

Figure S14: Box plots of the relative abundance of *Proteobacteria* for each genotype at each time point. The median, lower and upper quartiles are shown. Whiskers were calculated using the Tukey method; filled circles represent outliers. Asterisks indicate significant differences (one-way ANOVA, followed by Tukey-Kramer multiple comparisons test, \* P < 0.05; \*\* P < 0.01; \*\*\* P< 0.001).