

Biofilm Production and Antifungal Susceptibility Patterns of Candida Species in Hospitalized Patients in Tertiary Care Hospital: Eastern OdishaAnsuman Dash¹, Dipti Pattnaik², Nirmala Poddar³, Madhushree Mishra⁴¹Professor, Department of Microbiology, Hi-Tech Medical College and Hospital, Rourkela, Odisha, India²Professor, Department of Microbiology, Kalinga Institute of Medical Sciences, Bhubaneswar, Odisha, India³Professor, Department of Microbiology, Kalinga Institute of Medical Sciences, Bhubaneswar, Odisha, India⁴Assistant Professor, Department of Dentistry, Hi-Tech Medical College and Hospital, Rourkela, Odisha, India

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Abstract**Background:** Biofilm formation is a major virulence factor of *Candida* species and plays a crucial role in persistent infections and reduced antifungal susceptibility. The increasing prevalence of *Candida non-albicans* species with variable biofilm-forming capacity and antifungal resistance poses significant therapeutic challenges, particularly in hospitalized patients.**Aim:** The aim of the study is to contribute clinically relevant data to the evolving understanding of biofilm-associated candidiasis and to support evidence-based therapeutic strategies in hospital settings.**Materials & Methods:** A prospective laboratory-based study was conducted in a tertiary care teaching hospital in eastern India from November 2013 to September 2015. A total of 400 clinical samples were processed, yielding 150 culture-positive *Candida* isolates. Species identification was performed using conventional phenotypic methods, chromogenic media, biochemical tests, and automated identification systems. Biofilm production was assessed using Congo red agar and tube methods. Antifungal susceptibility testing was carried out by the Kirby–Bauer disc diffusion method using polyenes and azole antifungal agents. Data were analyzed descriptively and expressed as frequencies and percentages.**Results:** *Candida non-albicans* species constituted 66% of isolates. Biofilm production was detected in 73 (48.7%) isolates by Congo red agar and in 132 (88%) isolates by the tube method. Strong biofilm formation (3+) was observed in 27.3% of isolates, with *Candida non-albicans* species demonstrating slightly higher overall biofilm positivity than *Candida albicans*. Antifungal susceptibility testing showed highest sensitivity to nystatin (89%) and amphotericin B (88%), whereas overall susceptibility to azole antifungals was comparatively low (34%). Voriconazole and clotrimazole were the most effective agents among azoles.**Conclusion:** The study highlights a high prevalence of biofilm-forming *Candida* species, particularly among non-albicans isolates, associated with reduced azole susceptibility. Routine assessment of biofilm production and antifungal susceptibility is essential to guide effective therapy and strengthen antifungal stewardship in hospital settings.**Keywords:** Biofilm formation; *Candida* species; *Candida non-albicans*; Antifungal susceptibility; Azole resistance; Hospital-acquired candidiasis.

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This is an Open Access article that uses a funding model which does not charge readers or their institutions for access and distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0>) and the Budapest Open Access Initiative (<http://www.budapestopenaccessinitiative.org/read>), which permit unrestricted use, distribution, and reproduction in any medium, provided original work is properly credited.**Introduction**

Candidiasis remains one of the most important opportunistic fungal infections worldwide, contributing substantially to morbidity and mortality among hospitalized and critically ill patients. Species of the genus *Candida* are leading causes of healthcare-associated fungal infections, particularly in intensive care units (ICUs), where invasive procedures, indwelling medical devices,

broad-spectrum antibiotic exposure, and host immunosuppression converge to facilitate infection [1,2]. In recent years, the global epidemiology of candidiasis has undergone a marked transition, with *Candida non-albicans* (NCA) species increasingly replacing *Candida albicans* as predominant clinical pathogens in many regions [3–5]. Beyond species distribution, biofilm formation has emerged as a

critical virulence determinant in *Candida* infections. Biofilms are structured microbial communities embedded within an extracellular matrix that adhere to biotic and abiotic surfaces, including vascular catheters, urinary catheters, prosthetic devices, and mucosal tissues [2,6]. *Candida* biofilms confer significant survival advantages, including enhanced resistance to host immune responses and markedly reduced susceptibility to antifungal agents compared with planktonic cells [1,6]. Consequently, biofilm-associated candidiasis is often persistent, difficult to eradicate, and associated with poor clinical outcomes.

Experimental and clinical studies have demonstrated that both *C. albicans* and NCA species are capable of biofilm formation; however, species-specific differences in biofilm architecture, metabolic activity, and resistance profiles have been described [4,7]. Notably, *Candida tropicalis*, which has emerged as a dominant NCA species in many Asian countries, is reported to form dense and metabolically active biofilms, often associated with high levels of antifungal tolerance [4,8]. These characteristics raise significant therapeutic concerns, particularly in regions where *C. tropicalis* predominates in bloodstream and device-associated infections.

Antifungal resistance further compounds the challenge of managing biofilm-associated candidiasis. Numerous surveillance studies have reported increasing resistance to azole antifungal agents among NCA species, while biofilm growth itself independently reduces antifungal efficacy through multiple mechanisms, including limited drug penetration, sequestration by extracellular matrix components, efflux pump overexpression, and persistence of metabolically quiescent cells [2,6,9]. Although polyenes such as amphotericin B and topical agents like nystatin often retain activity against *Candida* biofilms, their toxicity profiles and limited systemic applicability restrict widespread use [7,10]. The interplay between biofilm formation and antifungal susceptibility therefore has direct implications for therapeutic decision-making.

Indian and international studies have consistently highlighted regional variation in both species distribution and antifungal resistance patterns, emphasizing the need for locally generated data [8,11,12]. In many low- and middle-income countries, including India, routine laboratory surveillance of biofilm formation and antifungal susceptibility remains limited, and empirical therapy is often guided by historical assumptions rather than current epidemiological evidence [3,11]. This gap is particularly concerning given the rising burden of diabetes, prolonged hospitalization, and device-associated infections,

all of which predispose to biofilm-mediated candidiasis [13].

Understanding the relationship between biofilm-forming capacity and antifungal susceptibility across different *Candida* species is therefore essential for improving clinical management, optimizing antifungal stewardship, and guiding infection control strategies. Recent translational research has also emphasized the potential of antibiofilm-targeted therapies and combination antifungal regimens, underscoring the importance of baseline epidemiological and phenotypic data to inform future interventions [6,14,15].

Against this background, the present study was undertaken to evaluate biofilm production among clinical *Candida* isolates and to analyze their antifungal susceptibility patterns in a tertiary care setting.

By correlating biofilm-forming ability with species distribution and antifungal response, this study aims to contribute clinically relevant data to the evolving understanding of biofilm-associated candidiasis and to support evidence-based therapeutic strategies in hospital settings.

Materials & Methods

A prospective, laboratory-based observational study was conducted in the Department of Microbiology of a tertiary care teaching hospital in eastern India over a 23-month period (November 2013 to September 2015) at Hi-Tech Medical College & Hospital, Odisha. The study aimed to evaluate biofilm-forming ability and antifungal susceptibility patterns of *Candida* species isolated from various clinical samples. Ethical approval was obtained from the Institutional Ethics Committee prior to commencement of the study.

A total of 400 clinical specimens were collected from patients attending outpatient departments and from those admitted to hospital wards and intensive care units (ICUs). Specimens included urine, blood, sputum, pus, endotracheal aspirates, catheter tips, other body fluids, vaginal swabs, and throat swabs. Samples representing normal commensal flora were excluded. Of the processed specimens, 150 culture-positive *Candida* isolates were included for further analysis.

1. Isolation and identification of *Candida* species:

All specimens were processed using standard mycological techniques. Samples were inoculated on Blood agar, MacConkey agar, cystine–lactose–electrolyte-deficient (CLED) agar, and Sabouraud's dextrose agar (SDA) supplemented with antibiotics, and incubated at 37 °C for 24–48 hours.

Blood specimens were additionally processed using an automated blood culture system (BacT/ALERT

3D, bioMérieux), with subcultures performed for up to 10 days. Presumptive *Candida* isolates were identified by colony morphology, Gram staining, and lactophenol cotton blue mount. Species-level identification was carried out using germ tube test, chlamydospore production on corn meal agar (Dalmau plate technique), chromogenic medium (CHROMagar *Candida*), sugar assimilation tests, and confirmation by an automated identification system (VITEK, bioMérieux) where applicable.

2. Antifungal susceptibility testing: Antifungal susceptibility testing was performed for all *Candida* isolates using the Kirby–Bauer disc diffusion method on Mueller–Hinton agar supplemented with 2% glucose and methylene blue, in accordance with standard guidelines. Inoculum suspension was prepared by emulsifying 2–4 colonies in sterile saline to achieve uniform turbidity. Antifungal discs tested included amphotericin B (20 µg), fluconazole (10 µg), itraconazole (30 µg), ketoconazole (10 µg), voriconazole (1 µg), clotrimazole (10 µg), and nystatin (100 µg). Plates were incubated at 35–37 °C for 24–48 hours, and zones of inhibition were measured and interpreted as sensitive or resistant according to standard interpretative criteria.

3. Assessment of biofilm production: Biofilm formation was evaluated using two phenotypic

methods: Congo red agar (CRA) method: Isolates were inoculated onto brain heart infusion agar supplemented with glucose and Congo red dye and incubated aerobically at 37 °C for 24–48 hours. Biofilm production was interpreted based on colony colour: black or dark red colonies indicated strong biofilm production, pink colonies indicated weak biofilm production, and white or pale colonies were considered non-biofilm producers.

Tube method: A loopful of growth from SDA was inoculated into Sabouraud's dextrose broth supplemented with 8% glucose and incubated at 37 °C for 24 hours. After incubation, the broth was discarded, and the tubes were washed and stained with 1% safranin. Biofilm formation was assessed visually and graded as negative (0), weak (1+), moderate (2+), or strong (3+).

Statistical analysis: Data were entered into a spreadsheet and analyzed using standard statistical software. Results were expressed as frequencies and percentages. Comparative analysis of biofilm production and antifungal susceptibility between *Candida albicans* and *Candida non-albicans* species was performed descriptively.

Results:

Table 1: Antifungal Susceptibility Pattern

Species	Isolates	Amphotericin B	Azoles	Nystatin
<i>C. albicans</i>	51	45 (88%)	18 (35.3%)	46 (90%)
<i>C. tropicalis</i>	61	56 (92%)	17 (27.8%)	57 (93.4%)
<i>C. glabrata</i>	15	13 (86.6%)	9 (60%)	12 (80%)
<i>C. parapsilosis</i>	8	8 (100%)	2 (25%)	5 (62.5%)

Table 2: Azole Sensitivity Pattern

Species	FLC	IT	KT	VRC	CC
<i>C. albicans</i>	42	34	36	45	47
<i>C. tropicalis</i>	39	42	55	50	53
<i>C. glabrata</i>	14	12	15	13	13

Table 3: Biofilm Production (Tube Method)

Species	3+	2+	1+	0	Total
<i>C. albicans</i>	2	23	19	7	51
Non-albicans	39	41	8	11	99

Antifungal susceptibility patterns of the *Candida* isolates are summarized in Table 1. Overall, Nystatin exhibited the highest susceptibility rate (89%), followed closely by Amphotericin B (88%). Sensitivity to azole antifungal agents was comparatively lower (34%). *Candida tropicalis* demonstrated the highest sensitivity to Amphotericin B (92%) and Nystatin (93.4%), while *C. albicans* showed 90% sensitivity to Nystatin. Notably, *C. haemulonii* isolates exhibited complete resistance to Amphotericin B and azoles but remained sensitive to Nystatin.

A detailed analysis of azole susceptibility patterns is presented in Table 2. Among the azole group, Voriconazole showed the highest overall activity (86%), followed by Clotrimazole (85.3%). Fluconazole and Ketoconazole demonstrated moderate effectiveness (74%), whereas Itraconazole showed the lowest susceptibility (70%). Species-wise analysis revealed that *C. albicans* was most sensitive to Clotrimazole, *C. tropicalis* to Ketoconazole, and *C. glabrata* exhibited relatively better susceptibility across multiple azoles compared to other NAC species.

Biofilm-forming ability of *Candida* isolates assessed by the tube method is depicted in Table 3. Strong biofilm production (3+) was observed in 27.3% of isolates, while moderate (2+) and weak (1+) biofilm formation were seen in 42.6% and 18% of isolates, respectively. *Candida non-albicans* species demonstrated a higher overall rate of biofilm production (89%) compared to *C. albicans* (86.2%). These findings suggest a greater virulence potential of NAC species, which may contribute to persistent infections and reduced antifungal susceptibility.

Discussion

This study evaluated biofilm production and antifungal susceptibility among 150 *Candida* isolates from clinical specimens, revealing that robust biofilm formation (3+) occurred in 27.3% of isolates and that *Candida non-albicans* (NCA) exhibited slightly higher overall biofilm production than *C. albicans* (89% vs 86.2%), while polyenes (amphotericin B, nystatin) retained the highest in-vitro activity and azole susceptibility was comparatively lower (Tables 1–3). These findings align with a growing body of evidence that links biofilm formation with reduced antifungal susceptibility and clinical persistence of *Candida* infections [1–3]. A recent meta-analysis reported high rates of biofilm formation among bloodstream *Candida* isolates and strong associations between biofilm phenotype and resistance to fluconazole, voriconazole and echinocandins, supporting our observation of biofilm-associated therapeutic challenges [1].

Species-specific tendencies observed here—particularly the predominance of biofilm-forming NAC such as *C. tropicalis*—are corroborated by experimental and clinical studies that show *C. tropicalis* frequently produces dense biofilms with high extracellular matrix content, leading to elevated antifungal tolerance [4,5]. In vitro life-cycle analyse or demonstrates that *C. tropicalis* biofilms mature rapidly and achieve high biomass, a feature that is associated with resistance phenotypes in clinical isolates [4]. Mechanistically, biofilm resistance is multifactorial, encompassing limited drug penetration, extracellular matrix sequestration, quorum-sensing mediated phenotypic shifts, upregulation of efflux pumps, and persist cell formation, as extensively reviewed in recent literature [2,6].

The antifungal susceptibility profile in our dataset showed high activity of nystatin (89%) and amphotericin B (88%), with much lower aggregate susceptibility to azoles (34%) (Table 1). Comparable surveillance studies from diverse settings report similar patterns, with polyenes often retaining activity when azole resistance increases among NCA species [7–9].

For example, multi-centre and single-centre Indian studies documented substantial fluconazole resistance among *C. tropicalis* and *C. glabrata*, while amphotericin B largely preserved activity in vitro [8,9]. These parallels emphasize that empirical azole therapy may be unreliable against biofilm-forming NAC in high-prevalence regions.

The clinical implications of biofilm-associated resistance are substantial. Devices such as urinary and central venous catheters are frequent niduses for biofilm establishment, and biofilm-embedded cells can seed bloodstream infections that are difficult to eradicate without device removal [1,10]. Our finding that urinary specimens were a major source of isolates in the parent dataset underscores the clinical relevance of catheter-associated biofilms in the studied population (see dataset).

Several translational studies spotlight potential strategies to overcome biofilm tolerance. Adjunctive approaches—combining antifungals with biofilm-disrupting agents, antifungal peptides, or novel small molecules—have shown promise in vitro and in animal models [11–13]. Recent reviews highlight that targeting and efflux mechanisms can sensitize biofilms to conventional agents [2,11]. Emerging antibiofilm agents, including antimicrobial peptides, lipopeptides, and nanoparticle formulations, have demonstrated eradication or disruption of mature *C. tropicalis* biofilms in preclinical studies, suggesting plausible future adjuncts to therapy [14,15]. Nevertheless, regional heterogeneity in species distribution and resistance complicates generalization. Some reports show higher echinocandin resistance or emerging polyene non-susceptibility in particular locales, highlighting the need for continuous local surveillance [7,9,16]. Our data thus contribute a timely regional snapshot but must be interpreted alongside contemporaneous surveillance to guide empirical therapy.

Several limitations warrant mention. First, biofilm assessment employed phenotypic methods (Congo Red Agar and tube method) rather than quantitative microtiter plate biomass/viability assays or confocal imaging; phenotypic scoring is semi-quantitative and may lack granularity. Second, antifungal susceptibility was assessed by disc diffusion rather than broth microdilution (CLSI/EUCAST MIC determination), which is the reference standard for correlation with clinical breakpoints.

Third, molecular characterization of resistance mechanisms (e.g., ERG11 mutations, efflux pump gene expression) and sequencing-based species confirmation were not performed for all isolates, limiting mechanistic inference. Fourth, the data derive from a single tertiary centre and a historical timeframe, which may not reflect evolving post-

COVID ICU epidemiology or emergent species such as *Candida auris*.

Future recommendations: To address these gaps, future studies should employ standardized quantitative biofilm assays (e.g., XTT/Alamar Blue, CFU enumeration, confocal microscopy) and MIC-based antifungal testing (broth microdilution) to better correlate biofilm biomass with antifungal susceptibility. Integrating molecular analyses including genotyping, resistance gene sequencing, and transcriptomics will illuminate mechanisms underpinning biofilm tolerance and resistance.

Multicentre prospective surveillance across geographic regions, including ICU networks and post-COVID cohorts, is needed to map species dynamics and resistance trends. Finally, clinical trials testing adjunctive antibiofilm strategies (matrix-degrading enzymes, AMPs, lipopeptides, or combination regimens) are warranted to translate *in vitro* promise into improved patient outcomes.

Conclusion

This study reinforces that biofilm formation is common among clinical *Candida* isolates particularly NAC species—and is associated with diminished azole susceptibility and reliance on polyenes *in vitro*. These findings underscore the urgent need for integrated diagnostic, stewardship, and therapeutic strategies targeting biofilm-associated candidiasis in hospital settings.

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