Table S4. *Induction of defense-related genes in Rio Grande and Ailsa Craig tomato lines upon Cmm infection*

GenBank accession	Gene product	Array ratio[a] Rio Grande (*Cmm*/mock)	qRT-PCR ratio[b] Rio Grande (*Cmm*/mock)	qRT-PCR ratio[b] Ailsa Craig (*Cmm*/mock)
U89256	Pti5, ERF/AP2 transcription factor	163	193	5
AF272366	*Verticillium* wilt disease resistance protein	88	3	2
AI776170	Cysteine protease	81	116	8
AJ133600	Extensin	20	105	94
BG629612	Chitinase	55	317	15
AY359965	EIX receptor 1	10	9	3
AW219676	Wound induced protein Sn-1	8	21	15
AY093595	Osmotin-like protein	10	2	4
CN385925	Patatin-like protein 3	5	4	2
M69248	Pathogenesis-related protein 6 (P6)	13	Not tested	26
U30465	Chitinase class 2	11	Not tested	19
Z15141	Endochitinase	10	Not tested	16
BM956714	Protease inhibitor/lipid transfer protein	6	Not tested	5
K03291	Wound-induced protease inhibitor II	24	Not tested	21
BT012691	2-oxoglutarate-dependent dioxygenase E8	22	Not tested	8
M69247	Pathogenesis-related protein PR1a (P4)	16	Not tested	13
BT013355	Pathogenesis-related protein 2 (P2)	3	Not tested	19

[a]Ratio between expression in *Cmm*- and in mock-inoculated stems as estimated by microarray analysis at 8 dpi. Data represent the mean of two biological replicates.

[b]Ratio between expression in *Cmm*- and in mock-inoculated stems as estimated by qRT-PCR at 8 dpi. Data represent the mean of three and four biological replicates in Rio Grande and Ailsa Craig plants, respectively.