

Exploring Gut Microbial Dynamics and Symbiotic Interaction in *Blattella germanica* Using Rifampicin
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Figure S1

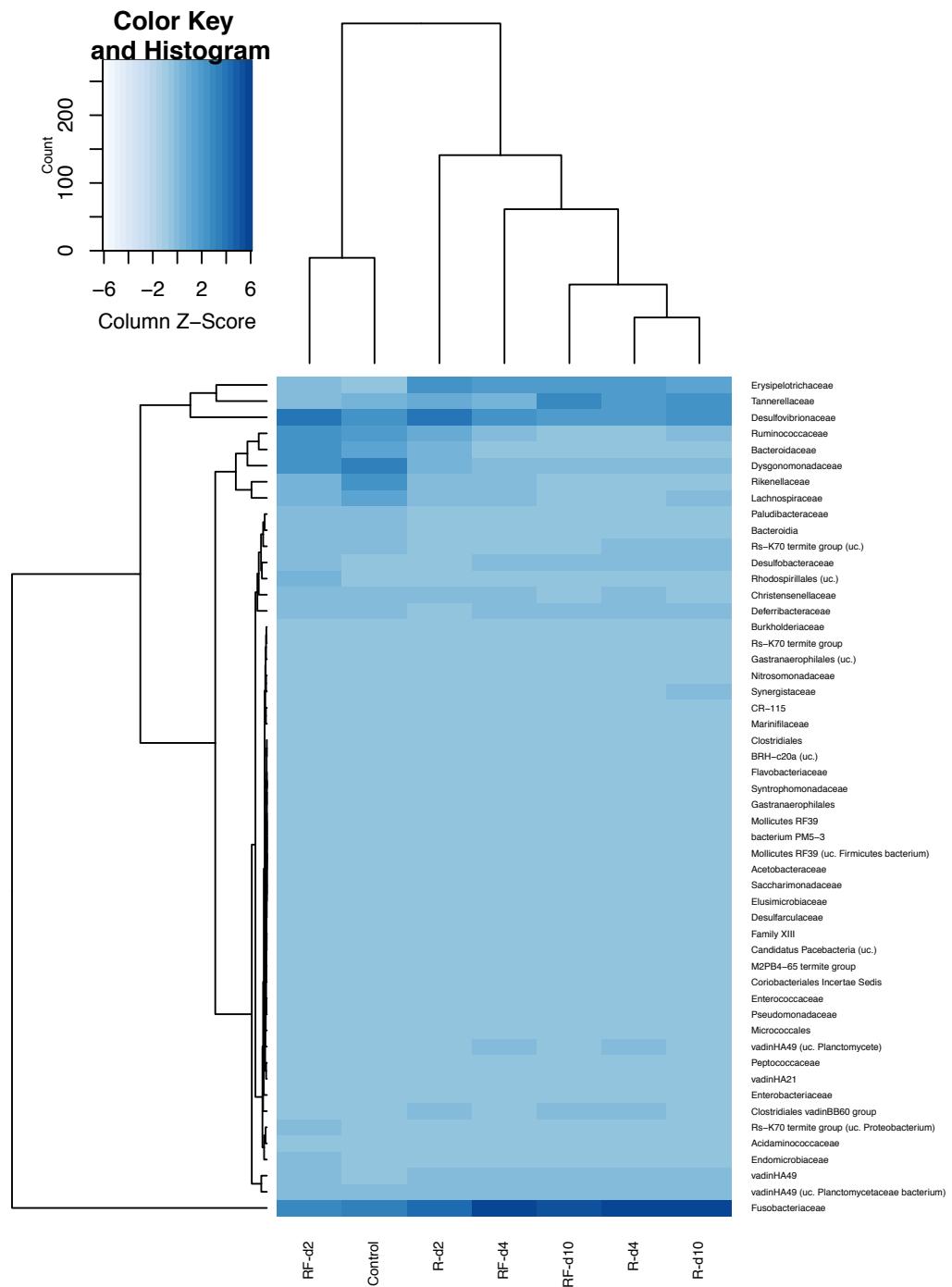


Figure S1. Heatmap and clustering based on taxon composition and Z-score transformed relative abundance of the gut microbiota corresponding to d2, d4 and d10 of treatment. Three populations were analysed: control (C, corresponding to grouped C-d2 to C-d10), treated with rifampicin (R), and treated with rifampicin and faeces (RF). Different experimental conditions are shown on the x-axis and the family of the taxa identified in each sample on the y-axis. The shade of colours was related to the Z-score of these taxa in each sample; light and dark blue indicate lower to higher abundance, respectively.