



INCIDENCE OF BIOFILM FORMATION IN VARIOUS CANDIDA SPECIES ISOLATED FROM CLINICAL SAMPLES AT GOVERNMENT MEDICAL COLLEGE, KOTA.

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Abstract

Candida species are typical skin and mucosal dwellers. Given the rise in these infections during the past ten years, the significance of epidemiological monitoring of yeasts implicated in pathogenic processes is undeniable. Clinical samples were obtained and cultured from the respiratory tract (sputum, bronchial wash, tracheal secretions), saliva, blood, urine, middle ear discharge, vitreous fluid, corneal ulcer, and plastic devices (endotracheal tube, catheter tip, suction tip). The isolated species of Candida were identified. From 270 different clinical sources, 100 isolates of Candida species were found. The most isolated species was Candida albicans (39.5 %), however the most common species was Candida non-albicans (60.5%). C. albicans (20.58%) and C. krusei (38.23%) were commonly isolated in blood cultures. The most common species on the mucosal surface was Candida albicans (63.27%). Hospitalized patients were more likely to have urinary tract infections brought on by yeasts, with C. krusei (50.0%) and C. albicans (25.0%) being the most often isolated. The pathogenicity is influenced by a number of virulence factors, including biofilm, proteinase, phospholipase, etc. Clinical decision-making benefits from early discovery of Candida virulence factors. Therefore, our goal has been to show how biofilm forms using the technique suggested by Branchini et al. (1994). The Staib et al. (1965) approach was used to determine the amount of proteinase generated by Candida. The phospholipase assay was performed using the Samaranayake et al. (2005) method. Conclusions: The findings points to the possibility that the pathogenic potential of the isolates may be reflected in the ability of Candida species to form biofilm. Strong slime production was demonstrated by C. tropicalis and C. krusei. Compared to Candida albicans, the non-Candida albicans produced more proteinase. In this investigation, Candida albicans produced more phospholipase than non-Candida albicans.

Keywords-Candida species, biofilm, Infection, fungi

Introduction

The skin and mucosa are typically home to Candida species. Given the rise in these infections over the past ten years, as well as the changes in species causing candidiasis and empirical antifungal treatment, the significance of epidemiological monitoring of yeasts implicated in pathogenic processes is undeniable.[1] Other Candida species have also become clinically significant

opportunistic pathogens, even though *Candida albicans* is the organism most frequently linked to serious fungal infections. To aid in colonization, invasion, and pathogenesis, the majority of pathogens, including *Candida* species, have evolved a potent arsenal of suspected virulence factors and particular tactics. The type of infection, the location and stage of the infection, and the type of host response can all affect the virulence factors that *Candida* species express to cause infections. The generation of acid proteinase, phospholipase, and biofilm formation are the primary virulence agents. While phenotypic switching or platelet coating may be employed to avoid the immune system, enzymes aid adherence once contact is established by breaking down cell membranes and extracellular proteins, allowing the yeast to penetrate the host. The development of clinical infection depends on biofilms, which are organized microbial populations connected to and enclosed in an exopolymeric material matrix [2]. A crucial part of the pathophysiology of candidiasis is played by the outermost layers of *Candida* cells, which are necessary for adhesion to the host surface.[3] Protection from the environment, nutrition availability, metabolic cooperation, and the acquisition of new genetic features are all benefits of creating a biofilm. During infection, pathogenic species of *Candida* release aspartyl proteinases in vivo.[4] When the organism is cultivated with foreign protein—typically bovine serum albumin—as the nitrogen source, the enzymes are secreted in vitro. The synthesis of proteases is thought to improve the organism's capacity to infiltrate and colonize host tissues as well as to elude the host immune system.[5] Phospholipase enzymes are linked to adhesion, penetration, and host cell membrane degradation. When microorganisms invade host cells, the outer cell membrane is penetrated and damaged. According to preliminary evidence, the primary processes for fungal virulence appear to be direct host cell injury and lysis.

MATERIALS AND METHODS

Patients receiving care in Kota government hospital and assisted living facilities provided 270 distinct clinical samples. Before collection, the patients had never been exposed to antifungal medications. 112 samples were taken from the respiratory tract (sputum, bronchial wash, tracheal secretion) and saliva, 130 from blood, 12 from urine, 2 from middle ear discharge, 1 from vitreous fluid, 1 from corneal ulcer, and 9 from plastic devices (endotracheal tube, catheter tip, suction tip).

In 10% KOH, every respiratory specimen and exudate was analyzed. The smears were also analyzed and stained with gram stain. As the primary isolation medium, Sabouraud's dextrose agar was used to inoculate the samples. SDA biphasic medium containing gentamicin and chloramphenicol was utilized for blood samples. For a week, or longer if necessary, the culture media was incubated at 37°C.

By evaluating the production of germ tubes, pellicles, assimilation, and sugar fermentation, the species was identified. To demonstrate the presence of chlamydospores, they were cultivated on cornmeal agar. The species was identified via culture on candid chrom agar.

Using a technique suggested by Pfaller MA et al., biofilm development was assessed for each isolate as well as the standard strains. [6] Ten milliliters of Sabouraud's liquid medium supplemented with glucose (final concentration of 8%) were added to a tube containing a loopful of organisms from the SDA plate. Following a 24-hour incubation period at 37°C, the tubes' walls were stained with safranin and the broth was aspirated out. A score of zero (0+), weak positive (1+), moderate positive (2+), or strong positive (3+) was assigned to biofilm formation.

The significantly modified Hendry AT et al. method [7] was used to detect the *Candida* proteinase using bovine serum albumin medium (dextrose 2%, KH₂PO₄ 0.1%, MgSO₄ 0.05%, agar 2% mixed after cooling to 50°C with 1% bovine serum albumin solution). Inoculating 10 µl aliquots of the yeast suspension (about 10⁸ yeast cells/ml) into the wells punched onto the medium's surface allowed for the detection of proteinase activity. For two days, the plates were incubated at 37°C. Following incubation, the plates were dyed with 1.25% amidoblack and fixed with 20% trichloroacetic acid. 15% acetic acid was used for decolorization. Degradation of the protein was shown by opaqueness of the agar, which corresponded to a zone of proteolysis surrounding the wells that could not be stained with amidoblack. Proteinase production was measured using the diameter of the unstained zones

surrounding the well. The ratio of the well's diameter to the proteolytic unstained zone's diameter was used to calculate the proteinase activity (Pz). There was no proteinase activity in the strain when $Pz = 1$. Therefore, low Pz indicates high enzyme synthesis. Phospholipase was estimated using a slightly modified version of Samaranayake et al.'s approach [8]. 13.0 g Sabouraud dextrose agar (SDA), 11.7 g NaCl, 0.111 g $CaCl_2$, and 10% sterile egg yolk made up the egg yolk medium. Twenty milliliters of the supernatant were added to the sterilized medium after the egg yolk was centrifuged at 500 g for ten minutes at room temperature. By inoculating 10 μ l aliquots of the yeast suspension (about 108 yeast cells/ml) into the wells punched onto the surface of the egg yolk medium, extracellular phospholipase activity was found. Following a 48-hour incubation period at 37°C, the diameter of the precipitation zone surrounding the well was measured. The Pz value, or phospholipase activity, was calculated. There was no phospholipase activity in the strain when $Pz = 1$. Therefore, low Pz indicates high enzyme synthesis.

Results

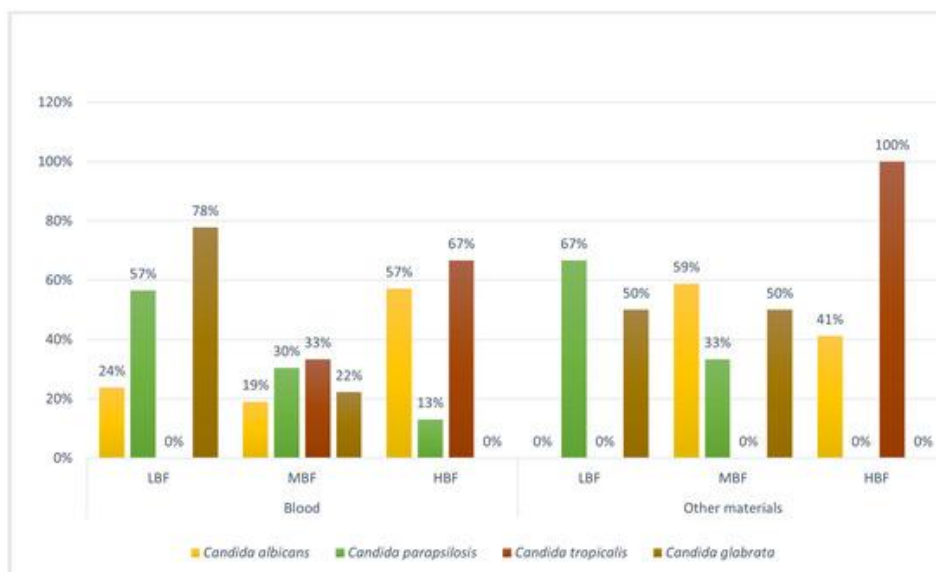
Of the 100 isolates, 39 were *Candida albicans*, 7 were *Candida glabrata*, 4 were *Candida guilliermondii*, 2 were *Candida kefyr*, 31 were *Candida krusei*, 4 were *Candida parapsilosis*, and 8 were *Candida tropicalis*. Table 1 displays the distribution of *Candida* species in various clinical samples.

Out of the 100 *Candida* species isolates that were recovered from the clinical isolates, 73 (73%) of them formed biofilm. Compared to the percentage of all non-*albicans* *Candida* species isolates that generated slime (90.32%, 55 of 61; $P < 0.0001$), only 51% (20 of 39) of *C. albicans* isolates produced biofilm. *C. tropicalis* and *C. krusei* exhibited strong biofilm formation. *C. albicans* was found to produce weak biofilms. In 89 (80.18%) of the isolates, proteinase activity was found. *C. albicans* (Pz 0.18), *C. kefyr* (Pz 0.16), and *C. guilliermondii* (Pz 0.17) produced the most proteinase, whereas *C. glabrata* (Pz 0.29) produced the least. Of the isolates, 49 (44.14%) had phospholipase activity. *C. guilliermondii* produces the most phospholipase (Pz 0.07), followed by *C. parapsilosis* (Pz 0.08). *C. tropicalis* is the least productive (Pz 0.27).

Figure 1 illustrates the generation of biofilm, proteinase, and phospholipase by *Candida* species isolated from clinical specimens.

Table 1: *Candida* species isolated from different clinical samples

Source of clinical isolates	Respiratory tract	Blood	Urine	Plastic devices	Eye	Middle ear discharge	Pus	Total
Positive isolates	42	30	12	9	2	2	3	100
<i>C. albicans</i>	27	6	3	3	0	0	0	39
<i>C. krusei</i>	9	11	6	3	1	0	1	31
<i>C. tropicalis</i>	3	2	1	1	0	0	1	8
<i>C. parapsilosis</i>	0	3	1	0	0	0	0	4
<i>C. guilliermondii</i>	1	2	0	0	0	1	0	4
<i>C. pseudotropicalis</i>	0	2	0	0	0	0	0	2
<i>C. glabrata</i>	0	4	1	0	1	1	0	7
<i>C. stellatoidea</i>	2	0	0	2	0	0	1	5



Percentage values of tertile groupings of high, medium, and low biofilm-producing strains of *Candida* spp. based on material. *C. tropicalis* and *albicans* appear to be the species with the highest number of strains in the HBF category, in contrast to *C. parapsilosis* and *glabrata*, which show high percentages of strains in the LBF category, confirming the following scale of producers: *C. tropicalis* > *C. albicans* > *C. parapsilosis* > *C. glabrata*. LBF: low biofilm forming; MBF: moderate biofilm forming; HBF: high biofilm forming.

Discussion

Humans and their surroundings are home to the asexual, diploid, dimorphic fungus *Candida*. There are comparatively few species of *Candida* that are harmful to humans. Numerous superficial and deep-seated mycoses, including cutaneous, mucocutaneous, subcutaneous, or systemic candidiasis, can be brought on by these organisms. *Candida* organisms are commensals; in order for them to function as pathogens, the host's regular defenses must be disrupted. Thus, immunocompromised conditions, diabetes mellitus, and iatrogenic factors such as antibiotic usage, indwelling devices, intravenous medication use, and hyperalimentation fluids are common risk factors for *Candida* infections. Due to an increase in individuals with impaired immune systems, advanced age, extended antimicrobial and severe cancer chemotherapy, invasive surgery, and organ transplantation, candidiasis has become a concerning opportunistic illness.

As demonstrated by Mujika et al., the current investigation demonstrated the preponderance of non-*Candida albicans* and the distribution of *Candida* species in various clinical samples. [1] *C. krusei* was the most prevalent isolate across all samples. The most common species found in respiratory tract samples was *Candida albicans* (41.37%). Patients over 60 who had productive coughs most often developed a subsequent *Candida* infection. 120 blood samples were taken from the dialysis and intensive care units. Non-*Candida albicans* were the most common species that were isolated from the blood samples. *C. krusei* was the most prevalent isolate. Microorganisms that enter the intracutaneous wound during or after catheter placement are the primary cause of the majority of catheter-related septicemias. [9–11]

The percentage of these infections caused by species other than *Candida albicans* is steadily increasing. [12–14] 84 people with diabetes who were older than 60 had their saliva samples taken. *C. albicans* was found in 23 (74.19%) and non-*Candida albicans* in 8 (25.8%) of the saliva samples from 31 patients. These patients showed symptoms of a urinary tract infection, and urine samples revealed *C. krusei* 6 (50.0%) and *C. albicans* 3 (25.0%). The majority of the isolates from the eye, middle ear discharge, and pus were not *Candida albicans* species. Patients' plastic devices, such as

catheter tips, suction tips, and endotracheal tubes, were gathered. These cultures mostly produced *Candida albicans* and *Candida krusei*.

A population of bacteria and their extracellular polymers adhered to a surface is called a biofilm. [15] A group of bacteria encased in the slime they secrete is called a biofilm. Since the capacity to create biofilms is linked to pathogenicity, it should be regarded as a significant virulence factor during candidiasis. Through their ability to evade host immune responses, endure antifungal therapy, and withstand competitive pressure from other organisms, biofilms may contribute to the maintenance of fungi's roles as commensal and pathogen. As a result, treating illnesses linked to biofilms is challenging. [16] High levels of antibiotic resistance in the related species are similarly linked to the development of biofilms. [17] Isolates of *Candida krusei* showed the highest frequency of biofilm positive, followed by *Candida tropicalis*, *Candida kefyr*, *Candida guilliermondii*, *Candida parapsilosis*, *Candida glabrata*, and *Candida albicans*. On the other hand, isolates of *Candida parapsilosis* and *Candida glabrata* were far less likely to form biofilms than the more pathogenic *Candida albicans*, according to Hawser and Douglas [18]. In this investigation, the species of *Candida* was more closely associated with biofilm development than the infection site. When the strains were categorized by the age of the patients and the infection site, there were no appreciable variations in the biofilm production. During infection, pathogenic species of *Candida* release aspartyl proteinases in vivo. The adhesion, tissue injury, and invasion of host immune responses are caused by secreted aspartic proteinases (Saps). During the infectious process, proteinases perform a variety of specialized tasks, such as breaking down molecules to obtain nutrients, breaking down or warping host cell membranes to promote adhesion and tissue invasion, and breaking down host immune system cells and molecules to prevent or fend off host antimicrobial attack. In this investigation, *Candida non albicans* 56 (50.45%) had a lower capacity to produce proteinase than *Candida albicans* 33 (67.34%). The amount of proteinase generated varied significantly between species ($P < 0.05$). A diverse collection of enzymes known as "phospholipases" are able to hydrolyze one or more ester linkages in glycerophospholipids. Direct host cell damage and lysis has been suggested as a key mechanism contributing to microbial pathogenicity because phospholipase targets membrane phospholipids and digests these components, resulting in cell lysis [19]. Phospholipase was generated by 23 (46.93%) of the *C. albicans* isolates and 26 (42%) of the non-*Candida albicans* isolates. In demonstrating that *C. albicans* isolated from the blood samples had higher extracellular phospholipase activity, the study's findings are consistent with those of Ibrahim et al. [20].

CONCLUSION

The frequency of invasive, often life threatening *Candida* infections has increased dramatically largely as a function of medical technology advances. Throughout the world, the dominance of *Candida albicans* has been challenged by the increased prevalence of serious infection caused by *Non albicans Candida* species.

In this study, there was a clear shift in prevalence of infection by *Candida albicans* to those of *Non albicans Candida* species in the various clinical samples isolated and also shows them as strong biofilm producers compared to *C. albicans* species. These data suggest that, biofilm formation as a potential virulence factor might have a higher significance for *non-albicans Candida* species than for *C. albicans* and also that the biofilm structure varies with the different species and strains of *candida*, the nature of the colonized surface and its localization. The key to successful management lies in early recognizing and diagnosis of species. Colony morphology on corn meal agar and CHROMagar are more reliable and rapid methods for speciation of *Candida* isolates. CHROMagar is superior in case of mixture of species in the same sample.

For Biofilm formation MTP method is gold standard followed by Tube method and CRA method.

So it has become imperative for us to take timely steps for early speciation and antifungal susceptibility pattern of the clinically significant *Candida* isolates prevalent in our society. This study can be further improve by identifying antifungal susceptibility pattern of *Candida* isolates so as to prevent development of resistance in *candida* species.

Further research to improve diagnostic, preventive and therapeutic strategies is necessary to reduce the considerable mortality and morbidity.

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