CORRECTION





Correction to: Brazilian fungal diversity represented by DNA markers generated over 20 years

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Due to a processing error, there was a mistake in Table 3. The first entry in the right column should read 109. The corrected table is given below.

The online version of the original article can be found at https://doi.org/ 10.1007/s42770-019-00206-y

Table 3 Fungal genera occurring in Brazil with the greatest number of ITS sequences and their genetic diversity based on OTUs (ITS 98% cut-

Genera ordered by number of ITS sequences		Genera ordered by number of OTUs	
Genera (phylum)*	n° of sequences	Genera (phylum)*	n° of OTUs
Colletotrichum (A)	969	Phyllosticta (A)	109
Candida (A)	536	Penicillium (A)	95
Phyllosticta (A)	467	Diaporthe (A)	90
Fusarium (A)	449	Candida (A)	83
Phakopsora (B)	444	Fusarium (A)	80
Trichoderma (A)	289	Colletotrichum (A)	76
Diaporthe (A)	257	Aspergillus (A)	63
Aspergillus (A)	204	Trichoderma (A)	44
Lasiodiplodia (A)	204	Rhizoctonia (B)	38
Fonsecaea (A)	200	Pluteus (B)	29
Penicillium (A)	199	Cora (B)	24
Puccinia (B)	168	Cryptococcus (B)	18
Cryptococcus (B)	94	Fonsecaea (A)	16
Rhizoctonia (B)	90	Ceratobasidium (B)	16
Cora (B)	70	Lasiodiplodia (A)	9
Ceratobasidium (B)	56	Puccinia (B)	4
Pluteus (B)	56	Phakopsora (B)	3

^{*}Genera with > 150 ITS sequences for Ascomycota (A) and > 50 ITS sequences for Basidiomycota (B)

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