



A STUDY ON IDENTIFICATION AND SPECIATION OF MEDICALLY IMPORTANT CANDIDA SPECIES ISOLATED FROM VARIOUS CLINICAL SAMPLES BY USING HICROME CANDIDA.

Microbiology

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ABSTRACT

Candida albicans remains the most frequently isolated species, but an increase in the prevalence of Non-*albicans* candida is a matter of concern for various laboratories, as NAC show less susceptibility to antifungal agents particularly azoles^{1,3}. Routinely used conventional methods are cumbersome and time consuming. Hence the present study aimed at species identification of candida isolated from various clinical specimens and also to evaluate the usefulness of HiCrome Candida differential agar as compared to routine conventional method for speciation of candida. A total of 51 non repetitive clinical isolates of Candida were obtained from various clinical samples. *Candida albicans* was the major species accounting for 23 (45.10%) of the total isolates. *Non albicans Candida* constituted 17 (33.33%) of *C. tropicalis*, 9 (17.64%) of *C. krusei* and 2 (3.92%) of *C. parasilosis*. 49 Candida species which were correctly identified by HiCrome Candida agar except 2 species of *C. parasilosis* (identified by conventional method) which were identified as *C. glabrata* by HiCrome Candida agar.

KEYWORDS

Non-*albicans* Candida, HiCrome Candida differential agar, conventional method, candida speciation.

INTRODUCTION

Candida species particularly *Candida albicans* remains the most common cause of superficial and deep fungal infections worldwide^{1,2}. But an increase in the prevalence of Non *albicans* Candida (NAC) during the last few decades is a matter of concern for various laboratories, as NAC show less susceptibility to commonly used antifungal agents particularly azoles ie fluconazole^{3,5}. Thus species identification of *Candida* isolates have become very necessary for providing effective antifungal therapy, which will in turn prevent the emergence of drug resistance and also nosocomial infection⁶.

A large variety of techniques from conventional to molecular methods are available for yeast identification^{4,5}. Routinely used conventional methods like gram staining of smear, germ tube test, colony morphology on Sabouraud's dextrose agar, urease test, growth on corn meal agar, sugar fermentation and assimilation test, growth at 45°C are labour intensive, cumbersome and time consuming ie may take 3-4 days⁷.

Now a days, newer methods like several chromogenic media, BiGGYagar system, API system, Vitek 2ID system and molecular methods, have been developed for rapid identification of *Candida* species⁸⁻¹⁰. HiCrome Candida agar is one of the selective and differential chromogenic media manufactured by Himedia, Mumbai, India. This media contain chromogenic substrates that react with enzymes secreted by microorganisms producing colonies with various pigmentation. These enzymes are species specific, allowing organisms to be identified to the species level by their colour and colony characteristics¹¹. They are used for rapid isolation and identification of candida species directly from clinical samples and also where multiple yeast species are present in the sample. On these media results are obtained in less time ie within 48hrs, can be easily interpreted and is also cost effective as compared to other newer methods like API system, Vitek 2ID system and molecular methods which are expensive¹²⁻¹⁵.

As many tests from conventional to molecular methods are available, clinical microbiological laboratories are facing an important challenge regarding selection of method that is cost effective, accurate, easily interpreted and also rapid for identification of *Candida*, which in turn is of great importance for clinicians of our hospital to select appropriate prophylactic and therapeutic antifungal drug. Keeping the above point in view, the present study aimed at species identification of *Candida* isolated from various clinical specimens and also to evaluate the usefulness of HiCrome Candida differential agar as compared to routine conventional method for speciation of *Candida*.

MATERIAL AND METHOD

Study design and setting – A prospective study was carried out in the

department of Microbiology, NIMS Medical College and Hospital, Jaipur (Rajasthan) from January 2017 to May 2017. A total of 50 non repetitive clinical isolates of *Candida* obtained from various clinical samples, which were received from different Wards, ICUs and OPDs of the hospital and were submitted to microbiology laboratory with all aseptic precautions were included in the study.

Specimen processing⁷

1. Direct examination of sample- Direct microscopic examination by 10% KOH mount and Gram staining reveals presence of oval budding yeast cells with / without pseudohyphae⁷.

2. Culture – All the clinical samples were inoculated on to Blood agar and SDA, both were incubated at 37°C for 48-72 hrs. All the samples which showed growth were identified by colony characteristics and by gram staining. Once the conformation of colonies was done, they were further speciated⁷.

3. Speciation of candida isolates (Fig 1-8)

(a) Standard conventional method - *Candida* isolates were further speciated by standard method which includes germ tube test, growth pattern on Corn Meal agar, sugar fermentation and assimilation test⁷.

(b) Growth on chromogenic agar – Isolated *Candida* species were also inoculated on to the chromogenic media (HiCrome Candida differential agar (M1297A); Himedia, Mumbai, India.) Inoculated plate was incubated at 37°C for 24 hours. These plates were further incubated for 48 hrs to get better well developed colored colonies. The color of the colony was interpreted as per color code provided on chromogenic media (Table 1) by two observer to avoid subjective variation¹¹. All the isolates with doubtful morphology were considered unidentified and excluded from the study.

RESULT

In the present study, a total of 51 (4.5%) *Candida* species were isolated from 1113 clinical specimens. In our study conventional method was considered gold standard method for speciation. Out of 51 *Candida* isolates, most common isolated species was Non-*albicans* *Candida* ie 28 (55%) followed by *Candida albicans* ie 23 (45%) (Fig 4). Out of 28 Non-*albicans* *Candida*, *C. tropicalis* was most frequently isolated (33%) followed by *C. krusei* (18%), *C. parasilosis* (4%) (Fig 9).

These 51 strains were also identified using HiCrome Candida differential agar. 49 *Candida* species which were correctly identified by HiCrome Candida agar except 2 species of *C. parasilosis* (identified by conventional method) which were identified as *C. glabrata* by HiCrome Candida agar (Fig 10). 100% sensitivity and specificity was observed for *Candida albicans*, *Candida tropicalis*, *Candida krusei*. 0% sensitivity and 100% specificity for *Candida parasilosis* (Table 2).

DISCUSSION

In the present study, out of 51 *Candida* isolates, 23(45.1%) were identified as *C. albicans* and 28 (54.9%) were NAC species. Our study showed that Non albicans *Candida* were isolated at higher rate (54.9%) than *C. albicans* (45.1%), which was in agreement with the studies conducted by Saroj Golia et al¹⁷ and N.Pahwa et al¹⁹ who also showed the Non albicans *Candida* incidence (64% and 58% respectively) to be higher than that of *C. albicans*. This change in pattern has been partly attributed to increased immune suppression resulting in higher number of susceptibility in immunocompromised patients and also due to the prophylactic use of antifungal agents in critically ill patients. Hospitalization (especially in ICU), placement of central venous catheters and the other indwelling devices, previous antimicrobial therapies have played significant role in this changing pattern of Candidiasis^{22,23}.

In our study, the most frequently isolated species was *C. albicans* in 45.1% of the infections, followed by *C. tropicalis* 33.3%, *C. krusei* 17.6% and *C. glabrata* 3.92% respectively. Hence *C. tropicalis* (33.3%) was the second most common species reported in the present study. This finding was comparable with other workers like L. Sumitra Devi et al.¹⁸ (22.9%), Divya Dadhich et al.²⁰ (26.4%), and Khan and Bobade O, et al.²¹ (21%) [table3].

In the present study, HiCrome agar showed 100% sensitivity and 100% specificity for *C. albicans*, *C. tropicalis* and *C. krusei*. Our study agrees with that of the studies conducted by D. Dadhich et al²⁰ and Nayak et al²⁴, who also showed cent percent sensitivity and specificity to these *Candida* species when compared with conventional method.

In our study low sensitivity was reported for *C. parasilosis* which correlates well with the study conducted by tha of Baradkar et al¹¹.

These values are in contrast to the study conducted by various other scientists^{24,25}. As both the species of *C. parasilosis* showed characteristic morphology on corn meal agar. Hence a combination of HiCrome and Corn meal agar can identify this *Candida* species with in 48 hours of culture.

CONCLUSION

On completion of this study it is concluded that *Non- albicans Candida* which was earlier considered to be non-pathogenic has emerged as important pathogen. It can no longer be discarded as a lab contaminant. Hence speciation of *Candida* species is of utmost importance in the present clinical scenario.

Speciation of *Candida* using conventional methods is quite cumbersome and time consuming. Therefore species level identification using HiCrome *Candida* agar medium is advocated. The advantages of HiCrome *Candida* agar is that it is easy to prepare i.e. boiling and dispensing in petri plates. It is technically simple and cost effective compared to technically demanding and time consuming conventional methods. It also facilitates identification of two or more different species present in a single clinical sample as is seen in our study. As a result it can be concluded that the use of HiCrome agar *Candida* is an easy reliable method for presumptive identification of most of the *Candida* species Furthermore, the species level identification of the *Candida* isolates along with their antifungal susceptibility patterns can greatly influence the treatment options for the clinician and may have an impact on the patient care.

Table 1: Colour of different Candida species on Hicrome agar

Species	Color of the colony
<i>C. albicans</i>	Light-green
<i>C. dubliniensis</i>	Dark-green
<i>C. glabrata</i>	Cream to white
<i>C. krusei</i>	Pink, fuzzy
<i>C. parasilosis</i>	White to Cream
<i>C. tropicalis</i>	Blue to purple

Table 2: Sensitivity and specificity of HiChrom agar for identification of various Candida spp.

Candida spp	No.of Candida spp. identified by conventional method	No.of Candida spp. identified using HiCrome agar	Sensitivity of HiCrome agar	Specificity of HiCrome agar
<i>C. albicans</i>	23	23	100%	100%

<i>C. tropicalis</i>	17	17	100%	100%
<i>C. krusei</i>	9	9	100%	100%
<i>C. parasilosis</i>	2	0	0	100%

Table-3 Candida species isolated in various studies

S. No.	Author	Year, Place	<i>C. albicans</i>	<i>C. tropicalis</i>	<i>C. Krusei</i>
1	Badiee P ¹⁶	Iran, 2011	48%	1.7%	16.1%
2	Saroj Golia ¹⁷	Karnataka, 2013	35.71%	26.78%	4.46%
3	L. Sumitra Devi ¹⁸	Haryana, 2014	51.6%	25%	16.6%
4	N Pahwa ¹⁹	Indore, 2014	42.2%	22.4%	3.4%
5	Divya Dadhich ²⁰	Kota, 2016	54%	22%	6%
6	Present Study	Jaipur, 2017	45.1%	33.3%	17.6%

Fig 1 –Microscopic appearance of various Candida species on Corn meal agar

- (a) *C. albicans* - elongated pseudohyphal cells with terminal chlamydospores.
- (b) *C. tropicalis* – blastoconidia formed along the abundant long branching pseudohyphal cells.
- (c) *C. krusei* - chains of blastoconidia extend from the junction of pseudohyphal cells.
- (d) *C. parasilosis* - blastoconidia occur singly, in clusters and in short chains along the pseudohyphae.

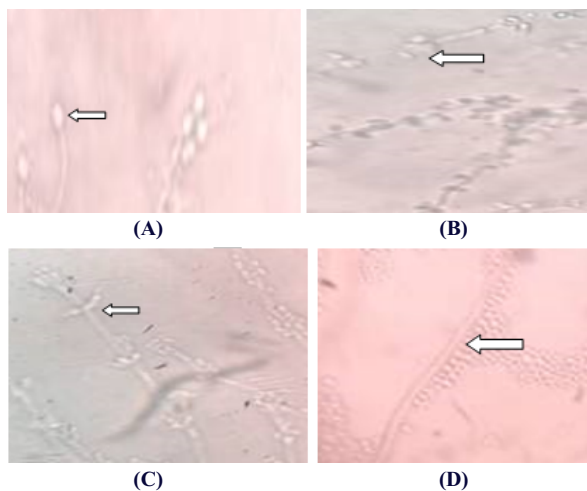


Fig 2 Differentiation of various species of Candida on HiCrome agar

- (a) *C. albicans* (b) *C. tropicalis* (c) *C. krusei*(d) *C. parasilosis*

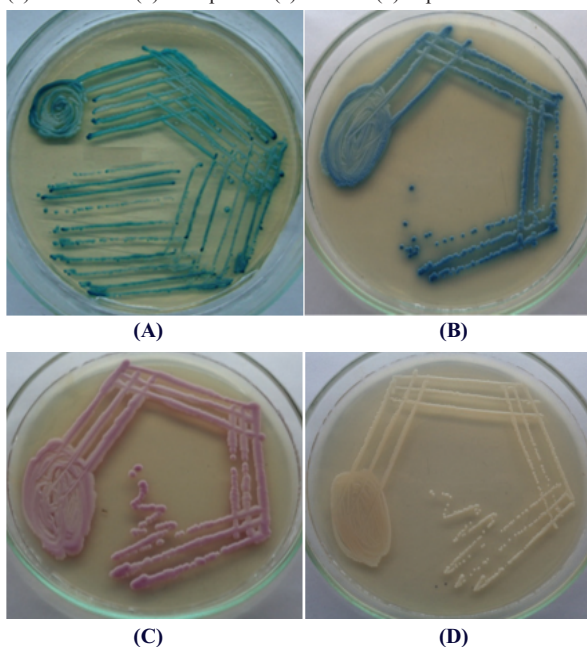
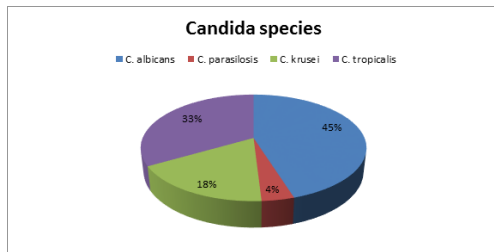
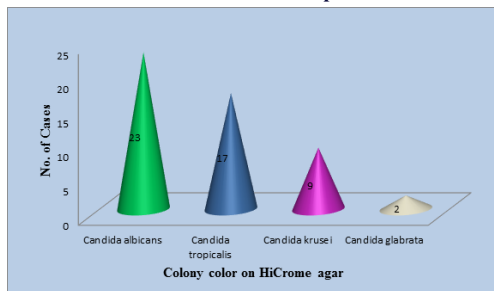


Figure 9 : Isolation of different species of Candida by Conventional methods**Figure 10 Isolation of different Candida species on HiCrome agar****REFERENCES**

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