

REVOLVING LABORATORY-STEWARDSHIP FOR PHENOTYPIC RESISTANCE PROFILES OF COMBINED ANTIFUNGAL AGENTS ON HUMAN CLINICAL *CANDIDA* SPECIES IN A DEVELOPING COUNTRY AND COVID-19 INDICATIONS

Adenike Ogunshe^{1*}, Yetunde Ekanola², Maryam Omolaja² and Tiwalade Adewale²

¹Pegasus-Zion Community & Environmental Health, Nigeria. ²Department of Biology, The Polytechnic, Ibadan, Oyo State, Nigeria *Correspondence author E-mail: adenikemicro@gmail.com; adenikemicro@yahoo.com ORCID 0000-0002-7741-9104

ABSTRACT: Background: Antimicrobial stewardship has always revolved round prudent antimicrobial regimens, including combined antimicrobial therapy, to curb antimicrobial resistance. But there is dearth of local microbiological data on combination antifungal therapy. Materials and Methods: 129 strains of human oral and vulvo-vaginal Candida species were assayed for phenotypic susceptibility / resistance profiles to 12 commonly-available-incountry antifungal drugs and 12 antifungal creams. **Results:** When tested singly, C. albicans strains were totally resistant to Primpex and Flucamed. Resistance rates of 46.6-95.2% were exhibited to remaining antifungal drugs, with multiple antibiotic resistance (MAR) of 41.7-100%. All C. glabrata strains were resistant to Fesovin; 55.7-96.2% strains exhibited resistance to other antifungals, and MAR was 41.7-100%. C. pseudotropicalis strains were totally resistant to Fesovin, Ketoconazole, Primpex and Flucamed; 36.8-84.2% strains resisted other antifungals, while MAR was 41.7-100%. C. tropicalis, exhibited resistance rates of 37.5-100% and MAR of 50.0-100%. The Candida species were resistant to all antifungal creams. Combined phenotypic assays of five most-resisted antifungal drugs and six most-resisted antifungal creams, on 40 most-resistant Candida strains gave 37.5% and 27.5% resistance (or 62.5% and 72.5% susceptibility) instead of 100% resistance (0.0% susceptibility) recorded during single antifungal drug and cream susceptibility assays. Conclusion: Appropriate laboratory-based direct-antifungal-drugs assay can serve as antimicrobial stewardship intervention, which can enhance ideal antifungal prescriptions that translates to better treatment regimens of mycotic infections, and confirm non-potent antifungals, including adulterated, fake, substandard or expired antifungals, common in some developing countries, including Nigeria.

KEYWORDS: Combined Antifungal Therapy, Commonly Available-in-Country Antimicrobials, CoviD-19, Mycoses, Laboratory-Derived Antifungal Prescriptions, Substandard Medications.

INTRODUCTION

Many species of *Candida*, once considered as minor pathogens have potentials of causing diseases commonly known as candidiasis, which range from non-life-threatening mucocutaneous illnesses to invasive conditions that may involve virtually any organ.¹⁻⁸



Globally, fungal-associated mortality is reportedly more than three times that of malaria, with an estimated 1.6 million deaths annually⁴. Even, most recent evidences on emerging clinical reports infer that severe and adverse CoviD-19 complications may be caused by fungal co-infections or secondary infections, such as oral candidasis, lung candidasis, and other respiratory system candidasis, etc., which is another important factor influencing reported mortalities.⁹⁻¹⁷ However, in addition to antifungal agents like azoles, polyenes, echinoderms, etc., candidasis have also been treated with various agents, such as, phytotherapeutic products, probiotics, as well as, disinfectants and germicides, especially for surface cleansing.¹⁸⁻²⁷ Meanwhile, antifungal resistance has become increasingly recognised as a major health concern, and significant increase in resistance to antifungal agents by pathogenic *Candida* species and associated effects have been repeatedly documented.^{22, 24, 27-30}

Past few decades have observed sustained medical implications of opportunistic infections, especially due to significant increase in incidence of invasive fungal infections, and worldwide increase in number of immuno-compromised patients, who are highly susceptible to opportunistic fungal infections.^{9, 10} But antifungal armamentarium for treatment of serious fungal infections remains limited, mostly in developing countries. It has therefore, been proposed that a possible approach to improve treatment outcomes, and overcome antifungal drug resistance and high mortality rates seen in severe fungal infections is, combination therapy, which can lead to increase in susceptibility and reduce resistance of microbial pathogens.^{10, 30-31}

Fungal diseases affect over a billion people, and kills more than 1.5 million,⁴ as most medically important *Candida* species overcome a broad range of host-imposed constraints, to increase their pathogenicity.^{32, 33} Thus, there has been increasing interest in the use of combination antifungal therapy, especially if the drugs have different mechanisms of action, and sometimes, combinations of new agents, in order to improve prognosis and treatment of fungal diseases.³⁴⁻³⁷ Research data on antifungal combination therapy are however, quite poorly reported in developing countries, including Nigeria, where increased adulterated and sub-standard antifungal agents are also common. The aims of this study therefore, is to investigate inhibitory potentials of combined in-country antifungals, using multi-resistant *Candida* species.

MATERIALS AND METHODS

Collection of Clinical Specimens:

Oral *Candida* species [*Candida glabrata* (52), *Candida albicans* (43), *Candida pseudotropicalis* (19) and *Candida tropicalis* (15)] used in this study were stock strains, originally obtained from human specimens at the Department of Medical Microbiology & Parasitology, University College Hospital (UCH), Ibadan, Nigeria. Stock strains were reactivated in sterile peptone water, and incubated at 27-30^oC for 24-48 hours before separately sub-cultured on sterile Sabouraud D extrose A gar (SDA, Lab M, Basingstoke, England), to which ofloxaciline antibiotics was added, to inhibit bacterial contaminations. Culture plates were incubated at 27-30^oC for 24-48 hours, and obtained pure culture of each *Candida* strain was then checked to confirm identity.



Determination of anti-candidal activities of antifungal agents against *Candida* strains (Modified Agar Well-Diffusion method):

Antifungal drugs used for bioassay with their respective active ingredients were Itranox (itraconazole); Fesovin (Griseofulvin); Gyno-Tiocosid (Ticonazole); Medcan (Fluconazole); Vulcan-50 (Fuconazole); Grufin (Griseofulvin); Lucon (Fluconazole); Grisovid-500 (Griseofulvin); Diflucan (Fluconazole); Ketoconazole (Ketoconazole); Primpex (Sulphamethoxazole): Flucamed (Fluconazole). Antifungal creams used for bioassay studies and their respective active ingredients were- Mycoten (Clotrimazole), Skmeal (Ketoconazole), Trosyd (Ticonazole), Lamisil (Terbinafine hydrochloride), caneXcream (Clotrimazole), Funbact-A (Clotrimazole, Betamethasone, Neomycin sulphate, and Clobetasol propionate), Fungur (Miconazole), Nizoral (Ketoconazole), Fungusol (Miconazole Nitrate), Ketofung (Ketoconazole), Whitfield's ointment (Salicylic acid and benzoic acid), Dakatrin (Miconazole nitrate).

Holes measuring 6.00 mm in diameter were aseptically bored and punched out of sterile SDA agar plates, followed by surface flaming of the agar plates. Each SDA agar plate was then seeded with each oral *Candida* strain. Using the modification of Tagg *et al.*³⁸ method, 500 μ l suspensions of each antifungal agent were separately dispensed into agar wells in seeded plates, followed by incubation at 30^oC for 24-48 hrs. Inhibitory activities depended on release of diffusible inhibitory metabolites from antifungal agents into assay agar, during incubation. Inhibitory zones surrounding the agar wells were recorded in mm diameter, while absence of zones or zones less than 10.0 mm in diameter were recorded as negative (resistant).

Candida strains that were totally resistant to antifungal drugs and creams were assayed for susceptibility trends to combined antifungal drugs and combined antifungal creams. Holes measuring 6.00 mm in diameter were aseptically bored and punched out of sterile SDA agar plates, followed by surface flaming of the agar plates. Each SDA agar plate was then seeded with each selected oral *Candida* strain, and 500µl suspensions of each set of combined antifungal drugs [afd1+afd2; afd1+afd3; afd1+afd4; afd1+afd5 / afd2+afd3; afd2+afd4; afd2+afd5 / afd3+afd4; afd3+afd5 / afd4+afd5 / afd1+afd2+afd3; afd1+afd2+afd4; afd1+afd2+afd5 / afd1+afd3+afd4; afd1+afd3+afd5; afd2+afd3+afd4; afd2+afd5 / afd2+afd3+afd5 / afd3+afd5

Bioassay was also determined on antifungal creams [afc1+afc2; afc1+afc3; afc1+afc4; afc1+afc5; afc1+afc6 / afc2+afc3; afc2+afc4; afc2+afc5; afc2+afc6 / afc3+afc4; afc3+afc5; afc3+afc6 / afc4+afc5; afc4+afc6 / afc5+afc6 / afc1+afc2+afc3; afc1+afc2+afc4; afc1+afc2+afc5; afc1+afc2+afc6 / afc1+afc3+afc4; afc1+afc3+afc5; afc1+afc3+afc6 / afc1+afc4+afc5; afc1+afc4+afc6 / afc1+afc5+afc6 / afc2+afc3+afc4; afc2+afc3+afc5; afc2+afc4+afc5; afc1+afc2+afc6; afc2+afc3+acf6; afc2+afc5; / afc1+afc3+afc4; afc1+afc3+afc5; afc1+afc3+afc6 / afc1+afc4+afc5; afc1+afc4+afc6 / afc1+afc5+afc6; afd2+afd3+afd5 / afd2+afd4+afd5 / afd4+afd5] and separately dispensed into wells in seeded agar plates, followed by incubation at 30°C for 24-48 hrs. Inhibitory zones surrounding the agar wells were recorded in mm diameter, while absence of zones or zones less than 10.0 mm in diameter were recorded as negative (resistant).



RESULTS

As shown in Table 1, all the 42 oral C. albicans strains were totally resistant to Primpex [Sulphamethoxazole] and Flucamed [Fluconazole] drugs, while resistance rates of 38.1-95.2% were exhibited towards the remaining antifungal. All the 52 C. glabrata were resistant to Fesovin [Griseofulvin] drug, while 44.2-98.1% of C. glabrata strains exhibited resistance towards other antifungals but slightly less resistant to Vulcan-50 [Fluconazole] and Gruffin [Griseofulvin] (40.4-44.2%). C. pseudotropicalis strains were totally resistant to Fesovin [Griseofulvin], Ketoconazole, Primpex [Sulphamethoxazole] and Flucamed [Fluconazole]; significantly resisted (36.8-84.2%) by other antifungals, but least recorded resistance were 15.8% and 21.1% for Vulcan-50 [Fluconazole] and Gruffin [Griseofulvin]. Exhibited resistance rates for C. tropicalis were 31.3-100%.

Percentage of multiple antifungal resistance (MAR) exhibited by C. albicans, C. glabrata and C. pseudotropicalis strains were 41.7 - 100%, while MAR for C. tropicalis was 50.0-100%. Most of the Candida species displayed over 50.0% antifungal resistance (Table 2). None of the Candida strains of all the species was inhibited by any of the single antifungal cream (Table 3).

Table 4. shows the results obtained when 40 selected Candida strains that were totally resistant to five antifungal drugs (Gyno-tiocosid [Ticonazole], Medcan [Fluconazole], Primpex [Sulphamethoxazole], Ketoconazole [Ketoconazole] and Fesovin [Friseofulvin]) were screened for phenotypic susceptibility, as combined antifungal drugs, as well as, six combined antifungal creams (Whitfield's ointment, Dakatrin [Salicylic acid and Benzoic acid), Dakatrin [Miconazole nitrate], Nizoral [Ketoconazole], Mycoten [Clotrimazole], Funcbact-A [(Clotrimazole, Betamethasone, Neomycin sulphate, and Clobetasol propionate] and Lamisil [terbinafine hydrochloride]). 100% resistance recorded during single antifungal drug and cream susceptibility testing but 37.5% and 27.5% resistance rates respectively were recorded in the combined antifungal drug and cream susceptibility testing (Figure 1). Nine strains of the oral Candida species were minimally inhibited (11.0-19.0 mm in diameter); 13 strains were moderately inhibited (21.0-29.0 mm in diameter), while three strains were highly inhibited (≥30.0 mm in diameter) by five combined antifungal drugs. Eight strains of the oral Candida species were minimally inhibited (11.0-19.0 mm in diameter); 18 strains were moderately inhibited (21.0-29.0 mm in diameter), while two strains were highly inhibited (\geq 30.0 mm in diameter) by the six combined antifungal creams (Table 4).

DISCUSSION

Invasive infections by opportunistic Candida species, and incidence of more virulent clinically important Candida species, which represent infectious agents with higher risk for mortality (due to a variety of virulence factors that make them capable of infectivity, even, in otherwise intact immune system), has increased in recent decades.^{8, 20, 39, 40} These occur especially in people with critical underlying co-morbidities.^{40, 41} There have also been large numbers of cases that reported co-infection of mycoses with viruses, fungi, and bacteria, some of which originate from oral cavity.⁴² Antifungal profiles of Candida strains from CoviD-19 positive subjects and patients were not determined in this study, due to gross lack of biosafety for research purposes, and present biosecurity measures in handling isolations of oral fungal isolates from CoviD-19 patients, Notwithstanding, very high MAR rates [C. albicans, C.



glabrata, C. pseudotropicalis (41.7-100%), and C. tropicalis (50.0-100%)] were recorded for non-CoviD-19 clinical Candida species implicated in human oral candidasis.

Antifungal combination regimens could achieve more rapid antifungal effects, prevent emergence of antifungal resistance, and possibly allow reduction in the dosages of individual antifungal agents.⁴³⁻⁴⁶ But implication of obtained results in this study is that none of commonly-available test antifungal drugs and creams could inhibit at least 50% of just 129 clinical Candida strains of human origin. Relentless increase of invasive fungal infections and poor outcomes associated with available antifungal agents, and sometimes with non-effective actions, has therefore, prompted therapeutic strategies, through combination antifungal therapy, as there seems to be several potential advantages in antifungal combinations, in addition to widening spectrum and potency of drug activity. In this study, instead of 100% resistance rates recorded for the single antifungal drugs and creams susceptibility testing, 37.5% and 27.5% resistance rates were respectively recorded for most-resisted five antifungal drugs and six antifungal creams, combined for susceptibility testing.

Oral candidasis can be treated with antifungals belonging to different classes of drugs that target different cellular processes ⁴⁷ but it has been reported that after administration, intra-oral concentrations of antifungal drugs tend to be sub-therapeutic and transient, due to diluent effect of saliva and cleansing effect of the oral musculature. Hence, intra-orally, *Candida* may undergo just a brief exposure to antifungal drugs,²⁶ indicating further reduced potencies of oral antifungals. Furthermore, likelihood of oropharyngeal candidasis development in CoviD-19 patients, with a list of attributable risk factors for oral infections has been investigated and reported.¹⁵ It is thus, of further great concern that in this study, none of the classes of antifungal agents, as single drugs or creams had significant maximal *in vitro* inhibitory effects on most of the *Candida* species, and it was not even, until five and six antifungal drugs and creams were combined in equal proportions that inhibitory effects were observed in most of the antifungals. However, it must be borne in mind that, considering the potential life-threatening drug interactions and adverse side-effects like, kidney or liver toxicity associated with antifungal therapies, it would be clinically hazardous to administer five oral antifungals in treatment of oral candidasis because of such multiple resistance.

There has always been a growing universal concern regarding counterfeit medications, particularly, because counterfeit antimicrobial drugs are considered as threat to overall public health, with many devastating consequences for patients, like, increased mortality and morbidity, long duration of antimicrobial therapy, treatments failures, emergence of drug resistance, etc.⁴⁷ Apart from concerns on fake, adulterated or sub-standard antimicrobial drugs,^{33, 48} another factor of special concern is the issue of inconsistencies in bacteriostatic and/or bactericidal activities of antifungals produced as different batches.⁴⁹⁻⁵¹ These factors additionally contribute to relatively high rates of recorded antifungal resistance in several studies, including the present study. In countries with standard drug regulations, less-quality or adulterated drugs cannot be expressly imported, while fake, adulterated and sub-standard drug batches are rejected and/or recalled from the markets but this is not usually the case in Nigeria's situation. So, it is very difficult to regulate drug batches that do not meet standard quality criteria, and of course, not surprising that significant resistance rates were recorded in this study, even, for some combined antifungals, which is of severe clinical disadvantage in combined antifungal therapy.

African Journal of Biology and Medical Research ISSN: 2689-534X Volume 3, Issue 3, 2020 (pp. 96-110)



In order to curb the global menace of antifungal resistance, there are suggestions of development of new antifungal agents, and with novel modes of actions.⁵² But, even, if newly developed antifungal drugs are effective and safe for treatments of topical and invasive fungal infections, and also active against fungal strains that are resistant to current antifungal drugs or if there is upgrading of antifungal armamentarium, through improvement in existing antifungals and/or novel antifungal strategies are devisied;⁷ such therapies are very unlikely to be commonly-unavailable in developing countries. This is due to high costs and deficient healthcare policies, being the reason that most antimicrobials in such countries are generic, and not original brands / trademarks, and their pharmacological effects are mostly not exactly the same as those of their brand-name counterparts. There had also been reports of manufacturing of some poor-quality drugs (which cannot be sold or prescribed in countries of manufacture), particularly prepared for developing countries, as some unscrupulous business merchants, specifically request for production of poor-quality medications, which they import into developing countries like Nigeria.⁵¹ Another peculiar missing link about antimicrobial resistance in Nigeria, and possibly few other developing countries, is deliberate change of expiration dates labels on clinical medications.⁵³

Invasive fungal infections have major impacts on human morbidity and mortality, which are responsible for almost over a million deaths yearly, and even, with increasing rates; since many fungal species cause infections, with mortality rates exceeding 50%.⁵⁴ Fungal co-infections and their impact on CoviD-19 patients are still understudied, ^{12, 52, 54} but it is of more recent safety concerns that severe CoviD-19 infection alone can cause damage to multiple organ systems.^{54, 55} Thus, adequate antifungal dosing strategies are necessary, to ensure successful mycotic treatments,¹⁵ following antifungal susceptibility testing, which is crucial for ensuring appropriate antifungal therapy for patients' optimal treatment options.⁵⁶ In consideration of overall challenges on antimicrobial resistance facing many developing countries, this study tried to apply a standardised, easily-reproducible, and clinically-applicable susceptibility testing method for *Candida* strains, for detection of antifungal resistance, through direct choice of most-potent antifungal agents.

In spite of mycotic co-morbidity, Africa has been quite fortunate to be less affected by the severity of recent emergence of CoviD-19 pandemic, but, this should serve as a stark reminder that, there is a need for continued medical innovation, to ensure availability of safe and effective antifungal drugs, for treatment of patients with serious invasive fungal infections.¹³ The fact that fungal infections have been implicated as co-morbidities in CoviD-19 cases should attract much attention on antifungal therapy and resistance. Findings of this study also confirm some earlier submissions,^{33, 51} particularly on low quality and substandard/counterfeit antimicrobial drugs in developing countries, which can lead to increased morbidity, mortality, as well as, emergence of antimicrobial resistance, especially by invasive and non-invasive Candida strains in the environment.⁵⁷ There must also be consideration for emergence of antifungal-resistant *Candida* strains displaying cross-resistance to other clinical antifungals.⁵⁸ which are of significant consequences for local and global public health. Mycotic infections have not been attracting as much attention as bacterial and viral infections in Nigeria; and for antibiotic susceptibility tests, antibiotic discs are commonly used, with inhibitory zones usually denoted as, +, ++, +++ and ++++, which are for interpretations of the increasing zones of inhibitions on agar plates. These interpreted values form basis for prescriptions of antibiotics; however, they do not translate to curative potencies by corresponding antibiotic drugs or



creams, as there can be significant differences in the susceptibility / resistance values between antibiotic-discs and corresponding antibiotic drugs.^{59, 60}

Two major indications of CoviD-19 pandemic are that tremendous attention must be focused on fungal co-infections, and qualities of antifungals in the country. Direct antifungal susceptibility method, using in-country antifungal medications can therefore, ensure, not only appropriate guidance, in the choice of antifungal therapy, but also, knowledge of local epidemiology of antifungal resistance in the country. Most importantly, direct antifungal susceptibility method can expose lack of potent activities of fake antifungals, especially, as there are no general clinical and epidemiological surveillance systems in place, to determine prevalence of antifungal resistance, as well as, fake and substandard antimicrobials in the country. Such direct assay can firstly, enhance coordinated optimisation of antimicrobial interventions, referred to as, Antimicrobial Stewardship, which promotes selection of optimal antimicrobial drug regimen; and secondly, for assessing the consistency of different batches of antimicrobial drugs, particularly with regards to potency. Complete documented investigations into failure of drug batches that fail to meet expected specifications can also be carried out by the direct assay used in the current study; thereby, aiding in appropriate policies for poor quality antifungals. So, bodies like, National Agency for Foods and Drugs Administration and Control (NAFDAC), and the Federal Competition and Consumer Protection Commission (FCCPC) must put in place, novel regulations for antimicrobials and other medications produced or imported into Nigeria, to protect patients' safety and prevent antimicrobial (antifungal) resistance.

CONCLUSION AND RECOMMENDATIONS

Significant reduction in phenotypic resistance rates were recorded for combined antifungals, compared to single antifungals; indicating that combined antifungal therapy can enhance better treatments of mycotic infections. Direct-antifungal-susceptibility-drug-assay also confirmed lack of bacteriostatic and bactericidal potentials in most of the combined test antifungal drugs and creams. Adulterated, fake, substandard or expired antifungals, as well as, inconsistencies in bacteriostatic and/or bactericidal activities of antifungals produced as different batches, are common in developing countries, including Nigeria. By the findings of this study, it is therefore recommended, a need to constantly implement antimicrobial stewardship; through, standardised, easily-reproducible, and clinically-applicable, direct-antifungal-drug-assay, for detecting phenotypic susceptibility or resistance, which can translate to ideal laboratory-derived antifungal prescribing and therapy, especially because of co-morbidities like, CoviD-19 and fungal infections.

Acknowledgements

Adenike A.O. Ogunshe acknowledges the Department of Microbiology, University of Ibadan, for part of the laboratory studies.

Competing interests

Authors declare no competing interests.



REFERENCES

- [1] Pappas PG, Rex JH, Sobel JD, Filler SG, Dismukes WE, Walsh TJ, Edwards JE. Guidelines for treatment of candidiasis. Clin Infect Dis 2004;38(2): 161–189.
- [2] Brown AJP, Gow NAR, Warris A, Brown GD. Memory in fungal pathogens promotes immune evasion, colonisation, and infection. Trends Microbiol 2019;27(3): 219–230.
- [3] Moyes DL, Wilson D, Richardson JP, Mogavero S, Tang SX, Wernecke J, *et al*. Candidalysin is a fungal peptide toxin critical for mucosal infection. Nat 2016;532:64–68.
- [4] Bongomin F, Gago S, Oladele RO, Denning DW. Global and multi-national prevalence of fungal diseases-+estimate precision. J Fungi (Basel) 2017;3(4):57.
- [5] Verma N, Singh S, Taneja S, Duseja A, Singh V, Dhiman RK, *et al.* Invasive fungal infections amongst patients with acute-on-chronic liver failure at high risk for fungal infections. Liver Int 2018;39:503–513.
- [6] Bertolini M, Dongari-Bagtzoglou A. The relationship of *Candida albicans* with the oral bacterial microbiome in health and disease. Adv Exp Med Biol 2019;1197:69-78.
- [7] Kainz K, Bauer MA, Madeo F, Carmona-Gutierrez D. Fungal infections in humans: the silent crisis. Microb Cell 2020;7(6): 143–145.
- [8] Rosati D, Bruno M, Jaeger M, ten Oever J, Netea MG. Recurrent vulvovaginal candidiasis: an immunological perspective. Micr 2020;8(2): 144.
- [9] Rodrigues ML, Nosanchuk JD. Fungal diseases as neglected pathogens: a wake-up call to public health officials. PLoS Negl Trop Dis 2020;14(2): e0H007964.
- [10] Soare AY, Watkins TN, Bruno VM. Understanding mucormycoses in the age of "omics". Front Genet 2020;11: 699.
- [11] Al-Hatmi AMS, Mohsin J, Al-Huraizi A, Khamis F. COVID-19 associated invasive candidiasis. J Infect 2020; 23:23.
- [12] Arastehfar A, Carvalho A, van de Veerdonk FL, Jenks JD, Koehler P, Krause R, Cornely OA, Perlin DS, Lass-Flörl C, Hoenig M. COVID-19 associated pulmonary aspergillosis (CAPA)—from immunology to treatment. J Fungi 2020;6(2): 91.
- [13] Kennedy C. COVID-19 and fungal superinfections: the deadly, perfect storm. *In*: Xconomy San Diego, May 5th, 2020. Available from https://xconomy.com/sandiego/2020/05/05/covid-19-and-fungal-superinfections-the-deadly-perfect-storm/ Last accessed on 2020 Sept 16].
- [14] Lansbury L, Lim B, Baskaran V, Lim WS. Co-infections in people with COVID-19: a systematic review and meta-analysis. J Infect 2020;81: 266-275.
- [15] Mohamed A, Rogers TR, Fe Talento A. COVID-19 associated invasive pulmonary aspergillosis: diagnostic and therapeutic challenges. J. Fungi 2020;6(3):115.
- [16] Zhou P, Liu Z, Chen Y, Xiao Y, Huang X, Fan X-G, Bacterial and fungal infections in COVID-19 patients: a matter of concern. Infect Control Hosp Epidemiol 2020;1–2.
- [17] Zhou F, Yu T, Du R, *et al.* Clinical course and risk factors for mortality of adult inpatients with COVID-19 in Wuhan, China: a retrospective cohort study. Lancet 2020. 395(10229): 1054-1062.
- [18] Ogunshe AAO. Laboratory investigation on candidastic potentials of bleach and toilet soaps. Annal Biol Res 2015;6 (5):15-19.
- [19] Ogunshe AO. Candidaemia or candidasis: controversy of *Staphylococcus* sexually transmitted infection? Afr J Biomed Res 2016;19:1- 5.



- [20] Ogunshe AAO, Lawal OA, Iheakanwa CI. Effects of simulated preparations of plants used in Nigerian traditional medicine on *Candida* spp. associated with vaginal candidiasis. J Ethnobot Res Appl 2009;6: 373-383.
- [21] Ogunshe AAO, Omotoso OA, Akindele TM. Soaps and disinfectants / germicides as adjunct antimycotic cleansing agents in cases of vulvovaginal candidasis. Advan Biol Res 2011;5 (6):282-290.
- [22] Ekanola YA, Ogunshe AAO, Olugbodi RJ, Ajibade JG, Ojediran YO. Comparative phenotypic susceptibility profiles of clinical isolates of *Candida* species to some antimycotic agents. Int Res J Biochem Bioinform 2014:4: 4-11.
- [23] Kean R, Sherry L, Townsend E, McKloud E, Short B, Akinbobola A, et al. Surface disinfection challenges for *Candida auris*: an *in-vitro* study. J Hosp Infect 2018;98, 433–436.
- [24] Van Daele R, Spriet I, Wauters J, Maertens J, Mercier T, Van Hecke S, Brüggemann R. Antifungal drugs: what brings the future? Med Mycol 2019;57 (Suppl_3): S328–S343.
- [25] Ahangari F, Farshbaf-Khalili A, Javadzadeh Y, Adibpour M, Sadeghzadeh Oskouei B. Comparing the effectiveness of *Salvia officinalis*, clotrimazole and their combination on vulvovaginal candidiasis: A randomized, controlled clinical trial. J. Obstet. Gynaecol. Res. 2019;45(4):897-907.
- [26] Buggio L, Somigliana E, Borghi A, Vercellini P. Probiotics and vaginal microecology: fact or fancy? BMC Women's Health. 2019; 19: 25.
- [27] Barrientos-Durán A, Fuentes-López A, de Salazar A, Plaza-Díaz J, García F. (2020). Reviewing the composition of vaginal microbiota: inclusion of nutrition and probiotic factors in the maintenance of eubiosis. Nutrients. 12(2): 419. doi: 10.3390/nu12020419
- [28] Ellepola AN, Khajah R, Jayatilake S, Samaranayake L, Sharma P, Khan Z. Impact of brief exposure to antifungal agents on the post-antifungal effect and hemolysin activity of oral *Candida albicans*. J Appl Oral Sci 2015;23(4):412-8.
- [29] Naglik JR, Gaffen SL, Hube B. Candidalysin: discovery and function in *Candida albicans* infections. Curr Opin Microbiol 2019;52:100-109.
- [30] Arya NR, Rafiq NB. Candidiasis. In: StatPearls [Internet]. Treasure Island (FL): StatPearls Publishing; 2020. https://pubmed.ncbi.nlm.nih.gov/32809459/[Last accessed on 2020 Sept 17].
- [31] Robbins N, Caplan T, Cowen LE. Molecular evolution of antifungal drug resistance. Ann Rev Microbiol 2017;71: 753–775.
- [32] Vazquez JA. Combination antifungal therapy: the new frontier. Future Microbiol 2007;2(2): 115-139.
- [33] Ayukekbong JA, Ntemgwa M, Atabe AN. The threat of antimicrobial resistance in developing countries: causes and control strategies. Antimicrob Resist Infect Contr 2017;6: 47.
- [34] Campitelli M, Zeineddine N, Samaha G, Masla S. Combination antifungal therapy: a review of current data. J Clin Med Res 2017;9(6): 451–456.
- [35] Haidar G, Singh N. How we approach combination antifungal therapy for invasive aspergillosis and mucormycosis in transplant recipients. Transplant 2018;102(11): 1815-1823.
- [36] Rodrigues ML, Albuquerque PC. Searching for a change: the need for increased support for public health and research on fungal diseases. PLoS Negl Trop Dis. 2018;12.
- [37] Alves R, Barata-Antunes C, Casal M, Brown AJP, Van Dijck P, Paiva S. Adapting to survive: how *Candida* overcomes host-imposed constraints during human colonization. PLoS Pathog 2020;16(5): e1008478.



- [38] Tagg JR, Dajani AS. Wannamaker LW. Bacteriocins of Gram-positive bacteria. Bacteriol Rev 1976; 40:722-56.
- [39] Allert S, Förster TM, Svensson CM, Richardson JP, Pawlik T, Hebecker B, *et al.* (2018). *Candida albicans*-induced epithelial damage mediates translocation through intestinal barriers. MBio 9:e00915-1.
- [40] Ghaddar N, Anastasiadis E, Halimeh R, Ghaddar A, Dhar R, AlFouzan W, Yusef H, El Chaar M. Prevalence and antifungal susceptibility of *Candida albicans* causing vaginal discharge among pregnant women in Lebanon. BMC Infect Dis 2020;20:32.
- [41] Rudramurthy SM, Singh S. *Candida* infections in immunocompetent hosts: pathogenesis and diagnosis. Curr Fungal Infect Rep 2020;14: 233–245.
- [42] Bao L, Zhang C, Dong J, Zhao L, Li, Y, Sun J. Oral microbiome and SARS-CoV-2: beware of lung co-infection. Front Microbiol 2020;11: 1840.
- [43] Galgóczy L, Papp T, Vágvölgyi, *In vitro* interaction between suramin and fluvastatin against clinically important Zygomycetes. *Mycos* 2009;52:5, 447-453.
- [44] Galgóczy L, Lukács G, Nyilasi I, Papp T, Vágvölgyi, CS. Antifungal activity of statins and their interaction with amphotericin B against clinically important Zygomycetes. *Act Biolog Hungar* 2010;61(3):356-365.
- [45] Galgóczy L, Bácsi A, Homa M, Virágh M, Papp T, Vágvölgyi C. *In vitro* antifungal activity of phenothiazines and their combination with amphotericin B against different *Candida* species. Mycoses. 2011;54(6):e737-743.
- [46] Chaux GE. Combination antifungal therapy for invasive mold infections involving polyenes. Infect Dis Clin Pract 2010;18(1):7-15.
- [47] Bhattacharya S, Sae-Tia S, Fries BC. Candidiasis and mechanisms of antifungal resistance. Antibiotics (Basel). 2020; 9(6): 312.
- [48] Kelesidisa T, Falagas ME. Substandard/counterfeit antimicrobial drugs. Clin Microbiol Rev 2015;28(2): 443–464.
- [49] Ogunshe AAO. Effect of production batches of antibiotics on *in vitro* selection criterion for potential probiotic candidates. J Med Food 2008;11 (4): 753-760.
- [50] Ogunshe AAO, Adepoju AA, Oladimeji ME. Clinical efficacy and health implications of inconsistency in different production batches of antimycotic drugs in a developing country. J Pharm Appl Biosci 2011b;3(1):158-164.
- [51] Ogunshe AAO. Comparative bacteriostatic potentials of oral paediatric antibiotic suspensions sold in two countries. Arch Clin Microbiol 2014;5 4(3):
- [52] Rauseo AM, Coler-Reilly A, Larson L, Spec A. Hope on the horizon: novel fungal treatments in development. Open Forum Infect Dis 2020;7:1–19.
- [53] Ogunshe AAO, Adinmonyema PO. Evaluation of bacteriostatic potency of expired oral paediatric antibiotic suspensions and implications on infant health. Pan Afr Med J 2014;19:378.
- [54] Song G, Liang G, Liu W. Fungal co-infections associated with global COVID-19 pandemic: a clinical and diagnostic perspective from China. Mycopathol 2020;185, 599–606.
- [55] Ozaras R, Arslan O, Cirpin R, Duman H. (2020). Coinfections among COVID-19 patients: a need for combination therapy? J Microbiol Immunol Infect. Available from: https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7431324/ [Last accessed on 2020 Sept. 26].
- [56] Houšť J, Spížek J, Havlíček V. Antifungal drugs. Metabol 2020;10(3): 106.



- [57] Jones CL. 2020. The potential environmental role of fungi as a complication in COVID-19 infections J Bacteriol Mycol Open Access 8(1):6–13.
- [58] Hokken MWJ, Zwaan BJ, Melchers WJG, Verweij PE. Facilitators of adaptation and antifungal resistance mechanisms in clinically relevant fungi. Fungal Genetics and Biology 2019;132: 103254.
- [59] Ogunshe AAO, Obiekea CA. Food risks of multi-drug-resistant indicator bacterialcontaminated most massively-consumed ethnic fermented food-seasoning condiments. J Nutr Ecol Food Res 2014;2: 1–11.
- [60] Ogunshe AAO. Assessing food safety implications of multi antibiotic resistant fermented-food-condiment-environment-adapted-bacteria. J Adv Microbiol 2019;5(2): 1-13.

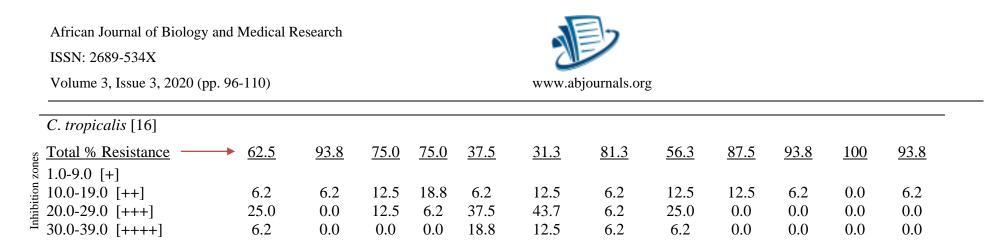
ISSN: 2689-534X



APPENDIX

Table 1: Antimycotic susceptibility / resistance patterns and profiles of oral *Candida* species to antifungal drugs

				An	tifungal d	rugs						
Candida species	IT	FV	GT	MC	VC-50	GF	LC	GS-500	DF	KZ	РХ	FD
C. albicans [42]												
Total % Resistance -	<u>→ 61.9</u>	<u>95.2</u>	<u>73.8</u>	<u>85.7</u>	<u>38.1</u>	<u>47.6</u>	<u>95.2</u>	<u>71.4</u>	<u>64.3</u>	<u>83.3</u>	<u>100</u>	<u>100</u>
₈ 1.0-9.0 [+]												
ត្ថ័ 10.0-19.0 [++]	7.1	0.0	9.5	9.5	14.3	11.9	4.8	16.7	16.7	7.1	0.0	0.0
© 20.0-29.0 [+++] 30.0-39.0 [++++] ≡ ≥40.0 [++++]	23.8	4.8	16.7	4.8	23.8	21.4	0.0	7.1	16.7	9.5	0.0	0.0
[™] 30.0-39.0 [++++]	7.1	0.0	0.0	0.0	21.4	16.7	0.0	2.4	2.4	0.0	0.0	0.0
≦ ≥40.0 [+++++]	0.0	0.0	0.0	0.0	2.4	2.4	0.0	0.0	0.0	0.0	0.0	0.0
C. glabrata [52]												
Total % Resistance -	→ 61 <u>.5</u>	<u>100</u>	<u>73.1</u>	80.8	40.4	44.2	<u>84.6</u>	<u>71.1</u>	<u>65. 3</u>	<u>98.1</u>	<u>92.3</u>	<u>94.2</u>
s 1.0-9.0 [+]												
s 1.0-9.0 [+] 10.0-19.0 [++]	7.7	0.0	19.2	19.2	17.3	15.3	9.6	11.5	19.2	0.0	3.8	5.8
.g 20.0-29.0 [+++]	23.1	0.0	7.7	0.0	26.9	32.7	1.9	15.4	13.4	1.9	0.0	0.0
5 20.0-29.0 [+++] 30.0-39.0 [++++] >40.0 [++++]	7.7	0.0	0.0	0.0	11.5	7.7	3.8	1.9	1.9	0.0	3.8	0.0
$\Xi \geq 40.0 \ [+++++]$	0.0	0.0	0.0	0.0	3.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0
C. pseudotropicalis [1	9]											
8 Total % Resistance -	<u>→ 36.7</u>	<u>100</u>	<u>63.1</u>	<u>78.9</u>	15.8	21.1	<u>73.6</u>	<u>63.1</u>	<u>68.4</u>	100	<u>100</u>	100
1.0-9.0 [+]												
	21.1	0.0	21.1	5.3	10.5	36.8	15.8	10.5	31.6	0.0	0.0	0.0
initial 10.0-19.0 [++] initial 20.0-29.0 [+++] initial 30.0-39.0 [++++]	21.1	0.0	15.8	10.5	31.6	26.3	5.3	21.1	0.0	0.0	0.0	0.0
∃ 30.0-39.0 [++++]	21.1	0.0	0.0	5.3	42.1	15.8	5.3	5.3	0.0	0.0	0.0	0.0



Keys: IT = *Itranox; GF* = *Gruffin; FV* = *Fesovin; LC* = *Lucon; GT* = *Gyno-Tiocosid; DF* = *Diflucan; MC* = *Medcan; KZ* = *Ketoconazole; VC-50* = *Vulcan-50; PX* = *Primpex; GS-50* = *Grisovid; FD* = *Flucamed*

Table 2: Multiple antimycotic resistance profiles of Candida species to antifungal drugs

Candida species	% multiple antimycotic resistance profiles
C. albicans [42]	41.7 [2 (4.8%)], 50.0 [3 (7.1%)], 58.3 [6 (14.3%)], 66.7 [6 (14.3%)], 75.0 [4 (9.5%)], 83.3 [8 (19.0%)], 91.7 [3 (7.1%)], 100 [10 (23.8%)]
C. glabrata [52]	41.7 [2 (3.8%)], 50.0 [4 (7.7%)], 58.3 [4 (7.7%)], 66.7 [11 (21.2%)], 75.0 [11 (21.2%)], 83.3 [7 (13.5%)], 91.7 [5 (9.6%)], 100 [8 (15.4%)]
C. pseudotropicalis [19	41.7 [1 (5.2%)], 50.0 [1 (5.2%)], 58.3 [7 (36.8%)], 66.7 [1 (5.2%)], 75.0 [4 (21.0%)], 83.3 [1 (5.2%)], 91.7 [2 (10.5%)], 100 [2 (10.5%)]
C. tropicalis [16]	50.0 [2 (12.5%)], 58.3 [1 (6.2%)], 66.7 [4 (25.0%)], 75.0 [3 (18.8%)], 83.3 [2 (12.5%)], 91.7 [3 (18.8%)], 100 [1 (6.2%)]



Table 3: Trends of antimycotic susceptibility / resistance patterns of Candida albicans to topical antifungal creams

						Antifungal creams							
Overall resistance	MY	SN	TY	LS	CC	FA	NZ	FG	KC	FC	WO	DR	%MAR
Total % resistance	100	100	100	100	100	100	100	100	100	100	100	100	100
Total % resistance	100	100	100	100	100	100	100	100	100	100	100	100	100
Total % resistance	100	100	100	100	100	100	100	100	100	100	100	100	100
Total % resistance	100	100	100	100	100	100	100	100	100	100	100	100	100

Keys:MY = Mycoten cream; NZ = Nizoral cream; SN = Skineal cream;FG =Fungusol cream; TY = Troyd cream; KC = Ketofung cream;LS = Lamisil cream; WO = Whitefield's ointment; CC = Care-cream;DR = Daktarin cream; FA =Funbact A cream; FG =

Table 4: Trends of antimycotic susceptibility / resistance patterns of Candida species to	D
combined antifungal drugs and topical creams	

Candida species	Single drugs	5 combined	Single	6 combined
	(5)	drugs	creams (12)	creams
C. albicans [9]	R [9 (100%)]	R [3 (33.3%)]	R [9 (100%)]	R [3 (33.3%)]
		<u>s</u> 20.0, 22.0,		<u>S</u> 19.0, 23.0,
		23.0, 25.0, 28.0,		25.0, 27.0, 28.0,
		31.0		30.0
C. glabrata [19]	R [19	R [6 (31.6%)]	R [19	R [4 (21.0%)]
	(100%)]	<u>S</u> 15.0, 16.0,	(100%)]	<u>S</u> 13.0, 16.0,
		17.0, 17.0, 17.0,		17.0, 17.0, 18.0,
		18.0, 18.0, 19.0,		20.0, 22.0, 23.0,
		20.0, 24.0, 27.0,		26.0, 26.0, 27.0,
		28.0, 28.0		27.0, 28.0, 29.0
C. pseudotropicalis [6]	R [6 (100%)]	R [3 (50.0%)]	R [6 (100%)]	R [2 (33.3%)]
		<u>s</u> 13.0, 23.0,		<u>S</u> 15.0, 19.0,
		28.0		20.0, 29.0
C. tropicalis [6]	R [6 (100%)]	R [3 (50.0%)]	R [6 (100%)]	R [2 (33.3%)]
_		<u>S</u> 28.0,		<u>s</u> 22.0, 25.0,
		30.0, 32.0		27.0, 33.0
Overall %	0.0%	62.5%	0.0%	72.5%
susceptibility				



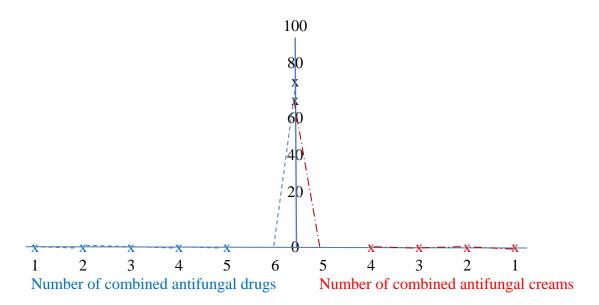


Figure 1: Susceptibility rates of combined antifungal drugs and creams

Copyright © 2020 The Author(s). This is an Open Access article distributed under the terms of Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International (CC BY-NC-ND 4.0), which permits anyone to share, use, reproduce and redistribute in any medium, provided the original author and source are credited.