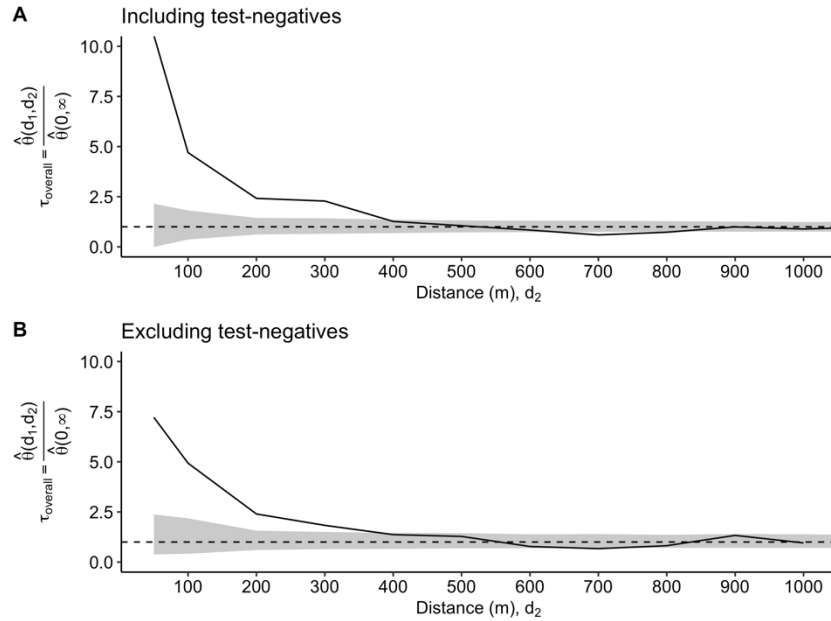# Disruption of spatiotemporal clustering in dengue cases by *w*Mel *Wolbachia* in Yogyakarta, Indonesia

Suzanne M Dufault, Stephanie K Tanamas, Citra Indriani, Adi Utarini, Riris Andono Ahmad, Nicholas P Jewell, Cameron P Simmons, and Katherine L Anders

Supplementary Material

Table S1. The distribution of virologically confirmed dengue cases and test-negative controls by AWED study arm.

	Intervention	Untreated
Test-negative controls	2838	3083
Dengue cases	67	318
DENV1	12	45
DENV2	20	134
DENV3	5	22
DENV4	17	70
Unknown	14	53

Figure S1. Estimated odds ratio, τ, comparing the odds of a homotypic dengue case pair within (d_1, d_2) versus the odds of a homotypic dengue case pair at any distance across the entire study area with A) test-negative controls contributing to the counts of heterotypic pairs, and B) test-negatives excluded and based solely on the heterotypic and homotypic counts of dengue pairs. Variability in estimation is visualized in two distinct ways. A) displays the pointwise 95% confidence interval (CI) based on 1,000 bootstrap resamples of the data and B) sows the pointwise 95% CI on the permutation-based null rejection region based on 1,000 permutations of the data.