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An observational study on the epidemiological and mycological profile of Candidemia in ICU patients

Authors' Contribution:

- A** Study Design
- B** Data Collection
- C** Statistical Analysis
- D** Data Interpretation
- E** Manuscript Preparation
- F** Literature Search
- G** Funds Collection

Aarti Kotwal^{1ABCEDEI}, Debasis Biswas^{1ABCEDEI}, Jagdish Prasad Sharma^{2BDE},
Alpa Gupta^{3BDE}, Parul Jindal^{2BDE}

¹ Department of Microbiology, Himalayan Institute of Medical Sciences, Swami Ram Nagar, Jolly Grant, Dehradun, India

² Department of Anesthesiology, Himalayan Institute of Medical Sciences, Swami Ram Nagar, Jolly Grant, Dehradun, India

³ Department of Pediatrics, Himalayan Institute of Medical Sciences, Swami Ram Nagar, Jolly Grant, Dehradun, India

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Summary

Background:

This study aimed to analyze the epidemiological and mycological profile of candidemia in intensive care unit (ICU) patients attending a tertiary care teaching hospital in the Himalayan region of northern India.

Material/Methods:

A 15-bed medico-surgical ICU and a 5-bed pediatric ICU. Ninety-one consecutively admitted ICU patients were screened for the presence of candidemia by performing blood cultures at periodic intervals.

Results:

The recovered *Candida* isolates were speciated and subjected to antifungal susceptibility testing using standard procedures. Forty-one of the recruited patients (45%) were found to be candidemic, with the majority of patients being in the extremes of age (13 neonates and 15 >65 years of age). Four risk factors were found to be significantly associated with the occurrence of candidemia in our patients – a period of hospitalization exceeding 7 days ($p=0.0008$), previous use of antibiotics ($p=0.001$), presence of chronic renal failure ($p=0.003$), and ongoing cancer chemotherapy ($p=0.041$). Ninety-six *Candida* isolates were recovered from the 41 culture-positive patients, with *Candida albicans* being the commonest isolate recovered ($n=75$, 78.1%), followed by *Candida tropicalis* ($n=15$, 16%), and *Candida glabrata* ($n=6$, 6.5%). Fluconazole resistance was observed among 26% of all *Candida* isolates and 17.3% of *C. albicans* isolates.

Conclusions:

Contrary to the majority of recent reports, species shift towards non-*albicans* candidemia has not been observed in our center, though the prevalence of azole resistance is alarmingly high even among the *C. albicans* isolates.

key words:

candidiasis • intensive care unit • antifungal drug resistance

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Author's address:

Debasis Biswas, Department of Microbiology, Himalayan Institute of Medical Sciences, Swami Ram Nagar, Jolly Grant, Dehradun, Uttarakhand-248140, India, e-mail: dbiswas71@rediffmail.com

BACKGROUND

Candidiasis is a common cause of bloodstream and invasive infection in critically ill and immunosuppressed patients throughout the world. In addition to its widespread occurrence, it is often acutely progressive, difficult to diagnose, unresponsive to treatment and associated with increased hospital stay and high mortality rates [1–5]. Although worldwide increase in the incidence of invasive *Candida* infections has been witnessed since the 1980s [6,7], the recent trends demonstrate a gradual change in its species distribution, with many countries experiencing a relative rise in the proportion of non-*albicans* *Candida* isolates [8–16]. In view of the intrinsic resistance to specific antifungal agents observed among several of these non-*albicans* *Candida* species, this changing trend bears important therapeutic implications. Moreover, there has been a documented increase in fluconazole resistance even among previously susceptible *Candida* spp., including *Candida albicans* (*C. albicans*) [17] *C. lusitanae*, *C. tropicalis* and *C. dublimiensis* [18,19], which has been partially attributed to the widespread use of fluconazole as empirical antifungal therapy since the 1990s [20].

The increasing population of immuno-compromised patients, together with the rising incidence of non-*albicans* candidemia and the emergence of acquired antifungal resistance, necessitates the judicious administration of antifungal prophylaxis in at-risk patients and empirical antifungal therapy in patients suffering from candidemia. Sensitivity profiles of the locally prevalent *Candida* strains and knowledge regarding risk factors relevant for the patient profile attending a particular healthcare facility are key determinants in the selection of appropriate patients and antifungal agents for antifungal prophylaxis and empirical therapy. A recent survey on the importance of appropriate empirical therapy in invasive *Candida* infection has shown that adequate empirical therapy is received by only a quarter of patients and that inappropriate therapy is associated with increased mortality [21,22].

India has a high prevalence of invasive candidiasis owing to the presence of a number of contributory factors including favorable climatic conditions, a large population of immuno-compromised hosts including people with HIV/AIDS and diabetes mellitus, and widespread access to antibiotics and steroids without prescription [23]. Despite the availability of a few studies from national laboratories located in metropolitan cities [11], lack of adequate number of diagnostic mycology laboratories precludes the availability of representative data on the epidemiological and mycological characteristics of invasive candidiasis occurring in vast stretches of the country. Nevertheless, because of the immense eco-geographical heterogeneity in the country and in view of the geographical and temporal variation often observed in the species distribution of *Candida* associated with bloodstream infections [10], there is a need to investigate and monitor local epidemiological patterns of candidemia in India.

Since there is scant data on candidemia occurring in the vast Himalayan region of northern India, this prospective study was designed with the objectives of studying the local epidemiology of *Candida* infection in this region and

determining the susceptibility of the *Candida* isolates to commonly used antifungal drugs.

MATERIAL AND METHODS

The study was conducted over a period of 18 months in a 15-bed general ICU and a 5-bed paediatric ICU, attached to the Himalayan Institute of Medical Sciences, a tertiary care teaching hospital in the Himalayan region of northern India. Febrile patients with bacteriologically sterile culture reports were recruited into the study. Known HIV-positive patients and patients with pre-existing fungal infection at the time of ICU admission were excluded from the study. The study protocol was approved by the institutional ethics committee and proper informed consent was obtained from each of the recruited patients.

Five milliliters and 2 ml of venous blood samples were collected from adult and paediatric patients, respectively, at the time of admission and on the 3rd, 7th, 10th and 14th days of ICU stay. The blood samples were inoculated in brain-heart infusion biphasic media and the culture bottles were tilted at periodic intervals until the appearance of fungal colonies. The bottles were incubated for 7 days before being declared negative. Growth on BHI Agar (brain-heart infusion agar) was sub-cultured on SDA (Sabouraud's dextrose agar). All *Candida* isolates, characterized by smooth, creamy and pasty appearance of colonies on SDA, were subjected to species identification using standard tests (Germ tube test, Sugar fermentation test, Sugar assimilation test) and studying the morphological characters on corn meal agar and color production on CHROM agar media. Recovery of any *Candida* species from at least 1 blood culture sample was taken as evidence of candidemia. The recovered *Candida* isolates were then subjected to antifungal susceptibility testing using commercially procured antifungal discs (Hi-media), as per standard CLSI guidelines (document M-44A) [24] For interpretation of sensitivity or resistance, measurement of inhibition halos recommended by manufacturers was taken into consideration. The antifungal agents used in the study included Amphotericin B (10 U), Nystatin (100 U), Clotrimazole (10 µg), Fluconazole (25 µg), Ketoconazole (10 µg) and Itraconazole (10 µg). Standard ATCC strains (*C. albicans* ATCC 90028, *C. parapsilosis* 22019 and *C. krusei* 6258) were used as control. Isolates resistant to Fluconazole by disc diffusion method were tested by broth macrodilution method according to CLSI guidelines (M27 A2) [25] using Fluconazole powder procured from Sigma-Aldrich.

Statistical analysis

To examine the relative risk of developing candidemia when exposed to a particular risk factor, we calculated an odds ratio and constructed 95% confidence intervals for the same. Chi-square test and Fisher's exact test (wherever appropriate) were performed to determine if the proportion of patients developing candidemia following exposure to a risk factor was significantly different from those not developing candidemia. Differences in antifungal sensitivity between *C. albicans* and non-*albicans* species were also examined for statistical significance using chi-square test and Fisher's exact test. Controlling for the type of azole tested the relationship between the *Candida* species and azole-sensitivity was examined using Mantel-Haenszel analysis. SPSS version

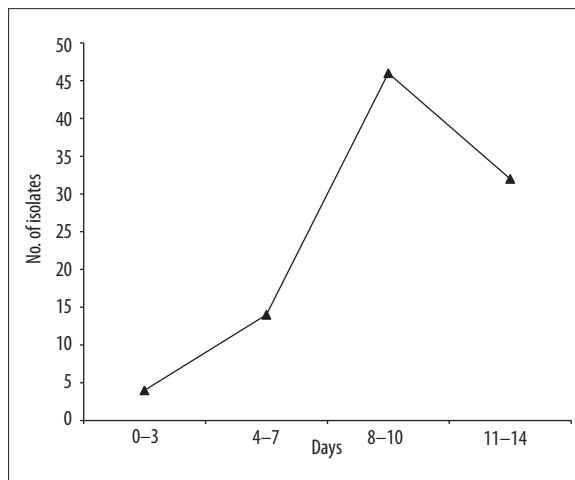


Figure 1. Time kinetics of recovery of *Candida* isolates.

17.0 was used for all statistical computations and $p < 0.05$ was taken as significant.

RESULTS

A total of 96 patients admitted in our ICUs were screened for the presence of candidemia – 32 neonates (mean age: 6 days), 8 children (mean age: 6 years), 29 adults (mean age: 36.5 years) and 27 patients above 65 years of age (mean age: 77.5 years). The mean age of the entire study group was 30 years (SD of 27 years). Of the recruited patients, 41 patients (32 males) were culture-positive, which included 23 adults and 15 children. The majority of the candidemic

patients were in the extremes of age, with 13 neonates and 15 over 65 years of age.

A total of 15 different co-morbid conditions and risk factors were identified among the recruited patients and analyzed for possible association with the development of candidemia (Table 1). Four of these risk factors were found to be significantly associated with the occurrence of candidemia in our patients – a period of hospitalization exceeding 7 days ($p=0.0008$), previous use of antibiotics ($p=0.001$), presence of chronic renal failure ($p=0.003$), and ongoing cancer chemotherapy ($p=0.041$). Among these risk factors, the odds of developing candidemia were highest for patients with chronic renal failure, with the risk being 8.3 times higher in these patients (95% confidence interval =6.9, 9.7). The odds ratios for the other significant risk factors were as follows: 5.5 for patients on antibiotics (95% confidence interval =4.5, 6.6), 5 for patients undergoing cancer chemotherapy (95% confidence interval =3.5, 6.4) and 4.6 for patients with a period of hospitalization exceeding 7 days (95% confidence interval =3.7, 5.6).

Ninety-six *Candida* isolates were recovered from the 41 culture-positive patients. The proportion of candidemic patients increased significantly beyond the 7th day of ICU stay ($p=0.0002$). The maximum number of *Candida* isolates was recovered between the 8th to 10th days of ICU stay (Figure 1). Thirty-one (75.6%) of the candidemic patients yielded multiple culture-positive samples. *Candida albicans* was the commonest isolate recovered ($n=75$, 78.1%), followed by *Candida tropicalis* ($n=15$, 16%) and *Candida glabrata* ($n=6$, 6.5%). Multiple *Candida* species were not recovered from any patient. We next analyzed whether the distribution of

Table 1. Distribution of co- morbid conditions and risk factors in the recruited patients.

Co- morbid condition/ Risk Factor	No. of patients	Candidemia present	p value	Odds Ratio	95% CI
Chronic Renal Failure	11	9	0.003	8.3	6.9, 9.7
Diabetes Mellitus	32	5	0.0002	0.14	-0.89, 1.17
Anaemia (Hb <8%)	22	5	0.04	0.32	-0.72, 1.37
Bronchial asthma	6	2	0.68	0.7	-0.83, 2.22
Cancer Chemotherapy	8	6	0.04	4.96	3.52, 6.41
Road Traffic Accident	29	6	0.005	0.24	-0.75, 1.23
Steroid Therapy	19	4	0.43	0.30	-0.8, 1.4
Low Birth Weight	15	3	0.003	0.08	-1.6, 1.86
Pre- maturity	10	3	0.24	0.38	-1.26, 2.02
Hospitalization >7 days	30	24	0.0008	4.64	3.7, 5.56
Antibiotic Usage	56	30	0.001	5.53	4.5, 6.6
IV Cannula	63	21	0.012	0.29	-0.7, 1.3
Urinary Catheter	52	9	<0.001	0.06	-1.0, 1.11
Endotracheal tube	22	2	0.0003	0.09	-1.19, 1.37
CVP line	13	3	0.149	0.375	-0.85, 1.6

Table 2. Antifungal sensitivity profile of the recovered isolates.

Isolates	Number (%)	Resistance (%)					
		Fu	Kt	It	Cc	Ns	AmpB
<i>C. albicans</i>	75 (78.1)	13 (20.6)	21 (33.3)	3 (4.8)	1 (1.6)	0	0
<i>C. tropicalis</i>	15 (15.7)	6 (50)	10 (83.3)	9 (75)	3 (25)	0	0
<i>C. glabrata</i>	6 (6.2)	6 (100)	6 (100)	6 (100)	6 (100)	0	0

Fu – Fluconazole; Kt – Ketoconazole; It – Itraconazole; Cc – Clotrimazole; Ns – Nystatin; AmpB – Amphotericin B.

risk factors was different between patients infected with *C. albicans* and those with non-albicans species. No statistically significant difference was observed between the 2 groups in terms of age, sex, length of hospitalization, administration of antibiotics or immunosuppressive agents, presence of indwelling devices and co-existence of the majority of comorbid conditions and risk factors.

Table 2 depicts the results of *in vitro* antifungal susceptibility testing performed by disc diffusion technique on the recovered isolates. In all, 26% (25 out of 96) of all *Candida* isolates and 17.3% (13 out of 75) of *C. albicans* isolates were resistant to Fluconazole. Resistance to the other azoles varied, with 38.5% of all isolates being resistant to Ketoconazole, 18.7% recording resistance to Itraconazole and 10.4% showing resistance to Clotrimazole. While the *C. glabrata* isolates were invariably resistant to every azole tested, similar resistance to all azoles was observed among 1 isolate (1.3%) of *C. albicans* and 1 isolate (6.6%) of *C. tropicalis*. For each of the azole antifungal agents, the frequency of resistance among *C. albicans* isolates was significantly lower than that among non-albicans isolates ($p=0.004$ for Fluconazole; $p<0.001$ for Ketoconazole; $p<0.001$ for Itraconazole; $p<0.001$ for Clotrimazole). Adjusting for the type of azole antifungal agent tested, resistance was significantly lower among the *C. albicans* isolates compared to the non-albicans isolates, χ^2 (1, $N=312$)=71.36, $p<0.001$. Resistance to Nystatin and Amphotericin B was, however, not observed in our study. Antifungal susceptibility testing of the resistant isolates by broth macrodilution method revealed 100% concordance with disc diffusion method. Eighteen out of 20 isolates subjected to broth macrodilution method had MIC ≥ 64 $\mu\text{g/ml}$ to Fluconazole.

DISCUSSION

In this paper we have shown that *C. albicans* is the predominant cause of candidemia in our centre and the degree of resistance is significantly high even among the *C. albicans* isolates recovered in this study. We also identified the comorbid conditions and risk factors associated with the development of candidemia in our patients. Interestingly, chronic renal failure was found to be the most important risk factor in this study.

Geographical variation is recognized to be an important feature in the species distribution of *Candida*. In sync with trends observed in the majority of studies from around the globe [10–16], a shift in the species distribution of *Candida* has been noted in several major Indian hospitals. Non-albicans *Candida* have been isolated from 30%-90% of cases

of invasive candidiasis [12–16,26,27]. However, in our study *C. albicans* was isolated as the most prevalent isolate with no shift from albicans to non-albicans candidemia. We hypothesize that, as suggested by Bassetti et al. [28], this could be accounted for by the fact that prophylactic use of fluconazole is not a standard practice in our ICU and none of the patients recruited in this study were receiving antifungal prophylaxis. Similar to our findings, Narain et al also reported predominant isolation of *C. albicans* in their study from Mumbai, India [29]. Our hypothesis is further supported by similarity of findings in reports from countries with restricted usage of antifungal agents. In a retrospective evaluation of candidemia spanning a period of 6 years, in 5 university hospitals in the Netherlands it was found that the proportion of bloodstream infections caused by *Candida albicans* remained stable throughout the study period, with no signs of increasing rate of infections due to non-albicans *Candida* [30]. Likewise, in a Swiss study no shift was observed from *C. albicans* fungemia to those caused by non-albicans *Candida* [31,32].

C. tropicalis was the most frequently recovered non-albicans isolate in our study, while *C. glabrata* is the commonest non-albicans species worldwide [27,33–36]. This is in agreement with previous reports from India, Singapore and Taiwan where *C. tropicalis* has been reported to be the commonest non-albicans *Candida* isolated [23,37,38].

A large number of risk factors have been incriminated in the development of candidemia in studies from across the world [30,39–45]. The variation in these risk factors between studies is reflective of the recruited patient profile and the nature of treatment practices and therapeutic interventions observed in the reporting institutions. Knowledge of these risk factors is helpful in adopting centre-specific strategies for selective administration of antifungal prophylaxis. The differences in risk factors observed by us and those reported by other authors could be due to the fact that procedures like organ transplantation and bone marrow transplantation are not performed in our center. Patients with hematological malignancies and neutropenic patients and procedures like total body irradiation, central venous catheterization, arterial line insertion, and cardiothoracic surgery are also fewer compared to those reported in most other studies.

All *C. glabrata* isolates recovered in our study were azole resistant. This is similar to data published in recent years in which azole resistance has been found to be higher among *C. glabrata*. In a Scottish study, among the isolates of candidemia 55% of *C. glabrata* isolates showed reduced susceptibility to fluconazole, but azole resistance among other

species of *Candida* was extremely low [40,38]. Similarly, Tan et al observed relatively higher fluconazole resistance among *C. glabrata* isolates [37]. The outstanding feature in the present study was the alarmingly high (17.3%) prevalence of fluconazole-resistance among the *C. albicans* isolates. This is in contrast to the findings in most of the studies that have reported 0–5.1% fluconazole resistance in *C. albicans* [37,46–48]. The reason for this difference remains unknown; this could be an interesting regional characteristic if this finding is validated in future studies. Work is currently underway in our laboratory wherein we are trying to corroborate these findings by blinded performance of broth dilution and disc diffusion in all our *Candida* isolates.

Unlike antibacterial susceptibility testing, antifungal susceptibility testing is not performed routinely and an empirical approach is usually followed in prescribing this class of drugs. The finding of 26% of fluconazole resistance among the *Candida* isolates in the present study underscores the importance of correct speciation and routine testing of antifungal susceptibility. The newer antifungals like voriconazole, posaconazole, and caspofungin, which have not been included in the present study, should also be tested for efficacy against the resistant isolates. Moreover, correlation needs to be explored between the results of antifungal sensitivity testing and host response to antifungal treatment.

CONCLUSIONS

This observational study, aimed at characterizing the profile of candidemia in the setting of a typical Indian ICU, could assist in alerting clinicians about the prevalence of this condition and in promoting adoption of important prophylactic and treatment guidelines for its improved management.

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