

Distribution and susceptibility profile of *Candida* isolates from HIV patients with oropharyngeal candidiasis

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Abstract

Background: Opportunistic infections are the leading cause of morbidity and mortality among immuno-compromised patients. Oropharyngeal candidiasis (OPC) dominates opportunistic fungal infections associated with HIV/AIDS.

Objective: We determined the distribution and prevalence of antifungal resistance in *Candida* isolates recovered from patients infected with HIV and presenting with OPC.

Methods: HIV-infected patients with provisional diagnosis of OPC were consecutively enrolled between May 2017 and June 2018. After patient data collection, oral swabs and blood specimens were collected for culture and CD4 T-lymphocyte estimation, respectively. Presumptive *Candida* isolates were speciated and their antifungal susceptibilities to fluconazole, flucytosine and amphotericin B, including minimum inhibitory concentration was determined using the E-test.

Results: Of 286 patients enrolled, 67.8% (194) cultured positive for *Candida* spp. The mean age of culture positive patients was 40.7 ± 15.2 with more female enrollment (63.4%, 123/194). The CD4 counts of culture positive patients were low (211.1 ± 235.6 cells/μL) and 68.6% (133) of them were on anti-retroviral therapy (ART) with 10.3% (20/194) having previous exposure to fluconazole. Seven different *Candida* species, with the following distributions were isolated: *C. albicans* (69.1%, 134), *C. tropicalis* (10.3%, 20), *C. glabrata* (6.7%, 13), *C. parapsilosis* (5.7%, 11), *C. krusei* (4.1%, 8), *C. dubliniensis* (2.6%, 5), and *C. lusitanae* (1.5%, 3). Of all *C. albicans* isolates tested, 29.1%, 1.5% and 2.3% were resistant to fluconazole, amphotericin B and flucytosine, respectively. Non-*C. albicans* isolates showed 45%, 3.3% and 8.3% resistance to fluconazole, amphotericin B and flucytosine, respectively.

Conclusion: *C. albicans* accounted for majority of oropharyngeal candidiasis (OPC), with non-*C. albicans* showing significantly higher resistance to fluconazole. Positive culture was independent of gender, previous exposure to antifungal drugs, ART status and duration. Without any contraindication, flucytosine and Amphotericin B may be considered for OPC not responding to fluconazole therapy.

Keywords: *Candida* species, HIV, oropharyngeal candidiasis, susceptibility profile

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INTRODUCTION

Opportunistic infections (OIs) are the leading cause of morbidity and mortality among immuno-compromised patients [1,2]. By far, oropharyngeal candidiasis (OPC) dominates opportunistic fungal infections associated with HIV/AIDS. An estimated 90% or more of HIV/AIDS-patients develop OPC, with 60% having at least an episode of infection in a year and 50% - 60% recurrent infections during the course of their illness [3,4]. In Ghana, OPC has been reported as the third commonest clinical oral infection among HIV/AIDS patients [5]. Although not considered life-threatening, OPC can gradually develop into severe complications such as local discomfort, malnutrition, wasting and early death [6]. Invasive

infections can also develop following the spread of oral infection into the bloodstream causing significant morbidity and mortality [7, 8]. *Candida albicans* in many studies have been reported as the major cause of OPC in HIV-patients [9-13]. A shift in distribution has however been observed over time, with some non-*C. albicans* species implicated as opportunistic pathogens [14,15]. The clinical significance of isolating these non-*C. albicans* species is that some have been found to possess intrinsic or acquired antifungal resistance, a situation that presents significant problems with patient management [16]. These epidemiological changes in the distribution of *Candida* species underscore the need for constant monitoring to determine the burden of antifungal resistance and to recommend possible prevention and control measures. This study was therefore conducted to determine the distribution and susceptibilities of *Candida* associated oral pharyngeal infections in HIV/AIDS patients and to provide knowledge that could further impact treatment guidelines.

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MATERIAL AND METHODS

Study participants

Confirmed HIV-patients with presumptive diagnosis of OPC across all ages and gender irrespective of their antiretroviral therapy (ART) status were recruited from selected ART centers in the Central Region of Ghana from May, 2017 to June, 2018 for this study.

Sampling

Oral swabs were aseptically taken from patients who consented to the study and transported on ice pack to our laboratory. Patient data, including demographics, year of HIV diagnosis, history of previous fungal infections, history of previous antifungal use, history of ART and duration of ART prior to sampling were collected. Blood samples were also collected in ethylenediaminetetraacetic acid tubes for CD4 T-lymphocyte estimation using the BD FACS count machine (Beckton Dickinson, UK). The oral swabs were aseptically cut into 10 mL Sabouraud brain heart infusion broth and incubated at $35 \pm 2^\circ\text{C}$ for 18 to 24 h. The broth was subcultured onto Sabouraud dextrose Agar (SDA) plates supplemented with broad spectrum antibiotics and incubated for 24-48 h to obtain pure yeast isolates. *Candida* species were isolated based on their colonial morphology on the SDA together with distinguishing characteristics microscopic and Gram-stain features. Pure fresh cultures were sub-cultured onto a differential *Candida* agar plate (HICROME, India) obtained from HIMEDIA laboratories (PVT, India), and incubated aerobically at 35°C for 48 h. Species were identified based on the colour and morphology of the colonies on the differential agar plate. Definitive identification was also done using HiCandida identification kit from HIMEDIA laboratories (PVT, India).

Antifungal susceptibility testing

Yeast isolates were evaluated against three antifungals (fluconazole, 0.016-256 $\mu\text{g}/\text{mL}$; amphotericin B, 0.002-32 $\mu\text{g}/\text{mL}$; and flucytosine, 0002-32 $\mu\text{g}/\text{mL}$) using the E-test (AB, Biodisk, Sweden) susceptibility testing method as described in the Clinical and Laboratory Standard Institute (CLSI) M44-A2 document [17]. Prior to antifungal susceptibility testing, all isolates in suspension were sub-cultured onto SDA plates to ensure their purity, viability and to obtain fresh isolates. About five distinct colonies obtained from 24 h incubation at 35°C was suspended in physiological saline (0.85% NaCl) vortexed for 15 sec and adjusted to 0.5 McFarland standard (corresponds to 1×10^6 to 5×10^6 colony forming units/mL) by adding sufficient sterile saline or more colonies. The dried surface of a sterile Mueller Hinton agar with 2% glucose and 0.5 $\mu\text{g}/\text{mL}$ methylene blue dye agar plate were inoculated with the saline-yeast mixture and antifungal E-test® strips applied. The plates were incubated at 35°C for 24 h. Each plate was examined after 24 h of incubation and read only when sufficient growth was observed. Results were interpreted based on CLSI interpretive criteria (M 60, 2017) [17].

Quality control strains were set-up in similar way for every batch of isolates that was tested using strains of American Type Culture Collection (ATCC) 90028, ATCC 22019 and ATCC 6258.

Statistical analysis

Data obtained from the study were entered into Microsoft Excel 2016 and imported into IBM SPSS Software version 25 for statistical analysis. Categorical variables were summarized as frequencies and percentages, and continuous variables as means and standard deviations. Pearson Chi Square test was used to determine associations between *C. albicans*-associated OPC and categorical variables using odds ratio (OR) and 95% confidence intervals (CI). Point biserial correlation was used to determine the relationship between *C. albicans*-associated OPC and continuous variables at an alpha level of 0.05. Variables that showed significant associations in bivariate comparisons were included in logistic regression analysis to determine possible factors that may predict for *C. albicans*-associated OPC. All $p < 0.05$ were considered significant.

RESULTS

Demographic characteristics

Overall, 197 of 286 (67.8%) patients we sampled between May 2017 and June 2018 were culture positive. The mean age of these culture positive patients was 40.7 ± 15.2 yr. with majority of them being females (63.4%, 123/194). The CD4 counts of the culture positive patients were low (211.1 ± 235.6 cells/ μL) and 68.6% (133/194) of them were on ART with 10.3% (20) having previous exposure to fluconazole.

Candida species identified and their distribution

Seven different *Candida* species with the following distributions were identified from the 194 culture positive patients: *C. albicans* (69.1%, 134), *C. tropicalis* (10.3%, 20), *C. glabrata* (7%, 13), *C. parapsilosis* (5.7%, 11), *C. krusei* (4.1%, 8), *C. dubliniensis* (2.6%, 5) and *C. lusitaniae* (1.5%, 3).

Oropharyngeal candidiasis and associated factors

The demographic characteristics of the study participants were statistically not different in gender, previous exposure to antifungal drugs, duration of ART and CD4 count despite being selected from different sites. However, *C. albicans*-associated OPC was significantly higher among HIV-infected patients on ART compared to ART-naive patients ($p = 0.018$). Details of the associations are presented in Table 1.

Antifungal susceptibility

Overall resistance by *C. albicans* isolates tested against fluconazole was 29.1% (39/134). Comparatively, the non-*C. albicans* isolates were significantly more resistant to fluconazole than the *C. albicans* isolates [non-*C. albicans* = 45% (27/60) vs *C. albicans* = 29.1% (39/134); $p = 0.033$]. Overall, 1.5% (2/134) of *C. albicans* isolates tested against amphotericin B were resistant. Non-*C. albicans* had more reduced susceptibility to amphotericin B (1.5% vs 3.3%; $p = 0.27$). Overall resistance in isolates of *C. albicans* tested against flucytosine was 2.7% (3/134). The highest resistance was shown by isolates of *C. krusei* (37.5%, 3/8). None of the *C. glabrata*, *C. parapsilosis*, *C. tropicalis* and *C. dubliniensis* isolates tested against flucytosine was resistant. Tables 2 and 3 show the antifungal profile of *Candida* species and their minimum inhibitory concentrations tested against fluconazole, amphotericin B and flucytosine.

Antifungal resistance and associated factors

Fluconazole resistance was significantly higher in participants previously exposed to fluconazole compared to fluconazole-naïve patients ($p = 0.001$). No significant association was however found between antifungal use and resistance to Amphotericin B ($p = 1.000$) or flucytosine ($p = 0.925$) (Table 4). Fluconazole resistance was significantly higher among ART-naïve patients compared to patients on ART ($p = 0.001$) (Table 5). Significantly ($p = 0.004$) high fluconazole resistance was observed among patients with lower CD4 count (≤ 200 cells/ μ L) compared to patients with higher CD4 counts (>500 cells/ μ L) (Table 6).

DISCUSSION

Several research findings point to a growing problem of antifungal resistance among *Candida* species especially the non-*C. albicans*. Additionally, there seem to be an epidemiological shift in *Candida* infections, from *C. albicans* to non-*C. albicans* species [18,19]. Not only has this shift been found to differ geographically but also among health settings and even cohorts of patients within a country [13]. This underscores the need for constant monitoring of etiologic agents of *Candida* infections and emergence of antifungal resistance. *C. albicans* accounted for majority of OPC in our study participants. This is consistent with findings obtained from similar studies in Ghana and

Table 1: Logistic regression model with respect to *Candida* infection in HIV patients

Parameter	<i>C. albicans</i> (n= 134)	Non- <i>C. albicans</i> (n= 60)	p value
	OR (95% CI)	OR (95% CI)	
Gender			
Males	1.01 (0.54-1.90)	0.99 (0.53-1.87)	0.98
Antifungal histamines			
Yes	1.05 (0.38 -2.88)	0.95 (0.35-2.61)	0.92
ART			
Yes	2.16 (1.14-4.10)	0.46 (0.24-0.88)	0.02
Duration on HAART			
< 3 months	1.03 (0.40-2.66)	0.97 (0.38-2.49)	0.99
3-6 months	0.58 (0.19-1.82)	1.72 (0.55-5.36)	0.35
CD4 Count			
≤ 200	0.85 (0.28-2.58)	1.18 (0.39-3.61)	0.77
201-499	0.92 (0.27-3.07)	1.09 (0.33-3.65)	0.89

* ART, antiretroviral therapy; HAART, highly active antiretroviral therapy; OR, odds ratio; CI, confidence interval

Table 2: Antifungal susceptibility profile of HIV/AIDS patients with OPC

Parameter	Fungi Species	Susceptible n (%)	S-DD. n (%)	Intermediate n (%)	Resistant n (%)	p value
Fluconazole	<i>C. albicans</i>	89 (66.4)	6 (4.5)	-	39 (29.1)	0.001
	<i>C. krusei</i>	0 (0.0)	0 (0.0)	-	8 (100.0)	
	<i>C. parapsilosis</i>	8 (72.7)	0 (0.0)	-	3 (27.3)	
	<i>C. glabrata</i>	0 (0.0)	4 (30.8)	-	9 (69.2)	
	<i>C. lusitaniae</i>	3 (100.0)	0 (0.0)	-	0 (0.0)	
	<i>C. dubliniensis</i>	5 (100.0)	0 (0.0)	-	0 (0.0)	
	<i>C. tropicalis</i>	12 (60.0)	1 (5.0)	-	7 (35.0)	
	Amphotericin B	<i>C. albicans</i>	132 (98.5)	-	-	
<i>C. krusei</i>		7 (87.5)	-	-	1 (12.5)	
<i>C. parapsilosis</i>		11 (100.0)	-	-	0 (0.0)	
<i>C. glabrata</i>		12 (92.3)	-	-	1 (7.7)	
<i>C. lusitaniae</i>		3 (100.0)	-	-	0 (0.0)	
<i>C. dubliniensis</i>		5 (100.0)	-	-	0 (0.0)	
<i>C. tropicalis</i>		20 (100.0)	-	-	0 (0.0)	
Flucytosine		<i>C. albicans</i>	130 (97.0)	-	1 (0.7)	3 (2.3)
	<i>C. krusei</i>	5 (62.5)	-	0 (0.0)	3 (37.5)	
	<i>C. parapsilosis</i>	11 (100.0)	-	0 (0.0)	0 (0.0)	
	<i>C. glabrata</i>	13 (100.0)	-	0 (0.0)	0 (0.0)	
	<i>C. lusitaniae</i>	3 (100.0)	-	0 (0.0)	0 (0.0)	
	<i>C. dubliniensis</i>	5 (100.0)	-	0 (0.0)	0 (0.0)	
	<i>C. tropicalis</i>	18 (90.0)	-	0 (0.0)	2 (10.0)	

* S-DD, Susceptible dependent dose; p value < 0.05 implies statistically significant.

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Table 3: Minimum Inhibition Concentration of antifungals to *Candida* species

Drug	Fungi	Sensitive		Resistant		p value
		MIC Range	Mean MIC (SD)	MIC Range	Mean MIC (SD)	
Fluconazole						
	<i>C. albicans</i>	0.016-2.00	0.58 (0.62)	16.00-256.00	173.54 (75.63)	0.04
	<i>C. krusei</i>	-	-	64.00-256.00	88.00 (67.88)	
	<i>C. parapsilosis</i>	0.016-2.00	0.61 (0.87)	16.00-64.00	37.33 (24.44)	
	<i>C. glabrata</i>	-	-	96.00-256.00	149.33 (67.88)	
	<i>C. lusitaniae</i>	0.023-1.50	0.84 (0.75)	-	-	
	<i>C. dubliniensis</i>	0.016-2.00	1.04 (0.94)	-	-	
	<i>C. tropicalis</i>	0.016-1.00	0.40 (0.37)	24.00-256.00	131.43 (117.20)	
Amphotericin B						
	<i>C. albicans</i>	0.002-1.00	0.11 (0.19)	1.50-4.00	2.75 (1.77)	0.02
	<i>C. krusei</i>	0.004-1.00	0.25 (0.36)	≥6.00	16.00 (0.00)	
	<i>C. parapsilosis</i>	0.004-0.75	0.11 (0.22)	-	-	
	<i>C. glabrata</i>	0.002-1.00	0.26 (0.40)	≥32.00	32.0 (0.00)	
	<i>C. lusitaniae</i>	0.004-0.05	0.02 (0.02)	-	-	
	<i>C. dubliniensis</i>	0.004-0.064	0.02 (0.02)	-	-	
	<i>C. tropicalis</i>	0.002-0.75	0.12 (0.23)	-	-	
Flucytosine						
	<i>C. albicans</i>	0.002-3.00	0.41(0.65)	≥32.00	32.0 (0.00)	0.001
	<i>C. krusei</i>	0.016-0.75	0.26 (0.30)	≥32.00	32.0 (0.00)	
	<i>C. parapsilosis</i>	0.004-0.75	0.14 (0.23)	-	-	
	<i>C. glabrata</i>	0.002-2.00	0.26 (0.54)	-	-	
	<i>C. lusitaniae</i>	0.094-0.50	0.26 (0.21)	-	-	
	<i>C. dubliniensis</i>	0.008-1.00	0.42 (0.53)	-	-	
	<i>C. tropicalis</i>	0.006-2.00	0.51(0.67)	≥32.00	32.0 (0.00)	

*MIC, minimum inhibitory concentration; SD, standard deviation

Table 4: Association between exposure to antifungal drug and resistance

Parameter	Antifungal History n (%)		p value
	Yes (n= 20)	No (n= 174)	
Fluconazole			
Sensitive	6 (30.0)	111 (60.3)	0.001
Resistance	14 (70.0)	52 (29.9)	
S-DD	0 (0.0)	11 (6.3)	
Amphotericin			
Sensitive	20 (100.0)	170 (97.7)	1.00
Resistance	0 (0.00)	4 (2.3)	
Flucytosine			
Sensitive	19 (95.0)	166 (95.4)	0.93
Resistance	1 (5.0)	7 (4.0)	
Intermediate	0 (0.0)	1 (0.6)	

*SDD, susceptible dependent dose.

Table 5: Association between ART intake and antifungal resistance

Parameter	ART intake		p value
	Yes (n= 133)	No (n= 61)	
Fluconazole			
Sensitive	89 (66.9)	28 (45.6)	0.001
Resistance	34 (25.6)	32 (52.5)	
S-DD	10 (7.5)	1 (1.6)	
Amphotericin B			
Sensitive	130 (97.7)	60 (98.4)	1.00
Resistance	3 (2.3)	1 (1.6)	
Flucytosine			
Sensitive	127 (95.5)	58 (95.1)	0.31
Resistance	6 (4.5)	2 (3.3)	
Intermediate	0 (0.0)	1 (1.6)	

*Antiretroviral therapy; SDD, susceptible dependent dose.

elsewhere [9,10,13]. The spectrum of non-*C. albicans* species and rate of isolation was however observed to differ from those reported in previous studies. We identified seven *Candida* species, compared to twenty and eight reported in Accra and Kumasi, respectively [9,10]. The 67.8 % rate of isolation reported in this current work is also lower compared to 75.3%, and 82.3% obtained in earlier studies [9,20]. Non-*C. albicans* species constituted about 31% of all isolated cases in our study, which was consistent with 30.5% reported in Accra by Kwamin et al. (2013) [9]. In Mexico [21], Tanzania [22], and USA [23], lower rates of 16.5%, 15%, and 22%, respectively, were recorded in studies done few decades ago. In Nigeria and Brazil [13,24], non-*C. albicans* species constituted more than 50% of all isolated *Candida* species.

Increased isolation of non-*C. albicans* species has serious clinical implications and underscores the need for routine identification before treatment, especially, since majority of them have been demonstrated to be less susceptible to commonly administered antifungals [19,21]. Positive culture was found to be independent of gender, previous exposure to antifungal drugs, whether participant was on ART or not and duration of patients on ART. A study by Kwamin et al. (2013) [9] involving two groups of HIV-positive patients, those on highly active ART (HAART) and HAART-naive also reported that the difference in the prevalence of OPC among patients on HAART and HAART-naive was insignificant. *C. albicans*-associated OPC was however found to be higher among patients on ART compared to ART-naive patients. The absence of information on ART regime and level of immune suppression of the patients makes it difficult to explain this trend. However, in Southern Brazil, OPC development was found to be associated with severe immunodeficiency and high viral loads irrespective of ART use. Alcohol consumption and smoking were also found as high-risk

Table 6: Association between CD4 counts level and antifungal resistance

Parameter	CD4 count, n (%)			p value
	≤ 200 (n= 112)	20 –499 (n= 48)	≥ 500 (n= 17)	
Fluconazole				0.001
Sensitive	59 (52.7)	32 (66.7)	13 (76.5)	
Resistance	50 (44.6)	10 (20.8)	4 (23.5)	
S-DD	3 (2.7)	6 (12.5)	0 (0.0)	
Amphotericin				0.78
Sensitive	109 (97.3)	47 (97.9)	17 (0.0)	
Resistance	3 (2.7)	1 (2.1)	0 (0.0)	
Flucytosine				0.81
Sensitive	105 (93.8)	46 (95.8)	17 (100.0)	
Resistance	6 (5.4)	2 (4.2)	0 (0.0)	
Intermediate	1 (0.9)	0 (0.0)	0 (0.0)	

factors [25]. Globally, prevalence of azole resistance has been estimated in a range of 9.3 to 56.7% among HIV/AIDS patients [26–30]. This varies from samples, patients and countries in terms of incidence and prevalence rates [31].

We observed that about 30% of *C. albicans* isolates were resistance to fluconazole with significantly higher resistance found among patients whose CD4 counts were low (≤ 200 cells/ μ L). This was consistent with findings of related study where significant correlation was established between CD4 T lymphocytes and fluconazole resistance. In that study, *Candida* isolates from HIV-patients with reduced CD4 cells (< 200 cells/ μ L) were significantly resistant to fluconazole [18]. Comparatively, we also observed that the Non-*C. albicans* species were significantly more resistant to fluconazole than the *C. albicans* species. This trend could be attributed to inherent resistance exhibited by isolates of *C. krusei* and *C. glabrata* to fluconazole [30].

Both *C. albicans* and non-*C. albicans* species displayed low resistance against amphotericin B and the difference in resistance between the two was not statistically significant. The relatively low resistance prevalence of *Candida* species to amphotericin B reported in the current study falls within 0% and 3% reported by Barchiesi et al. in 2003 [32] and in 2000 [33], respectively. The results also indicate that amphotericin B is still an effective drug for treating *Candida* species infections. The reason for the low amphotericin B resistance could be because usage of the drug is restricted in the country, and so is not easily obtained over the counter. Prevalence of flucytosine resistance among *C. albicans* and non-*C. albicans* was less than 9%. With the exception of *C. krusei* which showed a high resistance (37.5%), all the other non-*C. albicans* species (*C. glabrata*, *C. parapsilosis*, *C. tropicalis* and *C. dubliniensis*) were susceptible to flucytosine. Flucytosine is rarely administered in isolation; it is combined with other antifungal agents. Besides, resistance to the drug among *Candida* species has been generally low and differ geographically [36]. Resistance of 2.3% among *C. albicans* observed in this study is higher than the 0% and 0.6% reported by Barchiesi et al. (2000) [34] and Cuenca-Estrella et al. (2001) [35] respectively and could be attributed to geographical variations or co-resistance with other antifungal drugs. The current study had limitations. CD4 count, history of previous antifungal use, anti-viral

treatment and duration are important, however this information were accurately available in a limited number of patients. We sampled patients with provisional clinical diagnosis of OPC, and may have missed a limited number of patients without symptoms. Additionally, antifungal susceptibility profile of fungal isolates from HIV patients without OPC and therefore not on any antifungal therapy would have enriched the study.

Conclusion

C. albicans was the most commonly isolated *Candida* species in OPC among the HIV-infected participants, with non-*C. albicans* species also showing a rising trend. Non-*C. albicans* species were more resistance to fluconazole, and prior fluconazole therapy and ART were associated with reduced susceptibilities to fluconazole. Without any contraindication, flucytosine and Amphotericin B may be considered for OPC not responding to fluconazole therapy.

DECLARATIONS

Ethical considerations

Ethical approval was sought from the Ethics and Protocol Review Committee (EPRC) of the College of Health Sciences, University of Ghana (CHS –Et/M.4-P2.9/2017-2018). Written informed consent was also obtained from all participants before sampling.

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None

Competing Interests

None

Author contributions

HAQ conceived the study idea. JAO and HAQ contributed to the design and implementation of the research. JAO made significant input to the analysis of the results. JAO and HAQ wrote the manuscript. All authors agreed to the content of the final draft and gave consent to publish

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Availability of data

Data is available upon request from the corresponding author


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