Assessment of Virulence Factors and Antifungal Susceptibility of Candida Species Isolated from Catheter Associated Urinary Tract Infections

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ABSTRACT
Background: Catheter associated urinary tract infection (CAUTI) is a crucial attribute to virulence of organisms. Emergency of non albicans Candida (NAC) species become common due to several virulence factors, which play an important role in pathogenicity and their resistance to treatment.

Aim of the work: To investigate different virulence factors of Candida spp. isolated from CAUTI patients and evaluate their antifungal susceptibility.

Patients and Methods: Urine samples collected from 219 indwelling catheters of ICU patients. Candida isolates identified using Germ tube test, Chrom ID agar, Corn meal tween agar, VITEK 2 system and Integral System Yeast Plus (ISYP) used to detect antifungal susceptibility. Enzymatic, haemolytic activity and biofilm formation assessed by standard methods.

Results: The incidence of candiduria was 50/219 (22.8%). Candida tropicalis was predominant one (54.0%) followed by C. albicans (26.0%) and C. glabrata (20.0%). Females gender (64.0%), Diabetes (50.0%) associated with positive candiduria, while antibiotic therapy was predominant risk factor (78.0%). C. albicans have highest enzymatic activity for phospholipase (84.6 %), proteinase (84.6%) and haemolytic activity (100%), while C. tropicalis have highest biofilm formation (100%). NAC spp. showed higher rates of resistance to azole drugs (Clotrimazole 77.6%, Miconazole 70.3%, Econazole 63.2%, Nystatine 44.3%), while C. albicans didn’t show any resistance.

Conclusion: Candiduria is prevalent among catheterized patients with shift toward NAC spp. causing nosocomial UTI. A strong relationship between host risk factors (old age, antibiotic use, catheterization, female sex, ICU stay, diabetes) and expression of various virulence factors of Candida spp. causing candiduria and resistance to antifungal drugs.

Keywords: Candiduria; virulence factors; antifungal resistance.

INTRODUCTION
Urinary tract infection (UTI) is one of the most common infections in clinical life. They can be caused by a range of pathogens such as bacteria, fungi, parasites, and viruses. However, among the fungal agents, and the opportunistic fungal infections involving urinary system, Candida species are the most common causes of UTI.

Candida spp. can be colonized in the lower or upper urinary tract system and create pyelonephritis, cystitis, urethritis and prostatitis in men. They might even appear in the upper urinary tract from the bloodstream or raise the urinary tract from a focus of Candida colonization at near the urethra. Although C. albicans is the most commonly isolated species in candiduria, the incidence of non albicans species is ever increasing.

Predisposing conditions such as extensive use of broad-spectrum antibiotics, diabetes mellitus, urinary stasis, renal transplantation, and hospitalization can increase the risk of candiduria. Also, urinary catheters are a crucial attribute to virulence as they allow the yeast to attach to body sites and commence proliferation and colonization. Candida species have several virulence factors that facilitate proliferation; they may result in adhesion to the epithelium and invasion of the host tissue.

The extracellular hydrolytic enzymes including secreted aspartyl proteinase and phospholipases. Furthermore, the survival and ability of Candida albicans to establish infections within humans are mainly related to its ability to produce elemental iron through haemolysin production. Moreover, it has been reported that biofilm formation also plays an essential role in the pathogenicity of Candida...
species. The aim of this work is to investigate various virulence factors as phospholipase, proteinase and haemolytic activities in Candida species isolated from patients with catheter associated urinary tract infection, assess their ability to biofilm formation and evaluate the antifungal susceptibility pattern of the isolated Candida species.

PATIENTS AND METHODS

Study population: A total of 219 patients admitted to ICU of Tanta University Hospital and undergoing urethral catheterization for more than seven days with negative urine cultures at the time of admission were investigated. They were divided into 125 female and 94 male patients. Their age ranged from 50-90 years old with mean±SD (62.45±7.93). Verbal informed consent was taken from patients to participate in the study.

Inclusion Criteria: Hospitalized ICU patients on continuous catheterization for more than 7 days with negative urine cultures at the time of admission

Exclusion Criteria: Patients with history of UTI prior to admission and patients with duration of catheterization less than 7 days.

Ethical approval: written approval was obtained from the Ethical Research Board (ERB) of the Faculty of Medicine for girls, Al-Azhar University, Cairo, Egypt.

Collection of samples: After obtaining verbal informed consent and changing the catheter about 10 ml of urine sample was collected immediately from each patient using a sterile syringe through the sample port of the indwelling catheter under complete aseptic precautions. Each sample was transferred into two sterile screw capped labelled falcon tube. The tubes containing the samples were transferred on ice directly to our microbiology laboratory where processing of samples was carried out without any delay.

Samples Processing: Ten microliters from each urine sample were plated on Sabouraud’s dextrose agar (SDA) plate supplemented with gentamicin (50 mg/L) and chloramphenicol (0.05gms/L), blood agar and CLED medium using calibrated loop. Plates were incubated aerobically at 37ºC for 24-48 hours 9. Presumptive identification of Candida isolates were done morphologically as white to creamy, round, soft and smooth wrinkled colonies with characteristic yeast odor. Microscopic examination of Gram stained film prepared from suggestive colonies showed Gram-positive budding yeast cells with or without pseudohyphae.

Colony counts (CFU) were done for all positive urine samples. All Candida isolates with CFU ≥ 10^5 were purified by a repeated streaking method on SDA to obtain pure discrete colonies, that stored in 10% glycerol (Sigma-Aldrich, South Africa) at 80 °C until further investigation.

Species Identification: Isolated yeast colonies were speciated phenotypically by germ tube test, chromogenic medium (Chrom ID Candida Agar (CAN2)), chlamydospore formation on corn meal tween 80, and sugar assimilation by Integral System Yeast Plus Commercial Kit (Liofilchem, Ref.71822 - ITALY). and confirmed by Vitek 2 system. Standard strains of C. albicans (ATCC, 10231), C. tropicalis (ATCC750) and C. glabrata (ATCC, 2001) were used as controls in all investigation.

1. Germ tube test: Fifty microliters of each yeast cell suspension with a turbidity of 1.5 x 10^3 cells/ml equivalent to 0.5 McFarland standard was inoculated in 2ml Eppendorf tubes containing aliquots of bovine serum albumin and incubated aerobically at 37ºC for three hours. Then, centrifugation of the inoculated tubes was done. A wet mount film was prepared and examined under the microscope to look for the presence of germ tube. The germ tube appeared as short lateral hyphen filament (Figure 1). There is no point of constriction at the origin of the germ tube and is classically described as hand mirror appearance 11. C. albicans and C dubliniensis produce germ tubes. C. tropicalis may produce early pesudohyphae that are constricted at the point of origin with the yeast cell in contrast to germ tube (Figure 1). Other candida species give no hyphae or germ tube.

2. Chromogenic medium: Fresh Candida isolates were cultured on ChromID Candida Agar [(CAN2), bioMerieux} and incubated for 48 hours at 37ºC. Candida albicans appear as blue colonies, C.tropicalis appear pink while other Candida spp. appear as creamy white colonies (Figure 2).

Fig 1 : Germ tube test of isolated candida species
3. Chlamydospore formation: All Candida isolates were tested to produce chlamydospores on corn meal agar supplemented with 1% (v/v) Tween 80 (CMA-Tween 80) [12]. Positive testing isolates had feathery or spidery outgrowths with spores. Negative culture plates showed no visible sign of morphological change upon examination, but rather maintained their typical blastopore stage (Figure 3).

Sugar assimilation: by Integral System Yeast Plus Commercial Kit (Liofilchem, Ref. 71822 - ITALY) according to the manufacturer’s instructions. The results were comparable with the standard ATCC strains. The specific pattern of sugar assimilation was observed, then a specific code was calculated, and the species name was known from the codebook. All strains of C. albicans and C. tropicalis were identified through numerical code formation and color change in chromogenic well. Some C. glabrata isolates were misidentified by this system as it gives variable color change (Figure 4), and confirmed by Vitek 2 system.

Antifungal Susceptibility Test:
The antifungal susceptibility profiles of all Candida isolates were determined against a panel of antifungicals using the Integral System Yeast Plus (ISYP) (Liofilchem, Ref. 71822 - ITALY) according to the manufacturer’s instructions. Drug sensitivity was evaluated according to growth or inhibition of yeasts in media containing the antifungal and a growth indicator.

Detection of Virulence Factors Candida Species
Extracellular Phospholipase Activity: Candida species were investigated for extracellular phospholipase using the egg yolk agar plate method. Briefly, 10 µL of each standardized cell suspension carefully spotted on two plates of egg yolk agar medium and incubated 37 °C for 48 hours (Figure, 5A). Phospholipase activity (Pz) was measured and interpreted for each isolate as previously described by Price et al. (1982) [13]. Reference strains of C. albicans (ATCC, 10231), C. tropicalis ATCC750 and C. glabrata (ATCC, 2001) were used as controls.

Aspartyl proteinase activity: using Bovine serum albumin (BSA) agar plate's method (Figure, 5B. Briefly by adding 10 µL of each standardized cell suspension to a sterile paper disk, which was applied to the surface of BSA medium that was incubated at 37 °C for five days. Then, the plates were fixed with 20% trichloroacetic acid (TCA) and
stained with 1% Amido Black dye. Enzyme activities were scored according to the criteria by Patil et al (2014). Reference strains of C. albicans (ATCC, 10231), C. tropicalis ATCC750 and C. glabrata (ATCC, 2001) were used as controls.

Fig 5: Phospholipase (A) and Aspartyl proteinase (B) activities by Candida species

Hemolysin activity: Ten microliters of standardized cell suspension of each isolate was spot inoculated on an enriched sheep blood SDA agar medium in triplicate on three separate occasions. Plates were incubated at 37 °C for 48 hours. C. albicans (ATCC 10231) and Streptococcus pyogenes (group A) were used as positive controls for β and α hemolysis, respectively. The presence of a distinctive translucent area around the inoculum indicated positive hemolytic activity. Hemolytic index (Hi) equal the ratio of the diameter of the colony to total diameter of the colony plus the translucent area.

Biofilm formation: Biofilm production was determined by two methods:
- Tube method: A loopful from overnight culture of SDA plate for each isolate was inoculated into a polystyrene Falcon conical tube with a screw cap, containing 10 ml of Sabouraud-dextrose broth with glucose 8% and incubated for 24 h at 37 °C. Then the broth was gently aspirated followed by washing of the tubes with distilled water and staining with 0.1% crystal violet. Excess stain was washed with deionized water. Tubes were dried inverted and then examined for the presence of an adherent layer (Figure, 6). Biofilm production was scored as negative (−), weak (+), moderate (++) or strong (+++)17.
- Congo red Method: Congo red agar medium plates were inoculated with Candida isolates incubated at 37 °C for 24-48 hours aerobically. Positive result was indicated by red colonies, and negative results were indicated by white or light pink-colored colonies (Figure 7).

VITEK 2 System:
The ten isolates suspected to be C. glabrata were tested by VITEK 2 system according to the manufacturer instructions to confirm their identification.

Statistical Analysis: Statistical analysis was conducted, using the mean, standard Deviation, unpaired student t-test, and chi-square tests by SPSS V20.
RESULTS

Two hundred and nineteen patients were investigated for candiduria; 125/219 (57.1%) patients were female and 94/219 (42.9%) were males. The patients’ ages ranged from 50-90 years with mean±SD (62.45±7.93). Table (1) shows the frequency of different predisposing risk factors found among the studied patients.

Table 1: Frequencies of predisposing factors among the studied patients.

<table>
<thead>
<tr>
<th>Risk factors</th>
<th>No.(219)</th>
<th>(%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Urinary catheterization</td>
<td>219</td>
<td>100.0</td>
</tr>
<tr>
<td>Immunosuppressive therapy</td>
<td>22</td>
<td>10.0</td>
</tr>
<tr>
<td>Antibiotic therapy</td>
<td>186</td>
<td>84.9</td>
</tr>
<tr>
<td>Sepsis</td>
<td>12</td>
<td>5.5</td>
</tr>
<tr>
<td>Diabetes mellitus</td>
<td>84</td>
<td>38.3</td>
</tr>
<tr>
<td>Liver disease</td>
<td>73</td>
<td>33.3</td>
</tr>
<tr>
<td>Chronic kidney disease</td>
<td>28</td>
<td>12.8</td>
</tr>
<tr>
<td>Malignancy</td>
<td>22</td>
<td>10.0</td>
</tr>
</tbody>
</table>

Incidence of candida isolation: Fifty out of two hundred and nineteen (22.8%) urine samples were positive for Candida spp., while bacterial growth was detected in 132/219 (60.3%) and 37/219 (16.9%) was negative growth. {Figure 8, (A)}. Candida tropicalis was the most predominant one 27/50 (54.0%) followed by C. albicans 13/50 (26.0%), then C. glabrata 10/50 (20.0%), {Figure 8. (B)}.

Speciation of Candida isolates
(1) Germ tube test: C. albican form germ tube, while C. tropicalis form pseudohyphae, but other isolates didn’t form germ tube.
(2) Chrom ID Agar: C. albican produce blue colonies and C. tropicalis produce pink colonies while all other Candida species produce white colonies, Figure (2).
(3) Chlamydospores formation on CMA-Tween 80 agar: C. albican produce chlamydospore, C. tropicalis produce pseudohyphae, with clusters of blastoconidia at the center, while C. glabrata didn’t produce hyphae or pseudohyphae, Figure (3).
(4) Sugar assimilation: all strains of C. albican and C. tropicalis were identified through numerical code formation and color change in chromogenic well. The specific pattern of sugar assimilation was observed, then a specific code was calculated, and the species name was known from the codebook. The results were comparable with the standard ATCC strains. Some C. glabrata isolates misidentification by this system Figure (4) and confirmed by Vitek 2 system.

Table 2: Phospholipase activity of different Candida species

Phospholipase activity:

<table>
<thead>
<tr>
<th>Candida Species</th>
<th>( \text{C. albican} ) (n=13)</th>
<th>( \text{C. tropicalis} ) (n=27)</th>
<th>( \text{C. glabrata} ) (n=10)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>N %</td>
<td>N %</td>
<td>N %</td>
<td>N %</td>
<td></td>
</tr>
<tr>
<td>Negative</td>
<td>0.0</td>
<td>12.0</td>
<td>44.4</td>
<td>0.001**</td>
</tr>
<tr>
<td>+</td>
<td>0.0</td>
<td>15.6</td>
<td>55.6</td>
<td>0.001**</td>
</tr>
<tr>
<td>++</td>
<td>2.0</td>
<td>15.4</td>
<td>0.0</td>
<td>0.052*</td>
</tr>
<tr>
<td>Total</td>
<td>13.0</td>
<td>100.0</td>
<td>100.0</td>
<td></td>
</tr>
</tbody>
</table>

Table 3: Proteinase activity among the 50 Candida isolates, 34 (68.0%) were positive proteinase. The highest enzymatic activity was reported for C. albicans isolates as 84.6% (11/13) showed strong (+++) proteinase activity, 7.7% (1/13) showed moderate enzyme activity (++). While, 44.4% (12/27) of C. tropicalis had no phospholipase (-) and 55.6% (15/27) had weak enzymatic activity (+).

However, C. glabrata isolates had no enzymatic activity at all. The difference was statistically highly significant (P<0.001). Table (2)
Table 3: Proteinase activity of different Candida species

<table>
<thead>
<tr>
<th>Candida Species</th>
<th>C. albican (n=13)</th>
<th>C. tropicalis (n=27)</th>
<th>C. glabrata (n=10)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>N %</td>
<td>N %</td>
<td>N %</td>
<td></td>
</tr>
<tr>
<td>Interpret.</td>
<td></td>
<td></td>
<td></td>
<td>&lt;0.007**</td>
</tr>
<tr>
<td>Negative</td>
<td>0 0.0</td>
<td>10 37.0</td>
<td>6 60</td>
<td></td>
</tr>
<tr>
<td>+</td>
<td>1 7.7</td>
<td>17 63.0</td>
<td>4 40</td>
<td>&lt;0.004**</td>
</tr>
<tr>
<td>++</td>
<td>1 7.7</td>
<td>0 0.0</td>
<td>0 0</td>
<td>0.234</td>
</tr>
<tr>
<td>+++</td>
<td>11 84.6</td>
<td>0 0.0</td>
<td>0 0</td>
<td>&lt;0.001**</td>
</tr>
<tr>
<td>Total</td>
<td>13 100</td>
<td>27 100</td>
<td>10 100</td>
<td></td>
</tr>
</tbody>
</table>

*Non-significant **highly significant

Table 4: Haemolytic activity of different Candida species

<table>
<thead>
<tr>
<th>Candida Species</th>
<th>Number</th>
<th>Beta</th>
<th>Alpha</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>N</td>
<td>%</td>
<td>N %</td>
</tr>
<tr>
<td>C. albican</td>
<td>13</td>
<td>13</td>
<td>100</td>
<td>0 0</td>
</tr>
<tr>
<td>C. tropicalis</td>
<td>27</td>
<td>6</td>
<td>22.2</td>
<td>21 77.8</td>
</tr>
<tr>
<td>C. glabrata</td>
<td>10</td>
<td>0</td>
<td>0</td>
<td>10 100</td>
</tr>
</tbody>
</table>

** P value highly significant

Table 5: Biofilm formation by isolated Candida species

<table>
<thead>
<tr>
<th>Candida Species</th>
<th>C. albican (N=13)</th>
<th>C. tropicalis (N=27)</th>
<th>C. glabrata (N=10)</th>
<th>Total (N=50)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>N %</td>
<td>N %</td>
<td>N %</td>
<td>N %</td>
<td></td>
</tr>
<tr>
<td>Score of Biofilm</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>&lt;0.001**</td>
</tr>
<tr>
<td>Negative</td>
<td>2 15.4</td>
<td>0 0</td>
<td>1 10</td>
<td>11 22</td>
<td>0.133</td>
</tr>
<tr>
<td>+</td>
<td>9 69.2</td>
<td>0 0</td>
<td>9 90</td>
<td>18 36</td>
<td>&lt;0.001**</td>
</tr>
<tr>
<td>++</td>
<td>1 7.7</td>
<td>6 22.2</td>
<td>0 0</td>
<td>7 14</td>
<td>0.157</td>
</tr>
<tr>
<td>+++</td>
<td>1 7.69</td>
<td>21 77.8</td>
<td>0 0</td>
<td>22 44</td>
<td>&lt;0.001**</td>
</tr>
</tbody>
</table>

** P value highly significant

Antifungal susceptibility: Sensitivity of C. albicans was 100% to Amphotericin and Flucytosine, 76.9% to Nistatin and Voriconazole, 69.2% to Ketoconazole and fluconazole, 61.5% to Miconazole and itraconazole, 53.8% to Econazole, and 46.2% to Clotrimazole. Sensitivity of NAC was 27.0% to Amphotericin, 94.5% to Flucytosine, 9.2% to Nistatin, 10.8% to Voriconazole and Itraconazole, 75.6% to Ketoconazole, 13.5% to fluconazole, 8.1% to Miconazole and 2.7% to Econazole, and 5.4% to Clotrimazole, Figure (9).

Fig 9: Antifungal susceptibility pattern of the isolated C. albicans versus NAC.

While, C. albicans didn’t show any antifungal resistance, most of NAC spp. showed higher rates of resistance to several azole drugs (Clotrimazole 77.6%, Miconazole 70.3%, Econazole 63.2%, Nystatine 44.3%), Figure (10).
findings were reported by Melges et al., 2019 [18]. The average age of patients with candiduria was 64 years. Similar to other reports, most of patients in our study who confirmed with positive candiduria were between ages 50-70 years with predominance of female gender 64.0% (32/50). Jiménez-Guerra et al., 2018 [23] found that the average age of patients with candiduria was 64 years. Similar findings were reported by Melges et al., 2019 [18] who reported that the average age of patients with candiduria was 63 years and 57.0% were females.

Regarding the associated predisposing risk factors, antibiotics therapy is one of the most important risk factors exposing the patients to candiduria. This correlates with our results as all candiduria positive patients with Urinary catheterization 100% and 78.0% (39/50) were on antibiotic treatment. Our results are supported by Melges et al., 2019 [18] who reported that antibiotic therapy 89.2% and the use of an indwelling urinary catheter were 100% the most frequent predisposing factors. And Chandak et al., 2018 [21] who noticed that antibiotics were the most prominent associated risk factor 100%. Diabetes mellitus is a common disease also predisposes to Candida colonization. Our findings showed that 50.0% (25/50) of candiduria positive patients were diabetics in accordance with the results of Melges et al., 2019 [18] who reported that 34.4% of UTI patient with candiduria. Also, Chandak et al., 2018 [21] who found that 28.0% of candiduria positive ICU patients were diabetics.

Our findings showed that 22.8% (50/219) of catheterized ICU patients had candiduria and this was supported by the results of Montes et al., 2019 [22] who reported that 37.7% of Candida spp. isolated from UTI Sahai & Kumar., 2018 [23] who found that out of 100 patients, 26 patients were diagnosed to develop symptomatic CAUTI during their course of hospitalization.

Worldwide, NAC spp. which appear to be better adapted to the urinary tract environment have now replaced C. albicans as the predominant pathogen in nosocomial urinary tract infections Biagi et al., 2019 [24]. The current study revealed that 74.0% (37/50) of Candida isolates were NAC species, while 26.0% (13/50) confirmed as C. albicans. Chandak et al., 2018 [21] in their study found that NAC spp were predominantly high 74.0%, while C. albicans was 26.0%. And Sahai & Kumar., 2018 [23] who reported that NAC spp. contributed to 83.3% of the isolates and only 16.7% of isolates were C. albicans. However, in contrary, Garcia-Agudo et al., 2018 [19] reported that C. albicans was the predominant Candida species 61.9%.

Moreover, our results showed that the NAC species were predominantly isolated as C. tropicalis represented 54.0% (27/50) followed by C. glabrata 20.0% (10/50) and this was in agreement with Melges et al., 2019 [18] who reported that C. tropicalis was found most commonly 37.6%, followed by C. albicans 36.6% and C. glabrata 19.3%. And Chandak et al., 2018 [21] who showed that C. tropicalis was the most common amongst NAC representing 72.9% followed by C. glabrata 16.3%. However, in contrary, Montes et al., 2019 [22] reported that the most frequent species were C. albicans 42.9%. C. tropicalis 20.9% and C. glabrata 16.9%.

In the present work, all recovered Candida spp. were assessed for the production of extracellular enzymes as virulence factors including phospholipase, proteinase and haemolysin.

According to our results, C. albicans had the highest phospholipase activity as 84.6% (11/13) of the C. albicans isolates had strong phospholipase activity and 15.4% (2/13) had moderate enzymatic activity. While 44.4% (12/27) of C. tropicalis had no phospholipase (-). 55.6% (15/27) had weak enzymatic activity and all isolates of C. glabrata had no enzymatic activity at all. Similar results were reported by Dabiri et al., 2018 [26] who founded that the highest positivity of phospholipase production was related to C. albicans 56.8% followed by C. tropicalis 50.0% and C. glabrata isolates 20.0%.

However, in case of proteinase enzyme activity it was shown that C. glabrata was the lowest proteinase producer as 40.0% (4/10) showed only weak enzymatic activity and 60.0% (6/10) were negative followed by C. tropicalis as 63.0% (17/27) showed weak enzymatic activity and 37.0% (10/27) were negative. However the highest enzymatic activity were reported for C. albicans isolates as 84.6% (11/13) showed strong enzymatic activity, 7.7% (1/13) showed moderate activity and 7.7% (1/13) showed weak activity with highly statistical significant differences (P<0.001), in agreement with the results of Dabiri et al., 2018 [26] who reported that C. albicans showed the highest proteinase and phospholipase activity and biofilm formation ability. And Alenzi, 2016 [27] founded that Secretory aspartic proteinase activity was demonstrated in 48.0% of Candida isolates.

Meanwhile, haemolysin secretion followed by iron acquisition facilitates deeper tissue invasion by Candida, Tsang et al., 2007 [28]. In the present study C.
albicans produce beta haemolytic activity, this in agreement with Zarrin et al., 2015 and Sachin et al., 2012 who showed that all C. albicans isolates exhibited haemolytic activity. All C. tropicalis were haemolysin producer alpha and beta. This in agreement with Udayalax & Shenoy., 2016 who reported that all isolates of Candida spp. both C. albicans and NAC species, produced haemolysin (100%). While Sachin et al., 2012 found only 30.4% of C. tropicalis showed haemolytic activity.

One of the main virulence factors are surface molecules that permit adherence of the organism to the structures (e.g., human cells, extra cellular matrix, and prosthetic devices). A biofilm is a community of microorganisms and their extra cellular polymers that are attached to a surface.

Out of the 50 Candida isolates 22 (44.0%) isolates showed strong biofilm formation, 7/50 (14.0%) strains showed moderate biofilm formation and 18/50 (36.0%) strains showed weak biofilm formation. The results were comparable with the standard ATCC strains.

Biofilm production was reported in our study mostly to NAC species 97.3% (36/37) than C. albicans 84.6% (11/13) and this is also in accordance with Yarlagadda et al., 2017 who showed that biofilm production was detected most frequently among NAC species 48.0% than C. albicans 25.0%. As well as Sida et al., 2016 reported that biofilm production was found to occur by 65.2% of NAC species compared to 34.7% of C. albicans. Also, Shekar & Kabbin., 2015 reported that out of the 26 Candida isolates 14 (53.8%) were biofilm producer, 10/14 (71.5%) were NAC while 4/14 (28.5%) only were related to C. albicans. However, Al kilani et al., 2012 found that biofilm formation is more in C. albicans versus C. tropicalis 83.3% and 57.9%, respectively.

The major strong biofilm producer was C. tropicalis 77.7% (21/27) followed by C. albicans 7.7% (1/13) with highly significant difference (P<0.001). Biofilm formation also was reported in 63.6% of C. tropicalis versus (21.2%) for C. albicans in a study done by Chandak et al., 2018.

Several studies have reported that amphotericin B was the most effective drug against Candida isolates. Fluconazole is considered as a potentially useful drug in Candida UTI due to its high concentration in urine and better tolerance; however, increasing resistance to azoles is being reported.

In our study, the results showed that C. albicans was sensitive to all tested antymycotic drugs with maximum sensitivity 100% to Flucytosine and Amphotericin and the other night antymycotic showed either sensitivity or intermediate sensitivity without absolute resistance. In concordance with this finding, Sahai & Kumar., 2018 reported that all Candida isolates were sensitive to fluconazole, voriconazole, amphotericin B and itraconazole.

While NAC spp. were sensitive only to Fluconosine 94.59% and Ketonazole 75.67%, and it was highly resistant to other tested Azole including Clotrimazole 77.6%, Miconazole 70.3%, Econazole 63.2% and Nystatine 44.3%. Similar rates of fluconazole resistance were reported by Yarlagadda et al., 2017.

CONCLUSION

This study documents the prevalence of candiduria in catheterized patients and the change in trend with shift toward non albicans Candida species as the predominant pathogen causing nosocomial UTI. A wide range of intra- and inter- species enzymatic activities and biofilm forming ability have been accounted for testing the pathogenicity of Candida species in susceptible individuals, while correlation with clinical outcomes is still need to be validated. Biofilm formation is seen more frequently with non albicans candida species than with Candida albicans and as biofilm production may help maintain the role of fungi as commensals and pathogen, by evading host defence mechanisms, resisting fungal treatment and withstanding the competitive pressure from other organisms, these are difficult to treat. Studies of urine cultures in hospitalized patients should include the possible presence of candiduria. Species identification and biofilm detection must be performed for early and effective treatment of the patient.

REFERENCES


