

**Supplementary Table 2.** Oligonucleotide primers used in this study and their positions in each strain

Name	Sequence 5'-3'	Location in the <i>cyt b</i> gene
BDcy9-1F	CGTAGAGTTCGTCGGGATC	-314 to -296 <sup>a</sup>
BDcy1131R	TGTCCCCTTGTATCGCTCAT	1,131-1,150 <sup>a</sup>
BDcy807F	ACCTGGAAACATGACAAAGGG	807-827 <sup>a</sup>
BDcy2039R	TCTAGCCAAGGTAAGTGCGC	2,039-2,058 <sup>a</sup>
BDcy1710F	TCACTTCCTAGGGGCTGAGA	1,710-1,729 <sup>a</sup>
BDcy3221R	CAAAAGTAAAGTCACCCGTCTGA	3,221-3,242 <sup>a</sup>
BDcy3128F	TTTGGGGCTAGGGGTGAAAA	3,128-3,147 <sup>a</sup>
BDcy4461R	CCTGTGCGGGGAAGTATCAGT	4,461-4,480 <sup>a</sup>
BDcy3961F	TAGTGGGTAAAGATGCTCGGTT	3,961-3,982 <sup>a</sup>
BDcy5581R	ACCAAAAGTTCCTTCTCCTTCT	5,581-5,602 <sup>a</sup>
BDcy4986F <sup>c</sup>	TCCAAAAACAGCTGAAAAGTTG	4,986-5,007 <sup>a</sup> , 365-386 <sup>b</sup>
BDcy6122R	TGCTGTTTAGATCTATTCGCTGT	6,122-6,144 <sup>a</sup>
BDcy6514R	CAGGAAAGCAGTAGCCATCA	6,514-6,533 <sup>a</sup> , 1,893-1,912 <sup>c</sup>
BDcy687F <sup>c</sup>	CACGCTAACACTGCTTCTGC	6,387-6,406 <sup>a</sup> , 1,766-1,785 <sup>b,c</sup>
BDcy2467R <sup>c</sup>	TCGACATGAACACCTTCTTGGA	8,146-8,167 <sup>a</sup>
BDcy7586F	TCCCGCGAGTAACAATCTTT	7,586-7,605 <sup>a</sup>
BDcy8738R	ACCGACGGTCTGAATGCTAAA	8,738-8,758 <sup>a</sup>
BDcy9173R	AAAGGGTTACCTGACCCAGC	9,173-9,192 <sup>a</sup>
BDcy8731F	ATGTTTGTTAGCATTAGACCG	8,731-8,753 <sup>a</sup>
BDcy10057R	GTTGGATTAAATCATGTTGGGTAT	+340 to +362 <sup>a</sup>
Bdcy9-2F	TTTCTTTATGCAATCATGTGATAATG	-192 to -167 <sup>b,c</sup>
BDcy6806R	GGTTTTCAAAAATATACACAGGATTT	2,185-2,210 <sup>b</sup>
BDcy1542R <sup>c</sup>	TTCCCGTCTTGCTTCGCTA	2,602-2,621 <sup>b</sup>
BDcy7767R	CAGCATTTAGCCACCCATT	3,146-3,165 <sup>b,c</sup>
BDcy7586F <sup>c</sup>	TCCCGCGAGTAACAATCTTT	2,965-2,984 <sup>b,c</sup>
BDcy9347R <sup>c</sup>	GCAGGTGGAGTTTGCATAGG	6,136-6,155 <sup>b</sup>
BKey3292F	ACACATAGCTTCTACAGGGAGA	3,929-3,950 <sup>b</sup>
BKey5429R	ATTCAAGCGACAAAATTTTCTCT	5,429-5,451 <sup>b</sup>
BDcy9012F	TGAGTGCTATACCTTGAGTTGGA	3,330-3,352 <sup>b</sup>
BDcy9747R	TCTACAAAATAGAAAACAAAATCGC	+30 to +54 <sup>b,c</sup>
BDeye1F	AAGTCATCCTCTTTTAAACTGGTT	15-39 <sup>d</sup>
BDeye2F	AGCGATGCATTACAATCCTAGTG	150-172 <sup>d</sup>
BDeye2R	CAGTACATGCTTGACCTAGTTCT	1,050-1,072 <sup>d</sup>
BDeye6R	CTGACCCTGCAGTATCATGT	603-622 <sup>d</sup>

<sup>a</sup>Nucleotide range in the cytochrome b (*cyt b*) gene of *Botryosphaeria sinensis* GsYjBD22-1-1 and GwYwBD22-2-1.

<sup>b</sup>Nucleotide range in the *cyt b* gene of *Botryosphaeria kuwatsukai* CcYsBD22-3-1.

<sup>c</sup>Nucleotide range in the *cyt b* gene of *B. kuwatsukai* 22-MG32.

<sup>d</sup>Nucleotide range in the exon sequence of the *cyt b* gene from *Botryosphaeria* spp.

<sup>e</sup>Primers used for the detection of F129L, G137R, and G143A mutations in *cyt b*.