

1 **Table S1** Percent infection rate of “*Ca. L. asiaticus*” and *Wolbachia* and mean copy number of
 2 non-exposed *Diaphorina citri* using qPCR in both nymphs and adults, including individual guts
 3 and whole bodies¹

Non-Exposed					
	Stage	Bacteria	Ct < 35 (%)	Ct > 35 (%)	Mean Copy # ²
Whole Body ¹	Nymph (5 th)	" <i>Ca. L. asiaticus</i> "	0.0	100.0	0.0
		<i>Wolbachia</i>	100.0	0.0	110062.8
	Adult	" <i>Ca. L. asiaticus</i> "	0.0	100.0	0.0
		<i>Wolbachia</i>	100.0	0.0	201125.9
Gut ³	Nymph (5 th)	" <i>Ca. L. asiaticus</i> "	0.0	100.0	0.0
		<i>Wolbachia</i>	100.0	0.0	64300.1
	Adult	" <i>Ca. L. asiaticus</i> "	0.0	100.0	0.0
		<i>Wolbachia</i>	100.0	0.0	1506884.8

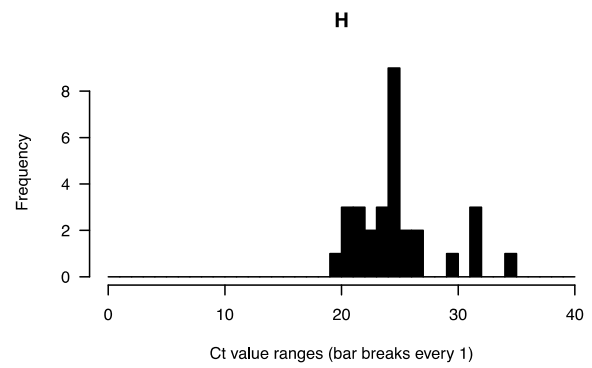
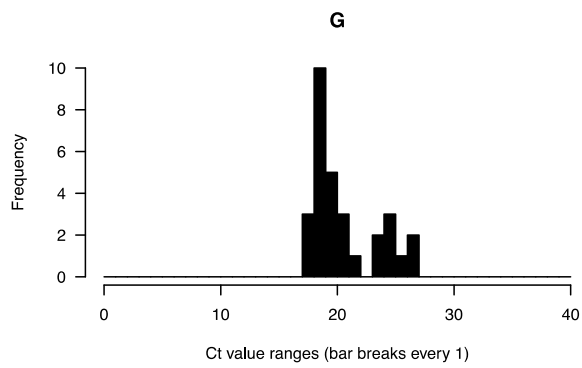
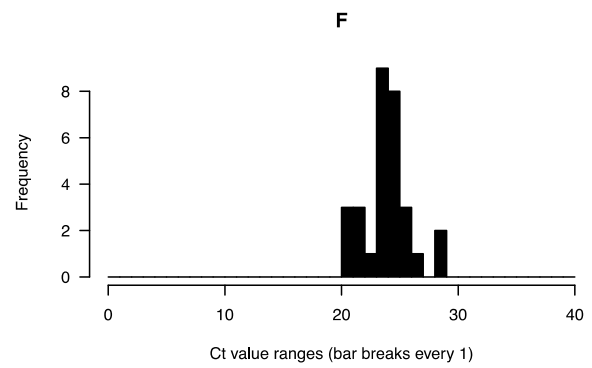
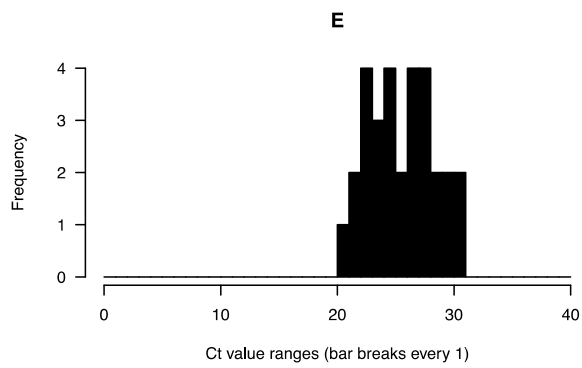
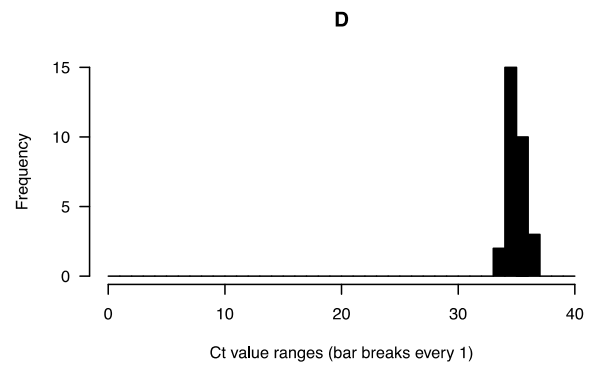
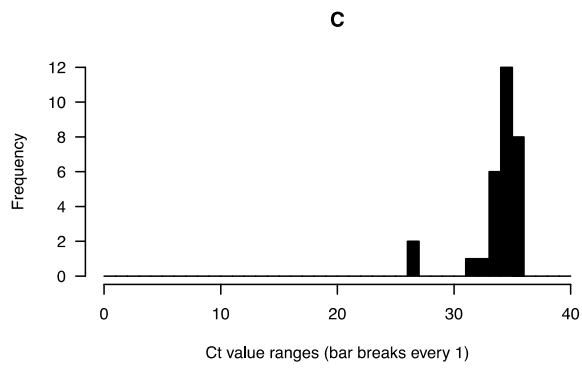
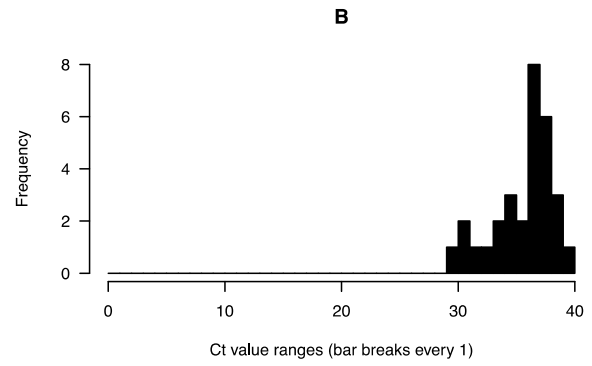
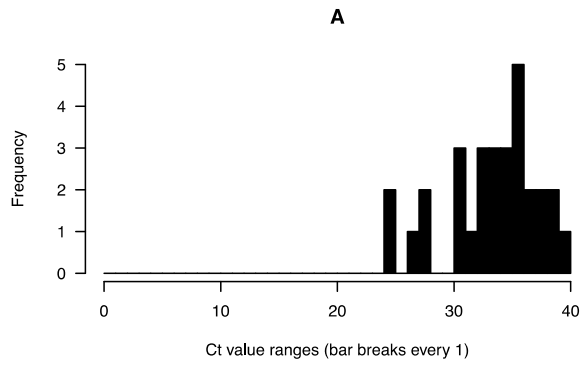
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 5 ¹Each sample was tested for both “*Ca. L. asiaticus*” and *Wolbachia* titer. For each life stage, 30
 6 samples were used and each sample had three technical replicates which were averaged for a final
 7 Ct-value. Those samples with an average technical replicate value of Ct < 35 contributed to the
 8 percent infection rate, (e.g. whole body nymphs tested had 21/30 samples with Ct < 35, or 70%
 9 “*Ca. L. asiaticus*” infection rate). Since all these samples are non-exposed it is expected that all
 10 would be 100% uninfected (Ct > 40) for “*Ca. L. asiaticus*”.

11 ²Mean copy number is the average of individual sample copy numbers with Ct < 35. Cycle
 12 threshold values was converted to copy number using a standard curve.

13 ³The individuals used for guts and whole bodies were taken from separate generations of the same
14 colonies. Exposed nymphs and adults were fed on “*Ca. L. asiaticus*”-infected citrus trees for one
15 or more generations.

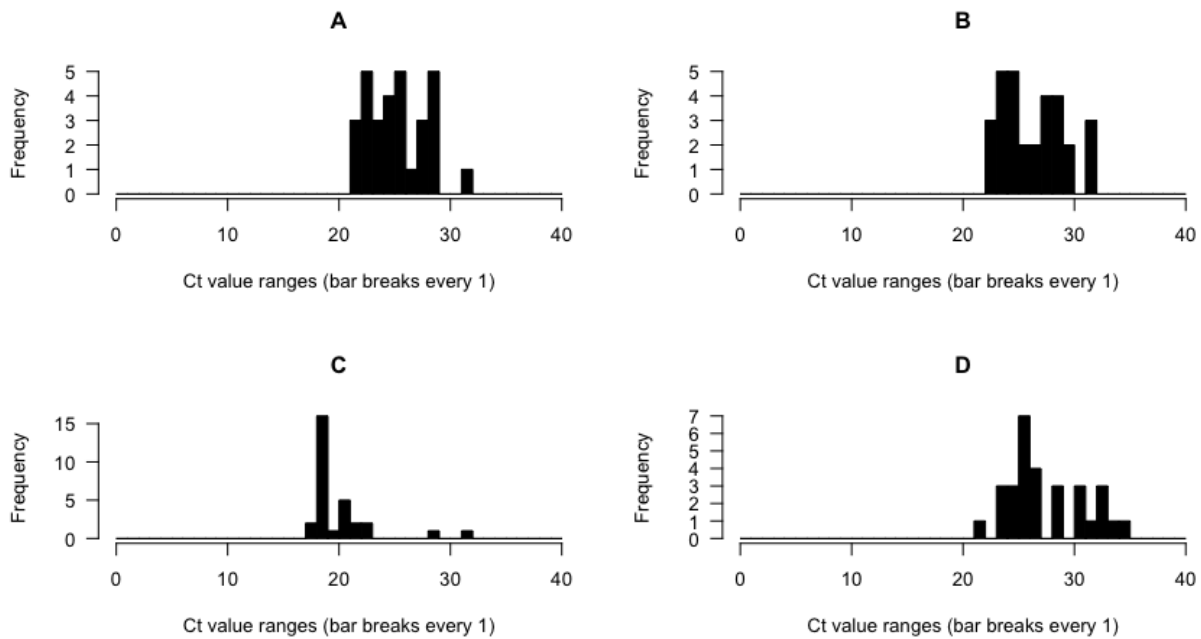
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19 **FIG S1** Frequency histograms of both “*Ca. L. asiaticus*” Ct-values and *Wolbachia* Ct values for
 20 “*Ca. L. asiaticus*”-exposed samples. The frequency of samples with Ct below 35 is visible for each
 21 treatment. The y-axis is frequency (or the number of samples falling within each given range of
 22 data values), while the x-axis is the range of Ct values from 0-40. The number of bars was chosen
 23 to show optimal data distribution. Graphs A-D are “*Ca. L. asiaticus*” Ct values, while graphs E-H
 24 are *Wolbachia* Ct values from only “*Ca. L. asiaticus*”(+) samples. Graphs A-B and E-F are gut
 25 samples (showing adults and nymphs, respectively), while graphs C-D and G-H are whole body
 26 samples (showing adults and nymphs, respectively).

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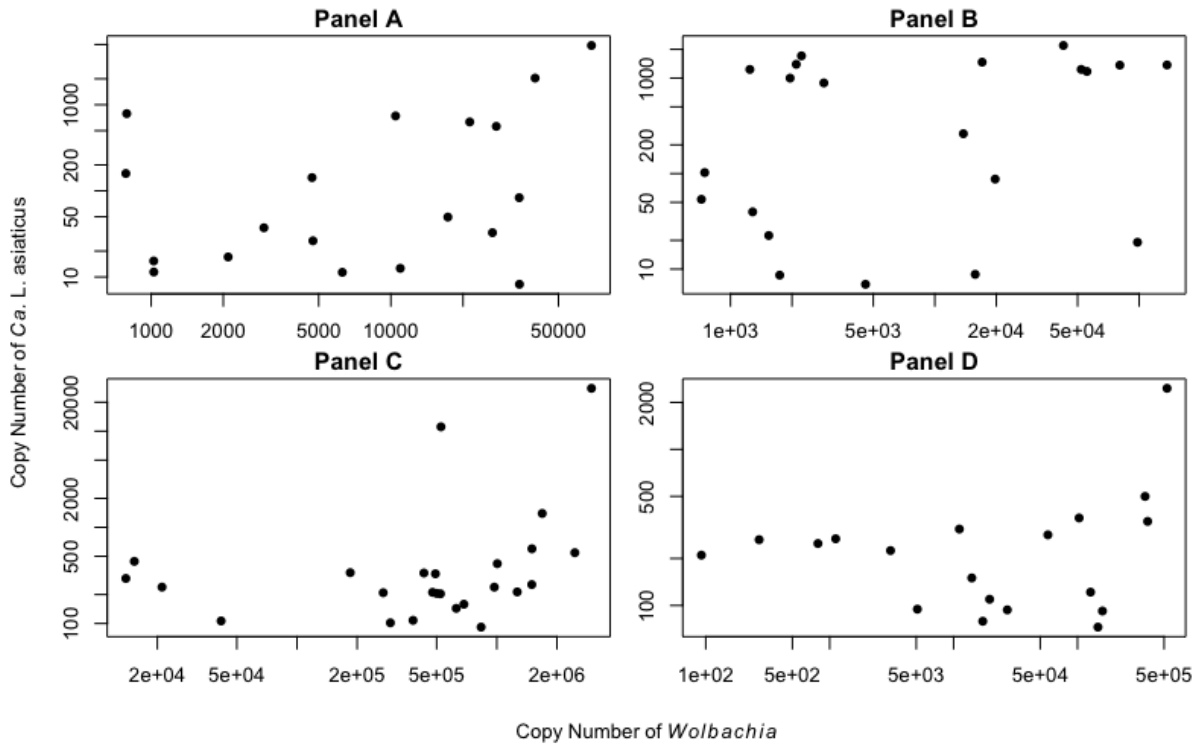


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29 **FIG S2** Frequency histograms of *Wolbachia* Ct values in non-exposed samples used in the Mann-
 30 Whitney-U statistical test. The frequency of samples with Ct below 35 is visible for each treatment.
 31 The y-axis is frequency (or the number of samples falling within each given range of data values),
 32 while the x-axis is the total range of Ct values displayed by the data. The number of bars was

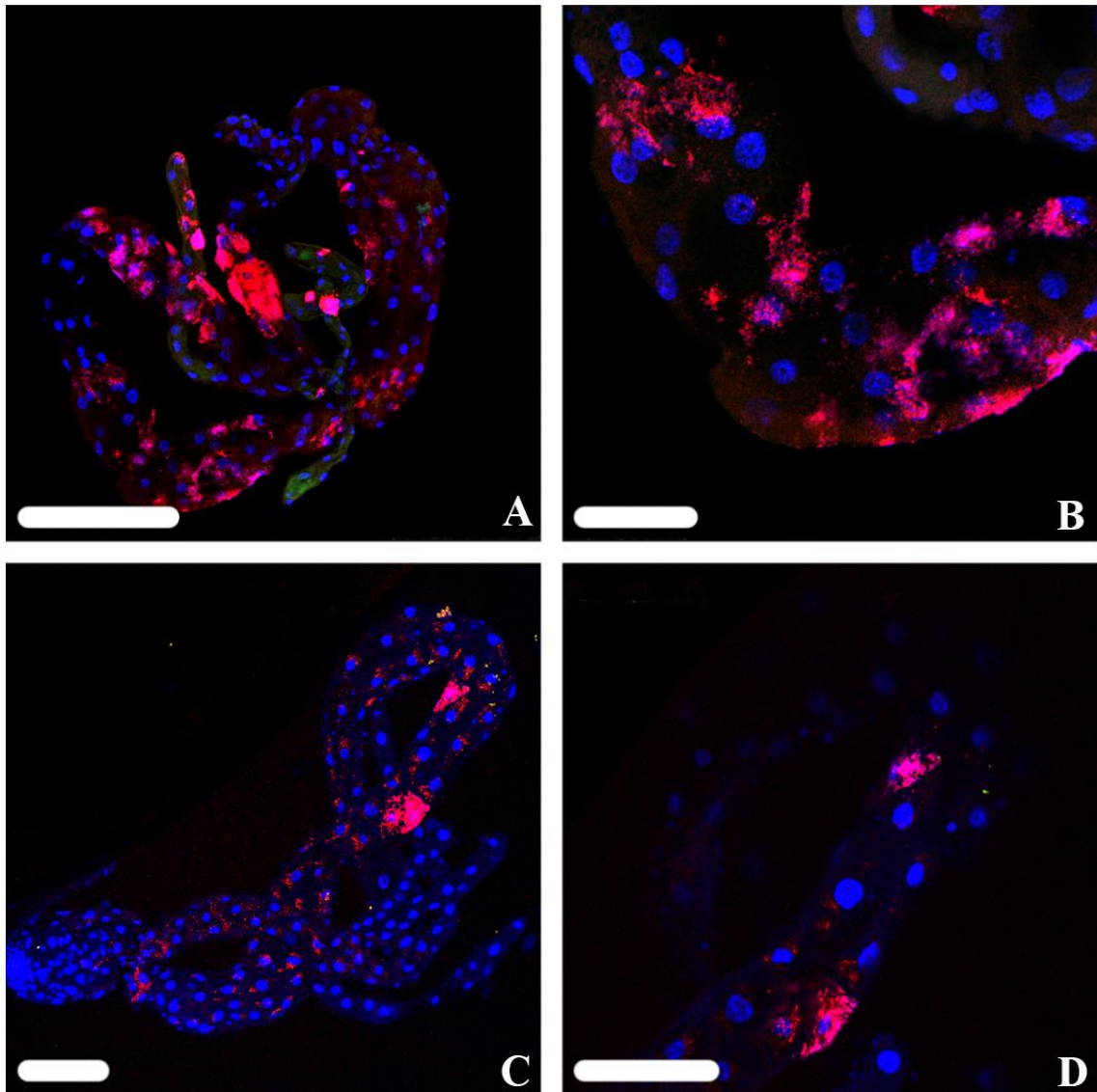
33 chosen to show optimal data distribution. Graphs A-B are whole body (adults and nymphs,
34 respectively), while graphs C-D are guts (adults and nymphs, respectively).

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37 **FIG S3** Distribution of “*Ca. L. asiaticus*” and *Wolbachia* copy numbers in “*Ca. L. asiaticus*”(+)
38 adult guts (panel A), “*Ca. L. asiaticus*”(+) nymph guts (panel B), “*Ca. L. asiaticus*”(+) whole body
39 adults (panel C) and “*Ca. L. asiaticus*”(+) whole body nymphs (panel D). The y-axis is individual
40 “*Ca. L. asiaticus*” copy number, and the x-axis is individual *Wolbachia* copy number, and both
41 axes are log scale. (Panel A, $p < 0.05$, $cor = 0.56$; Panel B, $p < 0.05$, $cor = 0.77$; Panel C, $p < 0.05$,
42 $cor = 0.76$; Panel D, $p = 0.49$, $cor = 0.16$).



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44 **FIG S4** Localization of *Wolbachia* (red) in non-exposed *D. citri* gut cells with DAPI staining of
45 nuclei (blue). Non-exposed adult guts (A-B) and non-exposed nymph guts (C-D) are stained using
46 Fluorescent *in situ* Hybridization (FISH). The scale bar lengths are as follows: A, C: 250 μm and
47 B, D: 75 μm.

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