

Supporting Information Table S5

PCR efficiencies calculated with qPCR of all species and primer pair combinations used to assess phylum-level PCR bias in the current study

Genus/species	Strain	PCR efficiency (%)		
		ITS1F/ITS2	ITS3/ITS4	ITS86F/ITS4
<i>Cladosporium cladosporioides</i>	MUCL 53652	55	85	92
<i>Cryptosporiopsis radicicola</i>	MUCL 53485	58	56	131
<i>Monilinia laxa</i>	MUCL 30841	74	72	78
<i>Arthroderma otae</i>	MUCL 39756	36	80	91
<i>Galactomyces geotrichum</i>	MUCL 52377	106	87	39
<i>Lentinula edodes</i>	MUCL 44827	70	39	77
<i>Agrocybe praecox</i>	MUCL 46727	73	98	112
<i>Coniophora marmorata</i>	MUCL 39471	73	64	69
<i>Suillus luteus</i>	UH-Slu-Lm8-n1	85	88	94
<i>Antrodia vaillantii</i>	MUCL 54533	51	88	101
<i>Rhizophagus clareus</i>	MUCL 46238	117	126	85
<i>Rhizophagus</i> sp.	MUCL 41833	116	82	74
<i>Mortierella verticillata</i>	MUCL 9658	106	74	86
<i>Absidia corymbifera</i>	MUCL 38907	60	74	65
<i>Mucor hiemalis</i>	MUCL 15439	88	98	93
Average		78	81	86
Standard error		6	5	5