



S6 Fig. Time-course analysis of *SIRDR1* expression levels. The expression levels of endogenous *SIRDR1* mRNA were analyzed by RT-qPCR. qPCR analysis was performed with the PCR primers for endogenous *SIRDR1* mRNA. Mean values are based on three biological replicates of the pooled sample of five individual plants. The relative expression levels were calculated for each time point with the value of EC plants inoculated with mock as a standard. The expression levels of endogenous *SIRDR1* mRNA were apparently different between Int- or RG1-infected *SIRDR6i* plants at later infection stage, or between RG1-infected EC and *SIRDR6i* plants at 15 dpi.