

**S2 Table. Oligonucleotide primers used in this study**

Targeted gene <sup>a</sup>	Source <sup>b</sup> Broad Institute / GenBank	Forward primer 5' - 3'	Reverse primer 5' - 3'	Expected amplicon size (bp)	Annealing temperature (°C)
pacC	GGTG_01809	caagcgcgatcacatcacct	gagaaccgagtcgctcggc	139	57
palB	GGTG_09525	gcagcaggacatctttgcag	gcggcatcaaagggatacat	243	63
palC	GGTG_06037	aagaagagcggcttccgtt	ggtatcattttgcttgaccatt	165	58
palF	GGTG_01551	atgactatggggcaatgggg	ctccagcttaggtcgaagcgaa	150	60
lac1	AJ417685/GGTG_02417	cgctcgagaggatctttgac	gtggatggggtgggtgag	146	62
lac2	AJ417686/GGTG_02239	tccgctggtacctcaactc	atgacgtaggccactgtc	143	62
gmk1	AF258529/GGTG_05241	agaagctcctggccttcaac	ttgctgaggtgtccttgtg	160	60
exo	CF554536/GGTG_01690	gcgggactggagtactacgg	gggtcctgccagatggagat	146	60
Wn1	Sabater-Muñoz et al., 2006	tcatttccttctgggcattc	gcacaaaacactgccatcaa	146	61
Wn2	Sabater-Muñoz et al., 2006	gtaacgctcgtcccactttcaat	ctaacgctcggcgtaaactaat	140	51

<sup>a</sup> *lac1*, laccase 1; *lac2*, laccase 2; *gmk1*, mitogen-activated protein (MAP) kinase; *exo*,  $\beta$ -1,3-exoglucanase; *Wn1*, 3' *Wunen* EST region amplified in qRT-PCR; *Wn2*, 5' *Wunen* EST region amplified in qRT-PCR (Daval et al., 2013).

<sup>b</sup> [http://www.broadinstitute.org/annotation/genome/magnaporthe\\_comparative/GenomesIndex.html](http://www.broadinstitute.org/annotation/genome/magnaporthe_comparative/GenomesIndex.html), GenBank accession number, or reference [24].