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CORRECTION

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Correction to: Molecular identification and antifungal susceptibility profile of yeast from vulvovaginal candidiasis

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Following publication of the original article [1], the authors identified an error in Table 3. The correct table (Table 3) is given in this erratum and the original article has been corrected.

The original article can be found online at https://doi.org/10.1186/s12879-020-04985-w.

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Table 3 In vitro antifungal susceptibility of 1844 clinical isolates of Candida species as determined by the CLSI method

***Candida* species (n)**	**Antifungal agents**							
		BUC	**CLO**	**FLC**	**ITC**	**MIC**	**TEC**	**VRC**
C. lusitaniae, n = 1	Range	0.03	0.03	0.125	0.03	0.03	0.03	0.03
C. Fabianii, n = 3	Range	0.03–0.25	0.03–0.06	0.5–1	0.03–0.25	0.125–1	0.03	0.03
	MIC50	0.125	0.03	0.5	0.06	0.125	0.03	0.03
	MIC90	0.25	0.06	1	0.25	1	0.03	0.03
Trichosporon asahii, n = 1	Range	0.03	0.03	0.25	0.03	0.06	0.03	0.03
Rhodotorula, n = 3	Range	0.03–0.5	0.03–1	4–128	0.03–8	0.25–8	0.03–0.5	0.03–1
	MIC50	0.06	0.06	64	2	1	0.06	0.03
	MIC90	0.5	1	128	8	8	0.5	1
Kodamaea ohmeri, n = 2	Range	0.125–0.5	0.03	0.25–2	0.125–0.25	0.25–0.5	0.03	0.03
	MIC50	0.125	0.03	0.25	0.125	0.25	0.03	0.03
	MIC90	0.5	0.03	2	0.25	0.5	0.03	0.03
Issatchenkia terricola, n = 2	Range	1–4	0.06–0.125	32–64	0.25–0.5	0.5	0.25	0.25
	MIC50	1	0.06	32	0.25	0.5	0.25	0.25
	MIC90	4	0.125	64	0.5	0.5	0.25	0.25
Torulaspora pretoriensis, n = 1	Range	0.25	0.03	8	0.5	0.5	0.125	0.125
*ATCC90028**	Range	0.015–0.5	0.015–0.5	0.125–2	0.015–4	0.008–0.015	0.015–32	0.015–8
	GM	0.04	0.03	0.21	0.08	0.06	0.03	0.03
	MIC90	0.125	0.03	0.5	0.25	0.015	0.03	0.03
***Candida* species (n)**	**Antifungal agents**							
		AmB	**FLU**	**NYS**	**TEB**	**AFG**	**CFG**	**MFG**
C. albicans n = 1272	Range	0.015–32	0.03–128	0.03–32	0.03–256	0.008–0.5	0.008–0.5	0.008–0.5
	GM	0.22	0.70	1.60	45.11	0.015	0.1	0.03
	MIC90	0.5	4	8	256	0.03	0.25	0.25
	R		3.3%			0	0	0
C. africana n = 49	Range	0.03–32	0.06–8	0.125–4	0.25–256	0.008–0.03	0.015–0.5	0.008–0.5
	GM	0.08	0.68	0.5	17.31	0.01	0.06	0.02
	MIC90	1	2	4	128	0.015	0.25	0.06
C. dubliniensis n = 1	Range	0.06	0.06	0.25	16	0.008	0.015	0.008
C. glabrata n = 267	Range	0.03–2	0.06–16	0.03–32	0.25–256	0.008–0.5	0.008–0.5	0.008–0.5
	GM	0.29	0.18	3.39	26.62	0.03	0.11	0.05
	MIC90	1	1	8	256	0.06	0.25	0.25
	R		0			0	0	0
C. nivariensis n = 9	Range	0.06–2	0.125–4	0.5–4	1–256	0.015–0.06	0.08–0.5	0.015–0.5
	MIC50	0.06	0.5	1	128	0.06	0.25	0.015
	MIC90	2	2	4	256	0.06	0.5	0.5
C. bracarensis n = 2	Range	0.06–1	0.125–2	0.25–8	8–256	0.015–0.03	0.125–0.5	0.015–0.5
	MIC50	0.06	0.125	0.25	8	0.015	0.125	0.015
	MIC90	1	2	8	256	0.03	0.5	0.5
C. parapsilosis n = 76	Range	0.03–2	0.125–8	0.03–32	0.25–256	0.008–1	0.008–1	0.008–1
	GM	0.19	0.14	0.59	0.62	0.69	0.60	0.54
	MIC90	1	0.125	4	32	0.5	0.5	0.5
	R		0			5.2%	5.2%	1.3%

Table 3 (continued)

***Candida* species (n)**	**Antifungal agents**						
		AmB	**FLU**	**NYS**	**TEB**	**AFG**	**CFG**	**MFG**
C. metapsilosis n = 20	Range	0.015–0.5	0.125–4	0.06–4	0.25–256	0.015–0.5	0.008–0.5	0.015–1
	GM	0.10	0.177	0.46	2.17	0.17	0.17	0.39
	MIC90	0.5	1	4	256	0.25	0.25	0.5
***Candida* species (n)**	**Antifungal agents**						
		AmB	**FLU**	**NYS**	**TEB**	**AFG**	**CFG**	**MFG**
C. orthopsilosis n = 6	Range	0.06–0.25	0.125–2	0.06–8	0.25–128	0.008–1	0.015–1	0.008–0.5
	MIC50	0.125	1	0.5	64	0.08	0.30	0.25
	MIC90	0.25	2	8	128	1	1	0.5
C. tropicalis n = 61	Range	0.03–1	0.125–32	0.03–8	0.25–256	0.015–0.125	0.008–0.5	0.008–0.5
	GM	0.19	0.23	0.54	60.02	0.03	0.24	0.04
	MIC90	0.5	1	4	256	0.06	0.5	0.5
	R		1.8%			0	0	0
C. krusei n = 54	Range	0.03–1	0.125–32	0.03–4	16–256	0.015–0.5	0.008–1	0.008–0.5
	GM	0.43	4.2	0.32	75.66	0.08	0.08	0.15
	MIC90	1	16	1	256	0.125	0.5	0.25
	R		2.9%			0	1.85%	0
Saccharomyces cerevisiae n = 12	Range	0.03–4	0.06–8	0.125–32	0.25–256	0.015–0.5	0.08–0.5	0.015–0.5
	GM	0.18	0.21	0.78	53.20	0.14	0.10	0.18
	MIC90	1	1	8	256	0.5	0.25	0.25
C. guilliermondii n = 2	Range	0.06–0.5	0.125–0.25	0.25–0.5	64–128	0.015–0.25	0.25–0.5	0.015–0.25
	MIC50	0.06	0.125	0.25	64	0.015	0.25	0.015
	MIC90	0.5	0.25	0.5	128	0.25	0.5	0.25
	R		0			0	0	0
C. lusitaniae n = 1	Range	0.125	0.125	0.25	0.25	0.03	0.25	0.015
C. fabianii n = 3	Range	0.06–0.25	0.125	0.25	128	0.015–0.03	0.015–0.25	0.015–0.06
	MIC50	0.06	0.125	0.25	128	0.015	0.015	0.03
	MIC90	0.25	0.125	0.25	128	0.03	0.25	0.06
Trichosporon asahii n = 1	Range	0.125	1	0.25	128	0.015	0.5	0.015
Rhodotorula n = 3	Range	0.03–0.5	0.125	0.06–0.125	8–256	0.06–0.5	0.25–0.5	0.25–0.5
	MIC50	0.03	0.125	0.06	8	0.06	0.25	0.25
	MIC90	0.5	0.125	0.125	256	0.5	0.5	0.5
Kodamaea ohmeri n = 2	Range	0.125–1	0.125	0.5	128	0.015	0.125–0.25	0.015–0.03
	MIC50	0.125	0.125	0.5	128	0.015	0.125	0.015
	MIC90	1	0.125	0.5	128	0.015	0.25	0.03
Issatchenkia terricola n = 2	Range	0.125–0.5	8	0.5	128–256	0.06	0.06	0.08
	MIC50	0.125	8	0.5	128	0.06	0.06	0.08
	MIC90	0.5	8	0.5	256	0.06	0.06	0.08

Table 3 (continued)

Candida species (n)	Antifungal agents						
		AmB	FLU	NYS	TEB	AFG	CFG	MFG
Torulaspora pretoriensis n = 1	Range	0.125	0.125	0.25	16	0.015	0.125	0.015
ATCC90028[a]	Range	0.03–2	0.125–8	0.25–16	1–256	0.008–0.015	0.015–0.5	0.008–0.015
	GM	0.22	0.64	1.31	88.22	0.01	0.09	0.01
	MIC90	1	2	8	256	0.015	0.5	0.015

GM geometry mean, *BUC* butoconazole, *CLO* Clotrimazole, *FLC* Fluconazole, *ITC* Itraconazole, *VRC* Voriconazole, *MIC* Miconazole, *TEC* Terconazole, *AmB* Amphotericin B, *FLU* Flucytosine, *NYS* Nystatin, *TEB* Terbinafine, *AFG* Anidulafungin, *CFG* Caspofungin, *MFG* Micafungin

[a] ATCC90028 was tested 57 times

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