Table S3. Genes identified in this work as having a general role in antibiotic susceptibility.

bnum	Annotation	References
b3791	rffA dTDP-4-oxo-6-deoxy-D-glucose transaminase	[1]
b3792	wzxE lipid III flippase	
b1904	yecR predicted protein	
b2616	recN protein used in recombination and DNA repair	[1,2,3,4]
b2494	yfgC predicted peptidase	[1,5]
b0462	acrB RND-type permease	[1,6,7]
b1130	phoP PhoP-Phosphorylated transcriptional dual regulator	[1]
b4396	rob Rob transcriptional activator	[8]
b3194	yrbE YrbF/YrbE ABC transporter	
b3191	yrbB predicted NTP-binding protein	
b3632	rfaQ lipopolysaccharide core biosynthesis protein; heptosyl transferase	[9]
b3630	rfaP lipopolysaccharide core biosynthesis; phosphorylation of core heptose	[9]
	rfaB UDP-D-galactose:(glucosyl)lipopolysaccharide-1,6-D-	
b3628	galactosyltransferase	[9]
b3631	rfaG lipopolysaccharide glucosyltransferase I	[9]
b2215	ompC outer membrane porin C	
b3780	rhIB degradosome	
b3956	ppc phosphoenolpyruvate carboxylase	
b3493	pitA PitA	
b2415	ptsH HPr	
b3806	cyaA adenylate cyclase	[10]
b0015	dnaJ chaperone with DnaK; heat shock protein	[11]
b1048	mdoG periplasmic glucan (MDO) biosynthesis protein	
b1049	<i>mdoH</i> membrane glycosyltransferase; synthesis of membrane-derived oligosaccharide (MDO)	
b2218	rcsC RcsC-P his	[12]
b1291	sapD peptide uptake ABC transporter	
b3290	trkA NAD-binding component of TrK potassium transporter	
b1290	sapF peptide uptake ABC transporter	
b3849	trkH TrkH potassium ion Trk Transporter	
b0464	acrR AcrR transcriptional repressor	[1,13]
b1676	pykF pyruvate kinase I monomer	<u> </u>

In finding references, emphasis was placed on global studies and work in *E. coli*. Expression changes in response to drug addition were not sufficient for inclusion. Results and Discussion describes how the class of genes was determined.

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