

**S1 Table. Quantitative data of CLas plasmid DNA with 16S primer in qPCR assay**

Calculated copies/ $\mu$ l <sup>a</sup>	Log SQ <sup>b</sup>	Primer/Probe concentration											
		50/100				100/200				150/300			
		Singleplex		Duplex		Singleplex		Duplex		Singleplex		Duplex	
		Mean Cq	SD <sup>c</sup>	Mean Cq	SD <sup>c</sup>	Mean Cq	SD <sup>c</sup>	Mean Cq	SD <sup>c</sup>	Mean Cq	SD <sup>c</sup>	Mean Cq	SD <sup>c</sup>
1.48E+05	5.169	19.23	0.090	19.28	0.095	19.81	0.075	20.01	0.111	19.79	0.025	20.16	0.023
1.48E+04	4.169	22.54	0.197	22.74	0.235	23.15	0.146	23.24	0.147	23.11	0.042	23.67	0.044
1.48E+03	3.169	25.73	0.175	25.86	0.121	26.40	0.112	26.40	0.153	26.42	0.044	26.57	0.471
1.48E+02	2.169	29.14	0.300	29.09	0.136	31.05	0.526	30.64	0.265	29.88	0.14	30.83	0.276
1.48E+01	1.169	33.89	0.729	33.57	0.700	33.10	0.485	33.20	0.140	34.33	0.124	35.47	0.375
1.48E+00	0.169	35.55	0.840	36.72	0.163	36.45	0.352	36.55	0.092	35.94	0.313	36.49	0.417
NTC <sup>d</sup>	0	0	0	0	0	0	0	0	0	0	0	0	0

<sup>a</sup> Values reflect copies/ $\mu$ l of calculated serial dilutions of positive plasmid DNA standard.

<sup>b</sup> SQ means starting quantity. Data represent the mean of each dilution tested in triplicate.

<sup>c</sup> SD means standard deviation.

<sup>d</sup> NTC means no template control.