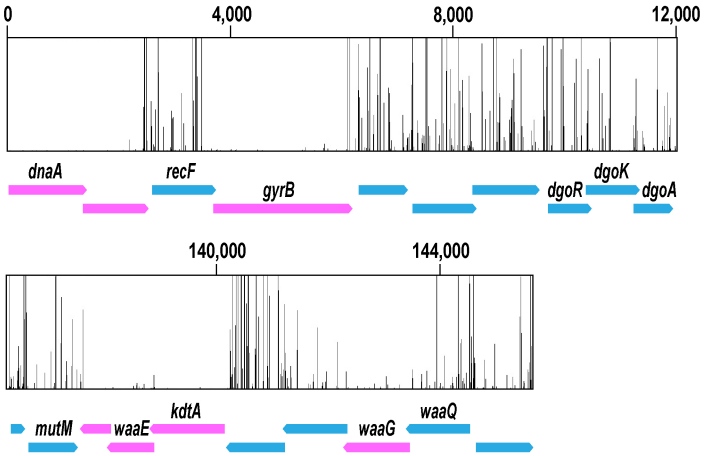
Figure S2


0
4,000
8,000
12,000
dnaA
recF
gyrB
dgoR
dgoK
dgoA
140,000
144,000
mutM
waaE
kdtA
waaG
waaQ