

Figure S1

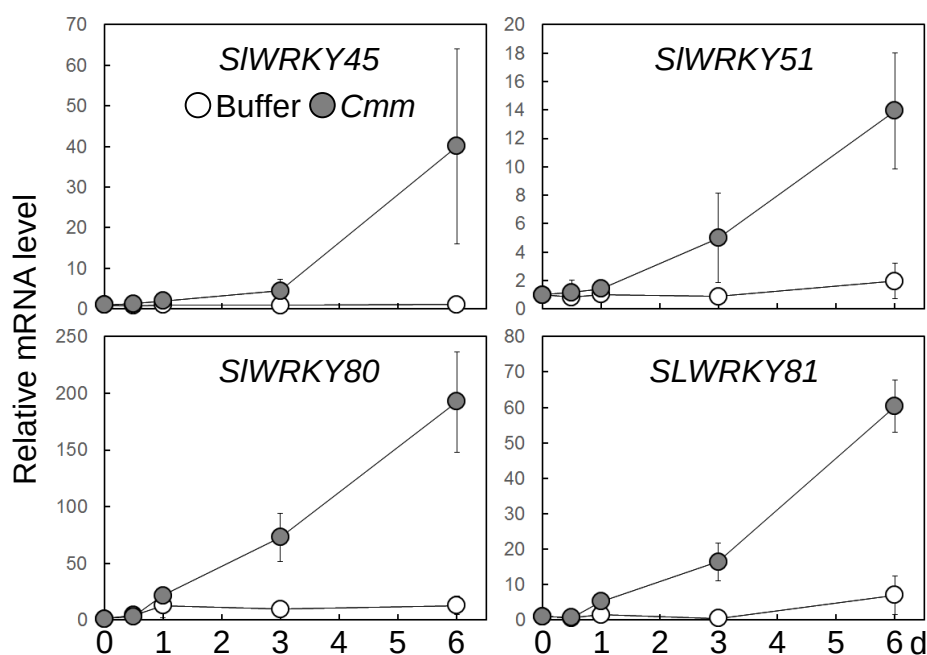


Fig. S1. Expression of WRKY genes in tomato plants after infection with *Cmm*. The transcript levels in tomato cotyledons were quantified by qRT-PCR analysis and expressed relative to the transcript level at 0 day, which was assumed to be one. Data are represented as mean values $\pm$ the standard deviation for three replicates.