

Table S1. Primers used in this study.

Method and target gene	Primer	Sequence (5'–3')
Mutagenesis and complementations		
<i>A. baumannii</i>		
<i>lpxO</i>	Ab_lpxO_UP_F1	GTA CAG CTC CGG TAC ACA TTT A
	Ab_lpxO_DWN_F1(BamHI)	¹ <u>GGA TCC</u> TGG TTC CTC ATG CTT GGA AT
	Ab_lpxO_UP_R1(BamHI)	¹ <u>GGA TCC</u> TAG GGT TGA TTC GGT ACC TTT G
	Ab_lpxO_DWN_R1	ATA GGA AAC TAG CCC GAC AAA G
	KpnLpxL1ComF	GCC GGT TAA AAA GCA GCC AA
	KpnLpxL1ComR	AGG AAA TAA ACG GCG GGC AT
Tn7 insertion	Ab_glmS_F1 Tn7_Right	TTT GCT GAT GAA AAT AGC GG CAC AGC ATA ACT GGA CTG ATT TC
<i>E. coli</i>		
Tn7 insertion	E.coli_glmS_Up E.coli_glmS_Down EcoliLpxL_check_F	TCG ACT GGG CGT ACA AAA CC CGG GAA ACC ATA CCG GAG TT GCA CAC TAA TTA TGC GCC CG
	E.coli_pTN7L_F1 E.coli_pTN7R_R1	ATT AGC TTA CGA CGC TAC ACC C CAC AGC ATA ACT GGA CTG ATT TC
qPCR		
<i>hprt</i>	mHPRT-F1	GAT CAG TCA ACG GGG GAC AT
	mHPRT-R1	GGT CCT TTT CAC CAG CAA GC
<i>tnfα</i>	mTNF-α-F1	TTC TGT CTA CTG AAC TTC GGG GTG ATC GGT CC
	mTNF-α-R1	GTA TGA GAT AGC AAA TCG GCT GAC GGT GTG GG
<i>il10</i>	mIL-10_F1	GGA CTT TAA GGG TTA CTT GGG TTG CC
	mIL-10_R1	CAT GTA TGC TTC TAT GCA GTT GAT GA
<i>creb</i>	mCREB1_F1	TAA CAG TGC CAA CCC CCA TTT
	mCREB1_R1	CCT GTA CCC CAT CCG TAC CA
<i>18S</i>	Galleria_18S_F1	ATG GTT GCA AAG CTG AAA CT
	Galleria_18S_R1	TCC CGT GTT GAG TCA AAT TA
<i>lysozyme</i>	Gall_Lysozyme_F1	TCC CAA CTC TTG ACC GAC GA
	Gall_Lysozyme_R1	AGT GGT TGC GCC ATC CAT AC
<i>galiomycin</i>	Galiomycin_F1	CCT CTG ATT GCA ATG CTG AGT G
	Galiomycin_R1	GCT GCC AAG TTA GTC AAC AGG
<i>gallerimycin</i>	Gallerimycin_F1	CGC AAT ATC ATT GGC CTT CT
	Gallerimycin_R1	CCT GCA GTT AGC AAT GCA CTC

¹ BamHI site underlined.