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Species Identification and Antifungal Susceptibility Profile of Candida Isolates from Various Clinical Specimen: A Three Years Study at a Tertiary Care Teaching Hospital in Coimbatore, TN

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ABSTRACT

In recent years, the incidence of fungal infections in general and candidiasis in particular has increased. Although traditionally, C. albicans is considered as the most pathogenic species, recent studies have highlighted emergence of non albicans Candida (NAC) spp. These 'cryptic' NAC spp. often develop resistance during course of therapy or are innately less susceptible to commonly prescribed antifungal drugs. The present study was conducted with an aim to identify Candida isolates up to species and to study their antifungal susceptibility profile. Candida isolates were identified up to species level by standard mycological protocol. Antifungal susceptibility testing of isolates was performed by disc diffusion method as outlined in Clinical Laboratory Standards Institute's (CLSI) M44-A guidelines. A total of 207 isolates were Candida spp. Predominance of NAC spp. was noted. C. tropical is was the predominant Candida spp. Fluconazole resistance was observed in 25.6% of Candida isolates. All isolates of C. krusei were resistant to fluconazole. Amphotericin B resistance was noted 4 (1.9%) isolates which included 2 isolates of C. krusei and single isolate each of C. Tropical is and C. glabrata. Non albicans Candida species have emerged as an important cause of infections. C. Tropical, C. krusei and C. glabrataare important NAC spp. As different Candida spp. varies widely in their susceptibility to commonly prescribed antifungal drugs, antifungal susceptibility testing often guides clinicians in selection of most appropriate antifungal therapeutic agent.

INTRODUCTION

Health and diseases have always been the most important matter of concern to human beings. The ever escalating want of better health resulted in discovery of various novel techniques for diagnosing and treating diseases. However, even in the golden era of medicine, where diagnostic and therapeutic modalities have progressed by leaps and bounds, infectious diseases still continue to be one of the major causes of morbidity and mortality in clinical setups worldwide. Infectious diseases are attributed to bacteria, fungi, parasites and viruses [1]. As compared to other counterparts, bacteria are commonly implicated in human infections. However in recent years, the incidence of fungal infections in general and candidiasis in particular has increased^[2]. Various factors like use of broad spectrum antibiotics and immunosuppressive drugs along with increase in immunocompromised individuals are considered important for increased incidence of fungal infections^[3]. Candida spp. is probably the only fungal pathogens that causes broad spectrum of clinical manifestation ranging from mucosal overgrowth to life threatening disseminated infections^[4]. Although traditionally, C. albicans is considered as the most pathogenic species, recent studies have highlighted emergence of non albicans Candida (NAC) spp^[5]. These 'cryptic' NAC spp. often develop resistance during course of therapy or are innately less susceptible to commonly prescribed antifungal drugs [6]. Therefore the present study was conducted at a tertiary care teaching hospital with an aim to identify Candida isolates up to species and to study their antifungal susceptibility profile.

MATERIALS AND METHODS

The present descriptive cross-sectional study was conducted in the Department of Microbiology in a private medical college in Tamil Nadu for a period of three years (January 2018 to December 2020). Candida spp. isolated from various clinical specimens were included in the study. Candida isolates were identified upto species level by standard mycological protocol for identification of yeast and yeast like fungi^[7]. Germ tube technique, carbohydrate assimilation test and colony color on Hichrom Candida agar (Himedia Laboratories Pvt Ltd Mumbai) were used for identification of Candida spp. Antifungal susceptibility testing of isolates was performed by disc diffusion method as outlined Clinical Laboratory Standards Institute's (CLSI) M44-A guidelines. Disc diffusion test was performed by Kirby-Bauer disc diffusion method on Mueller Hinton agar supplemented with 0.2% glucose and 0.5µg/mL methylene blue agar (MH-GMB). Isolates were tested for susceptibility against antifungal drugs like amphotericin B (20µg), fluconazole (25 µg), itraconazole (10 μg) and ketoconazole (10 μg). All antifungal discs were procured from Himedia Laboratories Pvt Ltd. Mumbai. Statistical analysis was done using statistical packages SPSS 19.0 (SPSS Inc. Chicago, IL). A P value <0.05 was considered to be significant. Pictorial presentation of the key findings was done by using appropriate statistical graph.

RESULTS AND DISCUSSIONS

During the study period, a total of 4853 clinical specimens were received in the Department of Microbiology. Out of these, 1432 (29.5%) were culture positive. The rates of isolation of bacteria and fungi were 79.2% (n=1134) and 20.8% (n=298) respectively. Among fungal pathogens, a total of 207 isolates were Candida spp. (Fig. 1). Isolation of Candida spp. was significantly high among various fungal pathogens (Fisher Exact test P value <0.05).

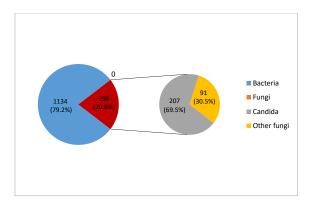


Fig. 1: Bacterial and Fungal Isolates from Various Clinical Specimens

The year distribution of Candida spp. is shown in (fig. 2). There increasing isolation trend observed for Candida spp.

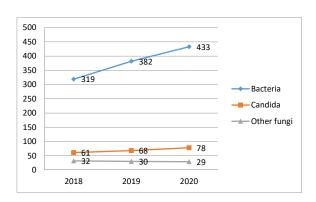


Fig. 2: Year Wise Distribution of Candida and Other Isolates from Clinical Specimens

The clinical specimen wise distribution of Candida spp. is shown in (fig. 3). The maximum number of Candida isolates were isolated from urine samples (37.7%) followed by vaginal swabs (25.6%). A total 25 (12.1%) Candida spp. were isolated from blood cultures.

Female gender, advanced age group, diabetes, use of broad spectrum antibiotics and presence of urinary catheter were important risk factors observed for isolation of Candida spp. from urine samples. Pregnancy, HIV infection and antibiotic therapy were major risk factors for vulvovaginal candidiasis whereas HIV infection and diabetes were the only risk factors noted in patients with oropharyngeal Candida infection. ICU stay, mechanical ventilator support, indwelling urinary catheter and central venous catheterisation were risk factors for Candida blood stream infection (BSI).

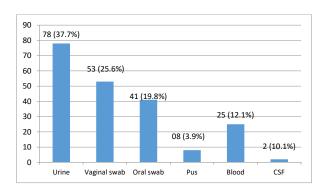


Fig. 3:The Clinical Specimen Wise Distribution of Candida Spp

In the present study, a total of 62 (29.9%) isolates were identified as C. albicans whereas 145 (70.1%) isolates belonged to NAC spp. Isolation of NAC spp. was significantly high (Fisher Exact test P value <0.05). The species wise distribution of Candida isolates is shown (table 1). C. tropical is (29.9%) was the predominant isolate in the present study. C. krusei was isolated from 25 (12.1%) clinical specimens.

Table 1: The Species Wise Distribution of Candida Isolates

Candida spp.	Number (%)
C. albicans	62 (29.9)
C. tropicalis	75 (36.2)
C. krusei	25 (12.1)
C. glabrata	21 (10.1)
C. kefyr	15 (7.2)
C. guilliermondii	09 (4.3)
Total	207

Antifungal susceptibility profile of Candida spp. by disc diffusion method is shown in (table 2). Fluconazole resistance was observed in 25.6% of Candida isolates. All isolates of C. krusei were resistant to fluconazole. Itraconazole resistance was observed in 10 (4.8%) isolates. A total of 7 (28%) isolates of C. krusei were resistant to itraconazole. Amphotericin B resistance was noted 4 (1.9%) isolates which included 2 isolates of C. krusei and single isolate each of C. tropicalis and C. glabrata.

In recent years, fungi, once studied in microbiology only as 'microbiological curiosities' without any clinical significance have emerged as an important cause of infections^[7,8]. Researchers from various parts of the world have reported increase in the incidence of fungal infections. Similarly in the present study, the rate of isolation of fungi from various clinical specimens was 20.8% and there was increasing trend observed for isolation of Candida spp. Candida spp. was the predominant fungal pathogen. Risk factors for Candida infections are similar to those for other fungal infections but may vary in cause and are mostly related to medical intervention and immune status of the patient^[8,9]. In the present study, 37.7% of Candida isolates were isolated from urine samples. As Candida is often found as commensal of genitourinary tract, its isolation from urine sample usually presents as a challenge to both clinicians and microbiologists as to whether ignore it as a mere colonization or, consider it as an indicator of lower or upper urinary tract infection including ascending pyelonephritis and renal candidiasis with sepsis^[9]. However, isolation of urine should never be overlooked as this condition may be the only and often the first indication of systemic Candida infection^[10]. In clinical practice, candiduria is a relatively rare finding in a structurally normal urinary tract and in healthy individuals[11]. Female gender, advanced age group, diabetes, use of broad spectrum antibiotics and presence of urinary catheter were important risk factors of candiduria in the present study. Similar findings were reported by other researchers like Kauffmann^[13], Bukhary^[11] and Achkeret^[12]. The waning of host immune defenses in geriatric age group may explain the high incidence of candiduria. As in females the Candida colonization of vulvo vestibular area with Candida spp. is frequent, they are more at risk of developing candiduria due to ascending infection^[14]. In diabetes glycosuria, impaired host defense mechanism, stasis of urine in neurogenic bladder enhance candiduria. As antibiotics suppress and change endogenous bacterial flora of the perineum, it favors both colonization and infection of urinary tract by Candida spp^[15]. In the present study, a total of 25 Candida spp. were isolated from blood cultures. In the United States, Candida spp. is the third most common cause of hospital associated blood stream infections. Candidemia is associated with high morbidity and mortality in both immunosuppressed and critically ill immuno competent patients [16,17]. It also significantly increases the duration of hospital stay and requirement of mechanical ventilation. In the present study, predominance of NAC spp. over C. albicans was noted. Several research studies have documented emergence of NAC spp. Although various factors like increased empirical use of azoles are cited

Table 2: Antifungal Susceptibility Profile of Candida Spp. by Disc Diffusion Method

Candida spp.	Fluconazole		Itraconazole	Itraconazole		Ketoconazole		Amphotericin B	
	S (%)	R(%)	S (%)	R (%)	S (%)	R (%)	S (%)	R (%)	
C. albicans (62)	51 (82.2)	11 (17.8)	61 (98.4)	01 (1.6)	50 (80.6)	12 (19.4)	62 (100)	-	
C. tropical is (75)	63 (84)	12 (16)	73 (97.3)	02 (2.7)	68 (90.7)	07 (9.3)	74 (98.7)	01 (1.3)	
C. krusei(25)	-	25 (100)	18 (72)	07 (28)	12 (48)	13 (52)	23 (92)	02 (8)	
C. glabrata(21)	16 (76.2)	05 (23.8)	21 (100)	-	18 (85.7)	03 (14.3)	20(95.2)	01 (4.8)	
C. kefyr(15)	15 (100)	-	15 (100)	-	14 (93.3)	01 (6.7)	15 (100)	-	
C. guilliermondii(09)	09 (100)	-	09 (100)	-	09 (100)	-	09 (100)	-	
Total (207)	154 (74.4)	53 (25.6)	197 (95.2)	10 (4.8)	171 (82.6)	36 (17.4)	203 (98.1)	04(1.9)	

for emergence of NAC spp., improvement in diagnostic mycological methods like use of chromogenic media with an ability to differentiate Candida spp. as well as introduction of molecular diagnostic techniques appears to be more important^[18,19]. C. tropical is was the predominant species. Similar to our observation, the predominance of C. tropical is among various Candida isolates was also reported by Pahwa^[20] and Deorukhkar^[21] C. tropical is alone, or in association with other species, is more frequently implicated in human infections^[22]. It is more frequently isolated from patients admitted to ICU, especially those requiring prolonged catheterisation, treated with broadspectrum antibiotics or with malignancies^[20]. C. krusei was isolated 25 (12.1%) clinical specimens. It is considered as an emerging NAC spp. C. krusei infections are highly uncommon outside clinical settings involving exposure to azoles and immuno compromised status^[23]. In the present study, antifungal susceptibility profile of Candida spp. was studied by disc diffusion method^[24]. Compared to the Clinical and Laboratory Standard Institute (CLSI) standardized micro broth dilution method, disc diffusion susceptibility testing is convenient and economical^[25]. In this study, fluconazole resistance was observed in 25.6% of Candida isolates. Of various antifungal drugs, resistance to fluconazole is of concern because it is one of the most widely used first line antifungal agents for treatment and prophylaxis of all forms of candidiasis^[26]. All isolates of C. krusei were resistant to fluconazole. Various national and international studies have reported total fluconazole resistance in C. krusei isolates. In the present study, Candida spp. were more susceptible to amphotericin B. Amphotericin B is generally regarded to have the broadest spectrum of antifungal activity and used in life-threatening disseminated infections^[27]. Although, amphotericin B resistance was low in this study, high cost and associated toxicity restricts its use^[27].

CONCLUSION

Non albicans Candida species have emerged as an important cause of infections. C. tropicalis, C. krusei and C. glabrataare important NAC spp. As different Candida spp. varies widely in their susceptibility to commonly prescribed antifungal drugs, antifungal

susceptibility testing often guides clinicians in selection of most appropriate antifungal therapeutic agent.

Conflict of Interest: Nil.

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