

**Table S1. Primers designed for validation of gene expression profile in *C. cinerea* transcriptome data**

<b>Primer ID</b>	<b>Primer sequences (5' to 3')</b>	<b>Amplicon size</b>
CC1G_07486FP	AGACGCGACGGACTACTTTC	101bp
CC1G_07486RP	TAGTGCCCTGGGTTCATACG	
CC1G_04403FP	AAGGTGGTCTTTGCCACAA	99bp
CC1G_04403RP	CCACCACGATCAACTGACGA	
CC1G_09477FP	ATGTCGCGACGGATTCTAC	99bp
CC1G_09477RP	ATTGCGGACCAGGAGATGGA	
CC1G_00058FP	TCCTGCTACGTGGCAAATGA	100bp
CC1G_00058RP	TGGATCATCGACGCATGGTT	
CC1G_00407FP	TTCAGTCACGCGTAGGGAAG	123bp
CC1G_00407RP	AACAGTTCTCCGCTTGTCCC	
CC1G_15600FP	CCTGGCAGTGTTGAAGACCC	161bp
CC1G_15600RP	CCGATGAGACCATGCAGAAGA	
CC1G_10354FP	TCGTTGATCACGTCAGTGGAA	144bp
CC1G_10354RP	ACGGGTACGGATGGAGACT	
CC1G_09588FP	TTAGGCGCGTGCCATTCAAG	105bp
CC1G_09588RP	TATCCCTGAGCGAACGACAC	
CC1G_00620FP	CCATTGACCGACGACCCTAC	96bp
CC1G_00620RP	TACGCCGATGGAGACACAGA	
CC1G_01088FP	AGCGCTGAGGAAGCAGAAAA	101bp
CC1G_01088RP	GCTCGTCGTCTGGATCACTT	
Beta-tubulin_FP	GAGCAGTCCTGGTTGACCTC	130bp
Beta-tubulin_RP	TGTCCTTTGGCCCAGTTGTT	

FP: forward primer; RP: reverse primer.