



Figure S6. Heatmap of Euclidean distances clustering calculated for normalized and transformed (regularized log transformation) expression data of all genes analyzed in each samples using (A) isolates as a condition and (B) times of incubation as a condition. The first part of sample names indicates isolate (CA1, CA2, and CA3) and the second part of sample names indicates time of incubation (24 h, 48 h, and 72 h).