

**S11 Table. Primer Sequences.**

Primer Name	Sequence
pBAD_F	GAATTCGAAGCTTGGCTGT
pBAD_R	GGTTAATTCCTCCTGTTAGCCC
ArmA_F	GGCTAACAGGAGGAATTAACCATGGA CAAGAATGACGTGGTCAAAAAA
ArmA_R	ACAGCCAAGCTTCGAATTCTTACTTTTGGAAGCCCGAAGT
NpmA_F	GGCTAACAGGAGGAATTAACCATGCT GATTCTGAAAGGAACTAAAACGG
NpmA_R	ACAGCCAAGCTTCGAATTCTTAGTGTTTCGATACATGACCGG
T7Nluc-F	GACGTTGTAAAACGA
pBAD-tetM_F	TGTTCAATAAAATAACTTAGgaattcgaagcttgctgtttgg
pBAD tetM_R	CCAATATTAATAATTTTCATggtaattcctcctgtagccC
tetM_F	gctaacaggaggaattaaccATGAAAATTATTAATATTGGAGTTTTAGCTCAT GTTGATGCAGG
tetM_R	aacagccaagcttgaattcCTAAGTTATTTTATTGAACATATATCTTACTTTA TCTATCCGACT
T7Nluc-R	AGGAAGCCCGGATAT
NLuc_BamHI-F	CGC <u>GGATCC</u> GTCTTCACACTCGAAGAT
NLuc_PstI-R	CGC <u>CTGCAG</u> TTACGCCAGAATGCGTTC
F16S+23S	CACCTCGCGACAGTGCCTAAAGCG
R16S+23S	GGTCAGGTGGGACCACCGCGCTAC
16SR*	CGCAAGACGCCTTGCTTTTCA

\*16S rRNA *rrsE* reverse primer from Orelle C, et al. (1).

## Reference

1. Orelle C, *et al.* (2013) Tools for characterizing bacterial protein synthesis inhibitors. *Antimicrob Agents Chemother* 57(12):5994-6004.