

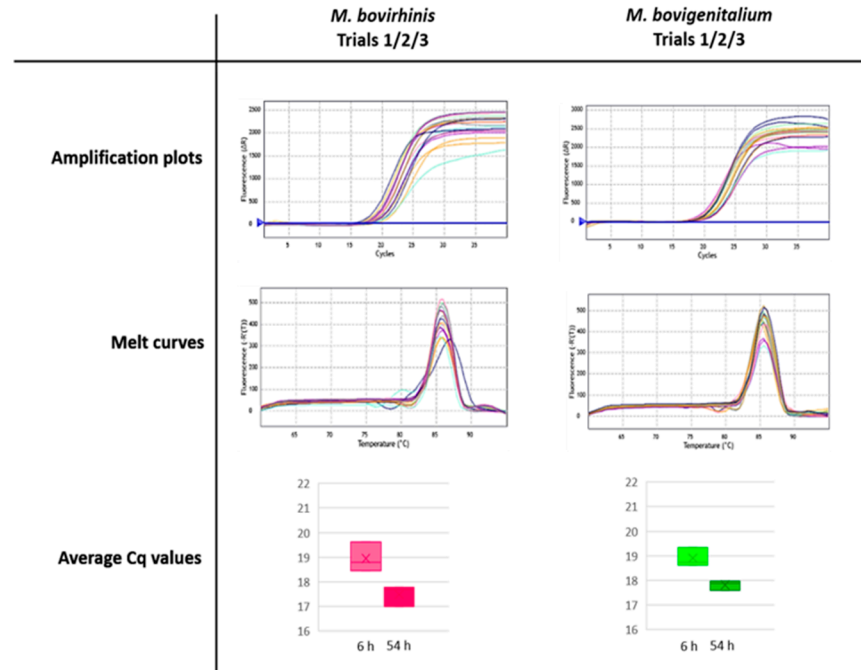
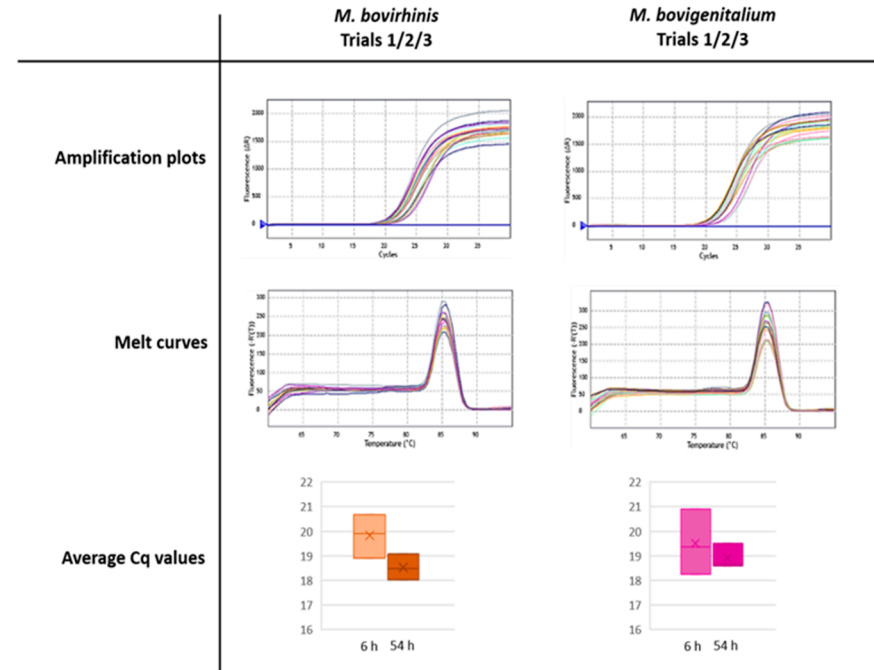
A**B**

Figure S4. qPCR amplification of eubacterial 16S rRNA gene (**A**) and bovine 18S rRNA gene (**B**) after MDBK cells infection with *Mycoplasma bovirhinis* and *Mycoplasma bovigenitalium* strains. Amplification plots resulted from triplicate qPCR using DNA extracted from cell lysates collected after 6 h and 54 h in three independent infection experiments. For each sample at each time point and at every trial, mean Cq values were calculated from triplicate qPCR reactions for each gene. Box and whisker plots show the variation of mean Cq values of the different samples. Amplification plots and melt curves were generated using Agilent Aria software version 1.71.