

ORIGINAL ARTICLE

Resistance of Oral *Candida albicans* Infection to Fluconazole and Nystatin among Healthy Persons after Treatment with Azithromycin and Hydroxychloroquine to Treat Suspected SARS-COV-2 Viral Infection

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ABSTRACT

Key words:

COVID-19; Antimicrobial Resistance; Azithromycin; Hydroxychloroquine

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Background: Oral candidiasis is considered the most common fungal infection in humans. Polyene or azole antifungal drugs can be used for *Candida* infection. In addition, Nystatin therapy is used in mild forms of oral candidiasis. However, the current literature reports conflicting results regarding azithromycin and hydroxychloroquine for COVID-19. **Objective:** We aim to evaluate the resistance of oral *Candida albicans* to fluconazole and nystatin in healthy individuals who have received therapy for suspected COVID-19 with azithromycin and hydroxychloroquine. **Methodology:** The current case-control study collected samples of oral candidiasis from two types of patients. The cases group (group 1) were patients with confirmed COVID-19; the control group (group 2) were healthy controls without previous COVID-19. The antifungal susceptibility was examined based on the principles of M44-A suggested by the Clinical and Laboratory Standards Institute (CLSI). **Results:** Thirty patients were included in the cases group. In terms of the antifungal susceptibility between the study groups, the study showed the details of fluconazole and nystatin resistance. Fluconazole resistance was detected significantly higher in 86.7% of the cases group compared with only 56.7% of the subjects in the control group ($P=0.010$). Similarly, nystatin resistance was significantly higher in 76.7% of the cases group, while the control group showed 23.3% drug resistance ($P< 0.001$). **Conclusion:** Treating COVID-19 patients with hydroxychloroquine and azithromycin negatively impacted the antimicrobial resistance (AMR) of oral candidiasis. Given the limited efficacy of both treatments, avoiding the broad-spectrum use of both agents in managing COVID-19 patients is advisable.

INTRODUCTION

Candida species are widely distributed in nature and are normal microflora of the skin and mucous. They can cause opportunistic infection, especially among immunocompromised people. Oral candidiasis is considered to be the most prevalent fungus infection in humans. Over 80% of oral candidiasis infections are caused by *Candida krusei*, *Candida albicans*, *Candida glabrata*, and *Candida tropicalis* infection represents the most common species isolated from the oral mucosa, found in 30% to 50% of neonates, children, and adults.¹

Cancer patients receiving chemotherapy and radiation therapy frequently develop oral candidiasis, representing an incidence of 90% to 95%.^{2,3} Oropharyngeal candidiasis commonly affects people with acquired immunodeficiency syndrome (AIDS) as they are incompatible with highly active antiretroviral therapy.⁴ When *Candida* penetrates the surface

epithelium, it causes severe systemic manifestations associated with substantial morbidity and mortality rates (79%); hence, efficient diagnosis and management are required.^{5,6}

Treating patients with oral *Candida albicans* infection often include polyene or azole antifungal drugs.⁷ Azoles drugs, such as fluconazole, are associated with better solubility and less nephrotoxicity than polyenes; however, they are compromised because of azole resistance, especially with *Candida albicans*.⁸ In addition, nystatin is used in mild forms of oral candidiasis and can be used in cases of azole resistance.^{9,10} Several mechanisms of antifungal drug resistance have been studied, mostly in vitro.¹¹⁻¹³

The severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), primarily reported in China in December 2019, is the cause of a worldwide pandemic, infecting more than 10 million persons, with a mortality rate of 1.4-6.9%.¹⁴ Opposing results regarding the use of azithromycin and hydroxychloroquine have been

reported. A prospective study enrolling 11 patients found no significant clinical benefit for hydroxychloroquine and azithromycin in patients with SARS-CoV-2.¹⁵ In contrast, another study found that hydroxychloroquine was associated with a significant viral load reduction on day 6 (70% [14/20]) compared to control (12.5% [2/16]).¹⁶

Therefore, the current study aims to assess the resistance of oral *Candida albicans* to fluconazole and nystatin among healthy persons after treatment with azithromycin and hydroxychloroquine for suspected COVID-19.

METHODOLOGY

Setting and patients:

The present study procedure was planned following the declaration of Helsinki's established principles and was approved by the Hospital's Ethical Committee at Misr University for Science and Technology (MUST). Participants were allocated from the University Hospital's Outpatient Dermatological Clinic. Samples from two different patient types with oral candidiasis were obtained for this observational investigation. Patients with confirmed COVID-19 who matched the following two major criteria comprised the first category (cases group): (1) Age above 12 years, (2) prior SARS-CoV-2 infection confirmed by PCR, (3) treatment with hydroxychloroquine and azithromycin were the first two criteria. The second kind, or control group, consisted of healthy patients who had never had COVID-19.

Sample collection:

We included participants who had not received antifungal therapy for the past three days to prevent any false negative results. Samples were collected using sterile swabs during the visit to the Outpatient Clinic. Samples were washed in distilled water and transported to the laboratory for further analysis. Samples were stored on Sabouraud Dextrose Agar culture media at 37°C. After 48-72 hours, the cultured plates were assessed. First, slides were developed from the structured colonies. *Candida albicans* organisms were identified by the standard methods if we found yeast cells.¹⁷

Antifungal susceptibility testing:

Antifungal susceptibility was evaluated based on the principles of M44-A proposed by the Clinical and Laboratory Standards Institute (CLSI). The standard strain of *Candida albicans* was ATCC 2091. The

isolates were subcultured overnight at 37°C on Sabouraud Dextrose Agar (SDA). The organism was suspended in saline, and the turbidity was set at 0.5 McFarland standards. The plates were loaded with the media and incubated at 37°C with the discs on top. After 24 and 48 hours, the zone of inhibition was captured on camera. Fluconazole 10 mg/disk and 100 units/disk of nystatin were utilised (Sigma Aldrich, USA).¹⁸

Statistical analysis:

Data were analysed using SPSS software, version 16. Chi-square was used to compare cases and controls regarding gender, Fluconazole resistance, and Nystatin resistance. In addition, T-test student was used for comparison between groups regarding age. A P-value of less than 0.05 was considered statistically significant.

RESULTS

During the study period, thirty patients were included in the case group. The patients were confirmed previous COVID-19 infection and history of hydroxychloroquine and azithromycin treatment during the management period of COVID-19 infection. The cases group showed an average age of 37± 9 years old. 36.7% of the included patients were males, and 63.3% were females. In the control group, the average age was 33 ± 8 years old. 46.7% of the included patients were males, and 53.3% were females. There was no significant difference between cases and control groups regarding age and gender distribution (P = 0.094 and P = 0.432, respectively). (Table. 1)

Table 1: Demographics Characteristics of the studied participants.

Parameters	Cases (n= 30)	Controls (n= 30)	P-value
Age, mean (± SD)	37 (9)	33 (8)	0.094
Gender			
Male	11 (36.7)	14 (46.7%)	0.432
Female	19 (63.3)	16 (53.3%)	

Regarding antifungal susceptibility between the study groups, Fluconazole resistance was detected significantly higher in 86.7% of the cases group compared with only 56.7% of the subjects in the control group (P = 0.010). Table. 2 shows the details of the fluconazole and nystatin resistance.

Table 2: Clinical data of the studied participants.

Parameters	Cases (n= 30)	Controls (n= 30)	P-value
Fluconazole resistance			0.010
Sensitive	4 (13.3%)	13 (43.3%)	
Resistant	26 (86.7%)	17 (56.7%)	
Nystatin resistance			< 0.001
Resistant	23 (76.7%)	7 (23.3%)	
Sensitive	7 (23.3%)	23 (76.7%)	

Similarly, nystatin resistance was significantly higher in 76.7% of the cases group, while the control group showed 23.3% of the drug resistance (P < 0.001) **Figure 1**.

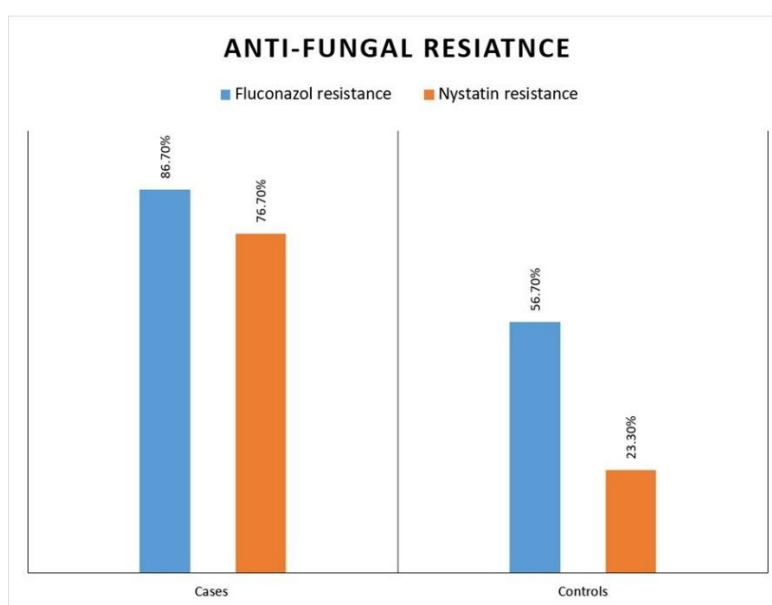


Fig. 1: Antifungal resistance in both cases and control groups

DISCUSSION

Antimicrobial resistance (AMR) is one of the long-term complications of the COVID-19 pandemic due to the suboptimal or inappropriate use of antimicrobials in the acute care setting.¹⁹ Recent studies reveal that fungal and bacterial co-infection is minimal in COVID-19 patients; however, broad-spectrum antimicrobial prescribing rates and use are substantial.²⁰ These prescribed antimicrobial agents include antibiotics such as azithromycin, antimalarial agent as hydroxychloroquine, and antivirals like remdesivir, ritonavir, and lopinavir.^{16,21} Many reports showed that these agents are used empirically without any confirmed diagnosis.²²⁻²⁴ Azithromycin is also extensively used with hydroxychloroquine, according to the WHO, even though they are not currently approved outside of COVID-19 clinical studies.²⁵ The inappropriate use of

these antimicrobial agents may lead to a serious condition, secondary bacterial infection, which was linked with a high mortality rate in patients with COVID-19.²⁶ Moreover, it was observed that patients with COVID-19 who were treated at crowded spaces like nursing homes and psychiatric hospitals were associated with highly resistant microbes, including vancomycin-resistant enterococcus, Methicillin-resistant *Staphylococcus aureus*, and carbapenemase-producing organisms, and fungi as *Candida*.²⁷

Generally, the risk of invasive *Candida* infection is very high in patients with critical illness who received an extensive course of antibiotics. This risk is directly proportionate with the type, duration, and dose of antibiotic.²⁸ Jensen et al.²⁹ showed that patients who received high-dose antibiotics were associated with a higher rate of invasive oral *Candida* than those who received low-dose (HR= 1.9, 95% CI: 1.01 – 3.6). In addition, the risk was substantially higher in patients

who received antibiotics for three days than those who used the high-dose antibiotics for 1-2 days (HR= 3.8, 95% CI: 1.6 – 9.3). In COVID-19 patients, Wang et al.³⁰ showed that out of 69 patients with COVID-19, 29 were subjected to sputum culture, but only five patients were associated with co-infection; of them, two patients had *Candida albicans*. Invasive candidiasis was shown to be prevalent in 14.1% of COVID-19 patients with ARDS, according to research conducted in the United Kingdom.³¹ In critically ill patients with COVID-19, three cases of candidemia were observed after receiving an IL-6 inhibitor, tocilizumab.³² Moreover, several case reports have documented the *Candida* co-infection in patients with COVID-19.³³

Our findings showed that fluconazole resistance was significantly higher in 86.7% of the cases group compared with only 56.7% of the subjects in the control group ($p= 0.010$). Similarly, nystatin resistance was significantly higher in 76.7% of the cases group, while the control group showed 23.3% drug resistance ($P < 0.001$). Shirvani and his colleagues³⁴ conducted another study to isolate *Candida* species from individuals with the COVID-19 disease and evaluate the susceptibility pattern of *Candida* species to routine antifungal drugs. The detected species showed significant resistance against Itraconazole and Fluconazole; however, the Amphotericin B and Voriconazole responded well. Their findings demonstrated that *Candida albicans* and *Candida parapsilosis* were the only detected species (96% vs 4%). In prior studies, multiple azole, multidrug, and fluconazole were found in 30%, 40%, and 100% of *Candida auris* isolates, respectively.^{35,36} Oral candidiasis can travel from the oropharynx to the oesophagus or systemically through the bloodstream or upper gastrointestinal tract when therapy is not given or is ineffective, such as in fluconazole-resistant *Candida* or immunocompromised people.³⁷ The resultant candidemia has been proven to cause considerable morbidity and death, with a 71–79% mortality rate.³⁸ An investigation in Spain found that COVID-19 individuals had candidemia, candiduria, and severe intraabdominal candidiasis.³⁹ As a result, it's critical to consider how these individuals will be treated in a hospital setting. The use of AuNPs, nanoparticles of gold, was proposed in cases of azole resistance in patients with invasive *Candida*.⁴⁰

To the best of our knowledge, this study is the first study to assess the resistance of oral *Candida albicans* to fluconazole and nystatin among healthy persons after treatment with azithromycin and hydroxychloroquine for suspected COVID-19. However, we acknowledge that our study has some limitations, including the small sample size and the single-centre setting, which may hinder the generalizability of our findings. In addition, we studied only one species of *Candida albicans*; however, according to the literature, it is the most prevalent species in COVID-19 patients.

CONCLUSION

Treating COVID-19 patients with azithromycin and hydroxychloroquine negatively impacted the antimicrobial resistance of oral candidiasis. Given the limited efficacy of both treatments, it is advisable to avoid the broad-spectrum use of both agents in managing COVID-19 cases. Azithromycin and hydroxychloroquine had a subsequent negative impact on the antimicrobial resistance of oral candidiasis. The resistance to Fluconazole and Nystatin was detected significantly higher in cases groups compared with the control group. The study results recommend avoiding broad-spectrum use of Azithromycin and Hydroxychloroquine agents in the management of COVID-19 cases with *Candida Albicans*.

Recommendations

Further large-scale studies are recommended to investigate more species of *Candida* and their response with broad-spectrum of antifungal agents.

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This manuscript has not been previously published and is not under consideration in the same or substantially similar form in any other reviewed media. I have contributed sufficiently to the project to be included as author. To the best of my knowledge, no conflict of interest, financial or others exist. All authors have participated in the concept and design, analysis, and interpretation of data, drafting and revising of the manuscript, and that they have approved the manuscript as submitted.

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