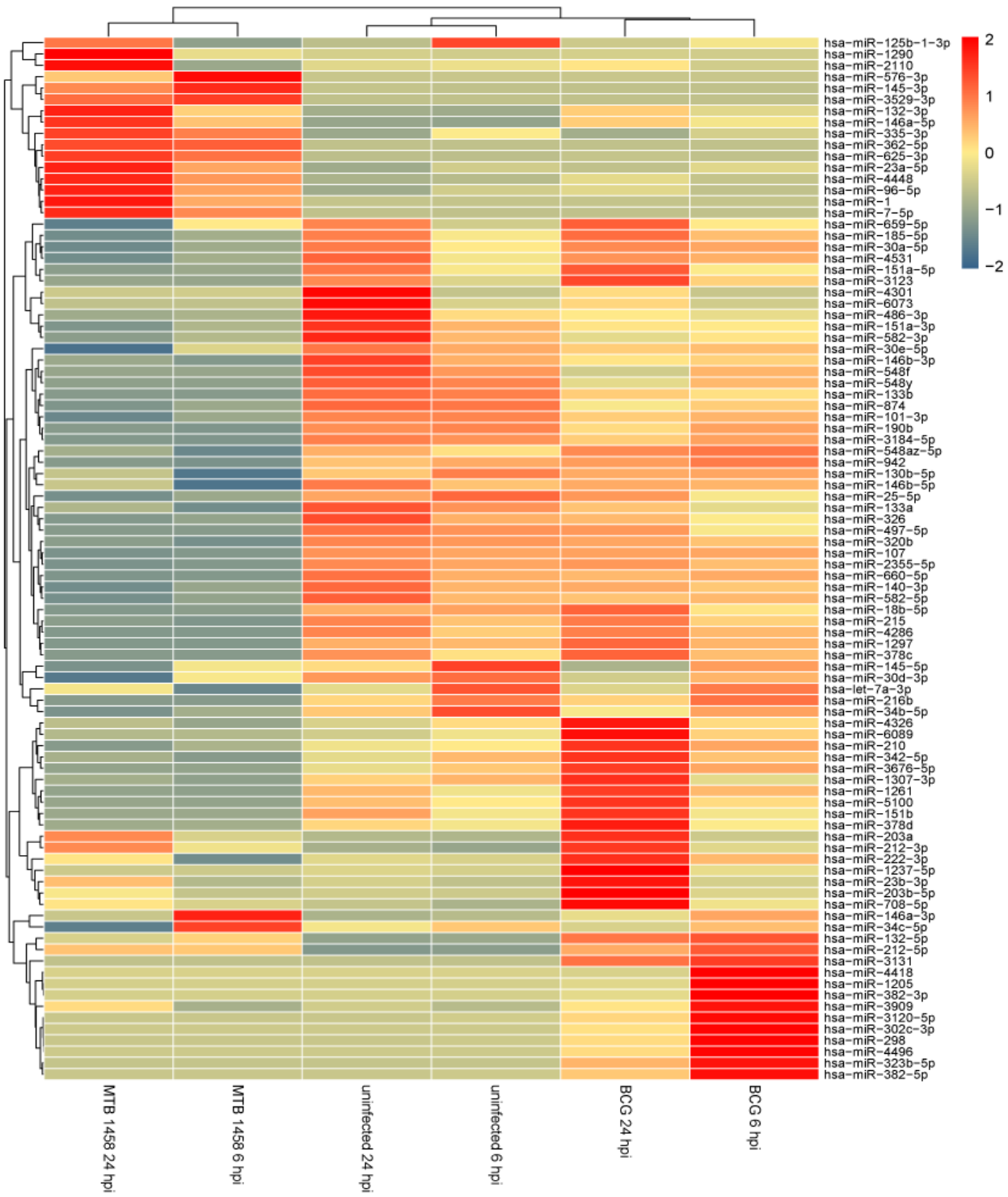


Figure S1. Hierarchical clustering analysis of differentially expressed miRNAs between MTB-1458-infected vs. uninfected cells and BCG-infected vs. uninfected cells

Hierarchical clustering analysis of differentially expressed miRNAs between MTB-1458-infected vs. uninfected cells and BCG-infected vs. uninfected cells. Each row represents miRNA; each column represents a sample. Red indicates higher expression in MTB-1458- or BCG-infected cells in comparison with uninfected cells; blue indicates lower expression in MTB-1458- or BCG-infected cells in comparison with uninfected cells.