



Figure S2. Heatmaps representing the expression patterns of genes involved in secondary metabolism. (a) Flavonoid biosynthesis pathway. (b) Terpenoid biosynthesis pathways, including terpenoid backbone biosynthesis (Terp. Backbone), monoterpenoid biosynthesis, sesquiterpenoid biosynthesis and diterpenoid biosynthesis pathways. The colour gradient represents mean expression levels (logTPM) of each gene for control (C), susceptible (S) and resistant (R) samples.