

**Table S3. Bacterial strains and plasmids used in this study**

Strain/Plasmid	Genotype/Resistance/Use <sup>a</sup>	Source
<b>Strains</b>		
<i>A. baumannii</i> strain 17978	wild-type	ATCC
<i>fepA</i> ( <i>AIS_0980</i> )	17978 $\Delta fepA::kan^R$	This study
<i>fepA+</i>	<i>fepA</i> with PABBR_MCS_ <i>fepA AIS_0981</i>	This study
<i>feoB</i> ( <i>AIS_0653</i> )	17978 $\Delta feoB::kan^R$	This study
<i>feoB+</i>	<i>feoB</i> with PABBR_MCS_ <i>AIS_0652_feoB</i>	This study
<i>ddc</i> ( <i>AIS_0568</i> )	17978 $\Delta ddc::kan^R$	This study
<i>ddc+</i>	<i>ddc</i> with PABBR_MCS_ <i>ddc</i>	This study
<i>pntB</i> ( <i>AIS_02435</i> )	17978 $\Delta pntB::kan^R$	This study
<i>pntB+</i>	<i>pntB</i> with PABBR_MCS_ <i>pntB</i>	This study
<i>Escherichia coli</i> EZTn5	Source of kanamycin cassette	S. Subashchandrabose
<b>Plasmids</b>		
pUC19	amp <sup>R</sup> , cloning mutant construct	Life Technologies
pAT02	Rec <sub>Ab</sub> system, recombineering	13
pKD4	kan <sup>R</sup> , source of kanamycin cassette	31
PABBR_MCS	amp <sup>R</sup> , cloning for complementation	13
PABBR_MCS_ <i>fepA AIS_0981</i>	amp <sup>R</sup> , complementation	This study
PABBR_MCS_ <i>feoB AIS_0652</i>	amp <sup>R</sup> , complementation	This study
PABBR_MCS_ <i>ddc</i>	amp <sup>R</sup> , complementation	This study
PABBR_MCS_ <i>pntB</i>	amp <sup>R</sup> , complementation	This study

<sup>a</sup>kan, kanamycin; amp, ampicillin/carbenicillin, R, resistant