

Original Research Article

<http://dx.doi.org/10.20546/ijcmas.2016.510.032>

Speciation and Antifungal Susceptibility of *Candida* Species Isolated from Clinical Samples – A Pilot Study

M. Saleem^{1*}, R. Gopal¹, T. Mangaiyarkarsi¹, S. Sunil¹, J. Krishnapriya¹ and R. Nagma²

¹Department of Microbiology, Sri Manakula Vinayagar Medical College and Hospital, Puducherry, India

²Department of Microbiology, SMVMCH, Puducherry, India

*Corresponding author

ABSTRACT

Infections due to *Candida* species are the most common fungal infections. The increase in the population at risk of developing invasive candidiasis viz. immunocompromised patients, patients with diabetes mellitus, prolonged use of antibiotics, immunosuppressive therapy has resulted in a rise in infections due to non-albicans *Candida* species which are resistant to fluconazole. All *Candida* species isolated from various clinical specimens during a period of 6 months (Jan 2016 to June 2016) were speciated by conventional tests and grown on SDA, chlamyospore agar and chrome agar. All *Candida albicans* isolates were tested for susceptibility to 'Fluconazole' by disc diffusion. A total of 54 specimens yielded *Candida* species. Non-albicans *Candida* species constituted the predominant isolates and the highest yield was from urine samples. *Candida albicans* was isolated from 20 of the 54 specimens. Only 65% of the *Candida albicans* were susceptible to fluconazole. Chrome agar was found to be very useful in isolation and speciation of the *Candida* species. Non-albicans *Candida* is the most common cause of candidiasis in our region. Resistance to fluconazole among *Candida albicans* is high (35%).

Keywords

Candida,
speciation,
fluconazole,
resistance,
non-albicans
Candida species

Article Info

Accepted:
14 September 2016
Available Online:
10 October 2016

Introduction

Candida is yeast-like fungi which are ubiquitous in nature (Edwards, 2012). The genus comprises of several hundred species, of which only a few cause disease in humans (Chander, 2012). The *Candida* species are commensals of the oropharynx, intestine, vagina and skin (Edwards, 2010). The incidence of *Candida* infections has increased at an alarming rate over the past

few decades (Raut *et al.*, 2009). *Candida albicans* the most common species isolated from clinical specimens earlier is replaced by non-albicans species in most of the hospital based studies. The increase in the population of immunocompromised individuals due to various causes, patients with diabetes mellitus, prolonged hospital stay and antibiotic therapy, invasive

procedures etc. have all led to the increase in the prevalence of candidiasis particularly with non-albicans species (Oliviera *et al.*, 2006). Although all species of candida can produce invasive infections, they show considerable difference in disease severity and susceptibility to antifungal agents (Shaheen *et al.*, 2006), hence it is necessary to identify the species to initiate appropriate antifungal agent for a better outcome.

Materials and Methods

All the candida species isolated from various clinical specimens received in the Microbiology lab during the study period of six months were speciated and susceptibility of *Candida albicans* to fluconazole determined. All the significant growth of candida species identified by Gram's stain from the colonies on blood agar (BA) or other routine media were subcultured on SDA with gentamycin and Hichrom candida agar (Himedia labs pvt. Ltd.) and incubated at 35°C for 24-48 hours. A germ tube test was performed from the growth on SDA after reconfirming the isolate as yeast by Gram's stain and also inoculated on chlamyospore agar, urease agar and sugar fermentation and assimilation test performed as per the standard procedures.

A disk diffusion test on Mueller-Hinton agar with 2% glucose and methylene blue (Himedia labs) was performed to determine the susceptibility of the *Candida albicans* species isolated to fluconazole as per the criteria given in the manufacturer's product insert. The species of candida was identified based on the macro and microscopic

morphology of the growth on the SDA, chlamyospore agar and Hichrom candida agar in addition to the conventional tests.

Results and Discussion

A total of 54 isolates comprising of five species of candida were isolated from the clinical specimens during the study period. Of the 54 samples maximum number of Candida species were isolated from the urine samples (81.48%). Candida species were isolated more from the females (61.11%) as compared to males (20.37%). Among the 54 samples Candida non-albicans was the most common causative agent comprising of *Candida glabrata* (25.92%), *Candida tropicalis* (16.6%), *Candida krusei* (11.11%) and *Candida parapsilosis* (9.25%) whereas *Candida albicans* showed a distribution of 37.03%. 35% of the *Candida albicans* species were found to be resistant to Fluconazole.

Candida species though part of the commensal flora can nevertheless produce invasive mycoses of endogenous or exogenous origin. The severity of the disease can vary widely from a localized micro-cutaneous lesion to a disseminated type of disease with varying morbidity and mortality. A host of factors including prolonged treatment with antibiotics, immunosuppressive therapy, immunosuppressive diseases like HIV, malignancy, diabetes mellitus, organ transplant recipients, invasive procedures predispose to the development of opportunistic invasive candidiasis (Tankhiwale *et al.*, 2012).

Table.1 Sample wise distribution of *Candida albicans* and non albicans species

S. No.	Samples	No. of samples	<i>Candida albicans</i>	Candida non-albicans
1	Urine	44	18	26
2	Sputum	1	1	-
3	Hvs	6	1	5
4	Blood	1	-	1
5	Pus	1	-	1
6	Wound	1	-	1

Table.2 Sex wise distribution of *Candida* species isolated.

Sample	Male	Female
Urine	11	33
Sputum	-	1
HVS	-	6
Blood	-	1
Pus	-	1
Wound	-	1
Total	11	43

Table.3 Distribution of *Candida* species

Species Isolated	No. of samples n=54
<i>Candida albicans</i>	20
<i>Candida glabrata</i>	14
<i>Candida tropicalis</i>	9
<i>Candida krusei</i>	6
<i>Candida parapsilosis</i>	5

Table.4 susceptibility of *Candida albicans* to Fluconazole

	Sensitive	Resistant
<i>Candida albicans</i> n=20	13	7

Fig.1 Candida species differentiation in CHROM agar.



In the present study out of 54 isolates 43 (79.6%) were from females i.e. a female preponderance which is similar to the study by Madhumati and Rajendran (Madhumati *et al.*, 2015). The highest number of isolates (44 out of 54) were from urine. Among the urinary isolates non-albicans candida constituted 26 out of 54 while *Candida albicans* to 18 which is similar to the study by Madhumati (Madhumati *et al.*, 2015), Bhaskar (Bhaskar *et al.*, 2015) and Ravinder Sandhu (Sandhu *et al.*, 2015). Among the candida species isolated from various specimens non-albicans candida accounted for 34 out of 54 of the total while *Candida albicans* to 20 out of 54 thus indicating that non-albicans candida are the predominant pathogens in our area as was the case in

studies by authors. *Candida glabrata* constituted the predominant species among the non-albicans (14 out of 34) species. Among the *Candida albicans* isolates 13 out of 20 were susceptible to fluconazole (65%) which is similar to a study by Ravinder Sandhu *et al.*, (2015) and Bhaskar *et al.*, (2015). There was a good correlation between conventional tests and CHROM agar for speciation.

In conclusion, the present study shows that non-albicans candida is the predominant pathogen in this region and that only 65% of *Candida albicans* are susceptible to fluconazole. The study also shows that the chromogenic agar is a very useful medium for isolation and species identification being

a simple and time saving method which can be used as a routine substituting for the laborious conventional fermentation and assimilation tests. The study also shows a decrease in the susceptibility of *Candida albicans* to fluconazole.

Acknowledgement

We thank the institutional ethics committee for permitting us to conduct this pilot study.

Conflict of interest: nil.

References

- Bhaskar, U.A., Yashavanth, R., Ronald, R. 2015. Identification of *Candida* species from clinical samples and their antifungal susceptibility patterns. *J. Evol. Med. Dent. Sci.*, 4(75): 12998-13004.
- Chander, J. 2012. Editor. Text book of Medical Mycology. 3rd Ed. New Delhi: Mehta Publishers, P 266-2.
- Edwards, J.E. 2010. *Candida* SPECIES. In: Mandell GL, Bennett JE, Dolin R, Editors. Mandell, Douglas & Bennett's Principles and Practice of Infectious diseases. 7th Ed. Philadelphia: Churchill Livingstone, P3225-38.
- Edwards, J.E. 2012. Candidiasis. In: Longo, Fauci, Kasper, Hauser, Jameson, Loscalzo, Editors. Harrison's Principles of Internal Medicine. 18th Ed. New York: The McGraw-Hill companies, 1651-2.
- Madhumati, B., Rajendran, R. 2015. "Evaluation of chrom agar in speciation of *Candida* species from various clinical samples in a tertiary care hospital". *Int. J. Curr. Microbiol. App. Sci.*, 4(9): 463-72.
- Oliviera, G.S., Ribiero, E.T., Baroni, F.A. 2006. An evaluation of manual and mechanical methods to identify *Candida* spp. from human and animal sources. *Rev. Inst. Med. Trop. S Paulo*, 48(6): 311-15.
- Raut, S.H., Varaiya, A. 2009. Differentiation of *Candida dublinensis* on Chrom Agar & Pal's Agar. *IJMM*, 27(1): 55-58.
- Sandhu, R., Dahiya, S., Sharma, R.K. 2015. "Isolation and identification of *Candida* and Non *albicans* *Candida* species using chromogenic medium". *Int. J. Biomed. Res.*, 6(12): 958-62.
- Shaheen, M.A., Taha, M. 2006. Species identification of *Candida* isolates obtained from oral lesions of hospitalized and non-hospitalized patients with oral candidiasis. *Egyptian Dermatol. Online J.*, 2(1).
- Tankhiwale, S., Gajbhiye, S., Powar, R. 2012. Fluconazole susceptibility testing of *Candida* species by disk diffusion and agar dilution method. *JMEDS*, 1(4): 527-32.

How to cite this article:

Saleem, M., R. Gopal, T. Mangaiyarkarsi, S. Sunil, J. Krishnapriya and Nagma, R. 2016. Speciation and Antifungal Susceptibility of *Candida* Species Isolated from Clinical Samples – A Pilot Study. *Int.J.Curr.Microbiol.App.Sci.* 5(10): 289-293.
doi: <http://dx.doi.org/10.20546/ijcmas.2016.510.032>