Figure S2: RT-PCR-based expression analysis of hydrophobin genes in mutant strains Δbhp1/bhp2, Δbhp3/bhp2 and Δbhl1.

M: Size markers, with relevant sizes indicated; Co: Resting conidia; My: mycelium (15 h.p.i.); To: Infected tomato leaves (48 h.p.i.); Sc: Sclerotia. See Fig. 2 for water and genomic DNA controls. An EF1α encoding fragment was amplified as positive control. White arrows indicate positions of bands based on cDNA. Undiluted first-strand cDNA was amplified with 35 cycles, except for ef1α cDNA, which was amplified from 1:10 diluted first-strand cDNA. The PCR bands based on genomic DNA that were obtained for Δbhp3 mutants in bhp3-specific reactions presumably result from a minor contamination of the knock-out strain with untransformed wild type nuclei.