

Brief Report

# The Mortality Attributable to Candidemia in *C. auris* Is Higher than That in Other *Candida* Species: Myth or Reality?

Carlos A. Alvarez-Moreno <sup>1,\*</sup>, Soraya Morales-López <sup>2</sup>, Gerson J. Rodriguez <sup>3</sup>, Jose Y. Rodriguez <sup>3</sup>, Estelle Robert <sup>4</sup>, Carine Picot <sup>4</sup>, Andrés Ceballos-Garzon <sup>4,5</sup>, Claudia M. Parra-Giraldo <sup>5</sup> and Patrice Le Pape <sup>4</sup>

<sup>1</sup> Facultad de Medicina, Universidad Nacional de Colombia, Clínica Universitaria Colombia, Clínica Colsanitas, Bogotá 111321, Colombia

<sup>2</sup> Grupo CINBIOS, Programa de Microbiología, Universidad Popular del Cesar, Valledupar 200004, Colombia

<sup>3</sup> Centro de Investigaciones Microbiológicas del Cesar (CIMCE), Valledupar 200002, Colombia

<sup>4</sup> Cibles et Médicaments des Infections et de l'Immunité, Nantes Université, CHU de Nantes, IICiMed, 10 UR1155, 44000 Nantes, France

<sup>5</sup> Unidad de Investigación en Proteómica y Micosis Humanas, Grupo de investigación en Enfermedades Infecciosas, Departamento de Microbiología, Facultad de Ciencias Pontificia Universidad Javeriana, Bogotá 110231, Colombia

\* Correspondence: caalvarezmo@unal.edu.co; Tel.: +57-31-4330-2367

**Abstract:** *Candida auris* has become a major health threat due to its transmissibility, multidrug resistance and severe outcomes. In a case-control design, 74 hospitalised patients with candidemia were enrolled. In total, 22 cases (29.7%) and 52 controls (*C. albicans*, 21.6%; *C. parapsilosis*, 21.6%; *C. tropicalis*, 21.6%; *C. glabrata*, 1.4%) were included and analysed in this study. Risk factors, clinical and microbiological characteristics and outcomes of patients with *C. auris* and non-*auris* *Candida* species (NACS) candidemia were compared. Previous fluconazole exposure was significantly higher in *C. auris* candidemia patients (OR 3.3; 1.15–9.5). Most *C. auris* isolates were resistant to fluconazole (86.3%) and amphotericin B (59%) whilst NACS isolates were generally susceptible. No isolates resistant to echinocandins were detected. The average time to start antifungal therapy was 3.6 days. Sixty-three (85.1%) patients received adequate antifungal therapy, without significant differences between the two groups. The crude mortality at 30 and 90 days of candidemia was up to 37.8% and 40.5%, respectively. However, there was no difference in mortality both at 30 and 90 days between the group with candidemia by *C. auris* (31.8%) and by NACS (42.3%) (OR 0.6; 95% IC 0.24–1.97) and 36.4% and 42.3% (0.77; 0.27–2.1), respectively. In this study, mortality due to candidemia between *C. auris* and NACS was similar. Appropriate antifungal therapy in both groups may have contributed to finding no differences in outcomes.

**Keywords:** *Candida auris*; mortality; candidemia; Colombia



**Citation:** Alvarez-Moreno, C.A.; Morales-López, S.; Rodriguez, G.J.; Rodriguez, J.Y.; Robert, E.; Picot, C.; Ceballos-Garzon, A.; Parra-Giraldo, C.M.; Le Pape, P. The Mortality Attributable to Candidemia in *C. auris* Is Higher than That in Other *Candida* Species: Myth or Reality? *J. Fungi* **2023**, *9*, 430. <https://doi.org/10.3390/jof9040430>

Academic Editors: Shankar Thangamani and Richard D. Cannon

Received: 13 February 2023

Revised: 12 March 2023

Accepted: 21 March 2023

Published: 31 March 2023



**Copyright:** © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (<https://creativecommons.org/licenses/by/4.0/>).

## 1. Introduction

Candidemia remains a worldwide public health concern. In Latin America, the incidence varies between 0.74–6.0 per 1000 hospital admissions and is associated with a high mortality rate (30–76%) [1]. Although there are geographical differences in the distribution of *Candida* species, *Candida albicans* remains the most frequently isolated. However, there is an increase in the incidence of invasive infections by non-*albicans* *Candida* species, including *Candida auris* [1,2]. *C. auris* is recognised as an emerging pathogen, which is difficult to identify, multiple-drug-resistant and highly transmissible. *C. auris* has a high potential for outbreaks in healthcare settings, possibly due to environmental contamination or transient colonisation by people or medical devices [3,4]. This species is known to form biofilms on inert surfaces, which can persist for extended periods and require stringent disinfection processes to remove [5,6]. To adapt to dry abiotic environments, *C. auris* activates stress-activated proteins [7] and produces hydrolytic enzymes that protect the yeast from

environmental stressors and disinfectants such as chlorhexidine and hydrogen peroxide. Chlorine-based products have been shown to be the most effective for environmental surface disinfection [8–10].

*C. auris* was described as a new species in 2009, when isolated from the external ear of a woman in Japan [11]. In 2011, it was described for the first time as a cause of fungemia in South Korea [12]. It has subsequently been isolated in at least 39 countries on 5 continents, causing isolated cases of colonisation to true outbreaks of invasive infections such as candidemia [13]. The first outbreak of *C. auris* in the Americas was reported in Venezuela (2012–2013) [14]. In Colombia, isolated cases have been reported since 2012 and, through a retrospective analysis, outbreaks since 2016 have indicated that *C. auris* could be considered to be endemic in several cities since 2013 [15–17]. In October 2022, the WHO issued its first fungal priority pathogen list, which included *C. auris* among others [18]. *C. auris* is commonly resistant to azole drugs and isolates that are resistant to all three main classes of antifungal agents have also been reported [19]. Mutations in the drug-target lanosterol 14- $\alpha$ -demethylase ERG11 gene such as Y132F, K143R and F126L frequently lead to azole resistance in *C. auris* strains. Additionally, mutations in the transcription factor TAC1 can cause an overexpression of CDR1, which also leads to high-level azole resistance in *C. auris* [20].

Mortality due to fungemia by *C. auris* is high. Recently, Chen et al. conducted a meta-analysis and found that the overall mortality of *C. auris* infections was 39% and 45% for bloodstream infections (BSI) [21]. However, the same authors conclude that the observed heterogeneities such as clade, BSI, drug resistance, continent and publication year were limitations. In general, mortality could be higher than that produced by other *Candida* species as a result of characteristics such as multiresistance, specific virulence factors, phenotypes, the ability to produce biofilms and a greater association with candidemia [13,17,21–23]. In addition, it has been difficult to confirm whether mortality from *C. auris* is related to the early selection of an adequate antifungal treatment due to its multidrug resistance. The objective of this study was to assess mortality due to *C. auris* fungemia and compare it with that resulting from NACS fungemia in the same period of time and in the same hospitals.

## 2. Materials and Methods

A case–case study was conducted in 12 hospitals in Valledupar, Colombia. We included all patients diagnosed with candidemia, defined by the presence of at least one positive blood culture for *Candida* spp. The participating institutions carried out laboratory surveillance for candidemia. All hospitals had automated blood culture systems. Blood cultures were collected under aseptic conditions and processed by conventional automated systems (BacT/ALERT 3D, bioMérieux, Marcy l’Etoile, France). The initial identification was performed with the method available at each institution of origin; i.e., Vitek 2 (bioMérieux, Marcy l’Etoile, France), Phoenix (Becton Dickinson, Franklin Lakes, NJ, USA), Austoscan-4 (Beckman Coulter, Fullerton, CA, USA) and Api Candida (BioMérieux, Marcy l’Etoile, France). The isolates were then sent to a regional reference laboratory where they were seeded in CHROMagar *Candida* medium (CHROMagar, Paris, France) and identified using MALDI-TOF mass spectrometry (Bruker Daltonics, Billerica, MA, USA). MALDI-TOF MS results were obtained according to the manufacturer’s specifications and all clinical isolates had a score above 2.0 [24]. *C. auris* identifications were confirmed by a single-tube PCR method based on the amplification of internal transcribed spacer (ITS) regions, as previously described by us [25]. Susceptibility to widely used antifungal drugs (AFG, anidulafungin; MCF, micafungin; CAS, caspofungin; 5-FC, flucytosine; PSC, posaconazole; VRC, voriconazole; ISV, isavuconazole; FLC, fluconazole; AMB, amphotericin B) was determined by Sensititre Yeast One<sup>®</sup> AST plates (ThermoFisher Scientific, Les Ulis, France). This method was used because it remains comparable with the CLSI reference method for testing the susceptibility of *Candida* spp. [26]. For the NACS isolates (*C. albicans*, *C. tropicalis*, *C. glabrata* and *C. parapsilosis*), CLSI M27-A3 breakpoints were used [27]. In the case of *C. auris*, breakpoints recommended by the US Centers for Disease

Control and Prevention (CDC) were applied. Resistance to FLC was set at  $\geq 32$   $\mu\text{g}/\text{mL}$ , AMB at  $\geq 2.0$   $\mu\text{g}/\text{mL}$  and CAS at  $\geq 2$   $\mu\text{g}/\text{mL}$  [28].

The clinical histories of patients with candidemia were reviewed; the information was collected in a case report and then tabulated into a database designed for this study. The demographic, clinical and microbiological variables as well as antifungal prophylaxis (FLC) and antifungal treatments were included. The risk factors for the development of fungemia that were evaluated were diabetes mellitus, haematological malignancy, solid organ malignancy, HIV infection, renal failure, need for haemodialysis, solid organ transplantation, extensive burns, need for mechanical ventilation, central venous catheter (CVC) or bladder catheter, red blood cell transfusion, history of abdominal surgery 30 days before the development of fungemia, parenteral nutrition, use of steroids defined as more than 20 mg of prednisone per day or an equivalent corticosteroid for more than 14 days before the onset of fungemia and previous use of antibiotics for more than 48 h in the 30 days before the onset of fungemia. The time to development of fungemia was defined as the number of days from hospitalisation to positive blood cultures for yeast. The time to start antifungal therapy was defined as the days from the first positive blood culture for yeast to the start of adequate antifungal therapy. An appropriate antifungal therapy was defined according to antimicrobial susceptibility tests. Additionally, information on outcomes such as mortality at 30 and 60 days from the onset of fungemia was included.

The data for categorical variables were expressed as a percentage and the continuous variables were expressed as the mean  $\pm$  SD or the median (interquartile range). A chi-squared test or Fisher's exact test (two-tailed) was used to compare the categorical variables and an unpaired Student's *t*-test was used to compare the continuous variables. A Kaplan–Meier analysis and log-rank test, recorded as a hazard ratio (HR: 95% CI), were used to compare both 30- and 90-day survival between the *C. auris* and NACS groups. Multivariate, backward, stepwise and logistic regression analyses were used to identify independent risk factors associated with day 30 mortality of patients; the results were presented as odds ratios (ORs) with 95% confidence intervals (95% CIs) and *p*-values  $< 0.05$  were considered to be significant. The statistical analysis was performed using EPI Info 7™ and GraphPad Prism 8.4.3. (Dotmatics, Boston, MA, USA) software.

### 3. Results

Seventy-four patients with candidemia were enrolled. The mean age of the patients was 38.2 years (range 5 days–88 years; SD 28.7); 44 patients were male. The average number of days from the start of hospitalisation until the onset of fungemia was 18.5 (range 0–55; SD 13.4). A total of 22 *C. auris* cases (29.7%) and 52 NACS cases (*C. albicans*, 25.6%; *C. parapsilosis*, 21.6%; *C. tropicalis*, 21.6%; *C. glabrata*, 1.4%) were included and analysed in this study.

The comparison of the main clinical and epidemiological variables is summarised in Table 1. Previous FLC exposure was significantly higher in *C. auris* candidemia patients (OR 3.3; 1.15–9.5). The in vitro antifungal susceptibility of the *Candida* species isolates is presented in Table 2. Most *C. auris* isolates were resistant to FLC (86.3%) and AMB (59%) whereas all were susceptible to echinocandins. Echinocandins (MCF and CAS) and 5-FC showed the lowest MICs among the antifungal drugs tested, with 100% of tested *C. auris* isolates having MICs of  $\leq 0.125$   $\mu\text{g}/\text{mL}$ . Regarding the NACS isolates, these were generally susceptible (FLC-resistant, 9.6%; AMB-resistant, 3.8%). No echinocandin-resistant isolates were detected.

**Table 1.** Epidemiological and demographic characteristics, underlying conditions, treatments and outcomes of patients with fungemia in 12 hospitals in Valledupar, Colombia.

	Total	%/(Range) SD	<i>C. auris</i>	%/(Range) SD	No. <i>auris</i>	%/(Range) SD	<i>p</i> -Value (OR)
<i>n</i>	74	100%	22	29.7	52	70.3	
Age (years), median	38.2	(5 days to 88 years old) SD 28.7	43.3	(8 months to 77 years old) SD 21.1	36	(5 days to 88 years old) SD 31.3	0.32 * (−7.3–21.8) *
Gender, male	44	59.5	12	54.5	32	61.5	0.3 (0.66–2.09)
Diabetes mellitus	9	12.2	4	18.2	5	9.6	0.6 (0.7–3.04)
Haematological malignancy	2	2.7	0	0.0	2	3.8	NA
Renal failure	21	28.4	9	40.9	12	23.1	0.1 (0.8–3.5)
Dialysis	11	14.9	6	27.3	5	9.6	0.06 (0.8–3.8)
Transplant	1	1.4	0	0.0	1	1.9	NA
Pancreatitis	3	4.1	2	9.1	1	1.9	0.08 (0.9–5.4)
Solid tumour	8	10.8	3	13.6	5	9.6	0.16 (0.8–44)
Extensive burns	1	1.4	0	0.0	1	1.9	NA
Mechanical ventilation	46	62.2	16	72.7	30	57.7	0.16 (0.76–3.1)
Blood transfusion	39	52.7	13	59.1	26	50.0	0.3 (0.68–2.8)
HIV	5	6.8	3	13.6	2	3.8	0.04 (0.99–5.32)
Central venous catheter	60	81.1	20	90.9	40	76.9	0.13 (0.74–3.3)
Urinary catheter	51	68.9	18	81.8	33	63.5	0.09 (0.81–3.4)
Abdominal surgery	24	32.4	7	31.8	17	32.7	0.09 (0.82–3.9)
Parenteral nutrition	29	39.2	6	27.3	23	44.2	0.48
Receipt of corticosteroids	8	10.8	2	9.1	6	11.5	0.13 (0.26–8.7)
Receipt of antibiotics	68	91.9	19	86.4	49	94.2	0.3 (0.07–2.9)
Previous use of fluconazole	23	31.1	11	50.0	19	36.5	0.02 (3.3–9.5)
Duration (days) of hospitalisation before candidemia, median (range)	18.4	(0–55) SD 13.4	21	(0–55) SD 17.4	17.3	(0–48) SD 11.3	0.29 (−3.1–10.3) *
Treatment							
No treatment	9	12.2	2	9.1	7	13.5	
Fluconazole	22	29.7	1	4.5	21	40.4	
Caspofungin	36	48.6	19	86.4	17	32.7	
Amphotericin B	4	5.4	0	0.0	4	7.7	
Combination of antifungals	3	4.1	0	0.0	3	5.8	

**Table 1.** *Cont.*

	Total	%(Range) SD	<i>C. auris</i>	%(Range) SD	No. <i>auris</i>	%(Range) SD	<i>p</i> -Value (OR)
Time to start antifungal (days)	3.6	(−1–34) SD 4.6	3.2	(1–6) SD 8.0	3.8	(−1–34) SD 5.4	0.68 (−2.9–1.98) *
Appropriate antifungal treatment	63	85.1	19	86.4	44	84.6	1.1 (0.27–4.8)
30 day mortality	28	37.8	7	31.8	21	40.4	0.66 (0.24–1.97)
90 day mortality	30	40.5	8	36.4	22	42.3	0.77 (0.27–2.1)

\* Two-tailed *p*-value and 95% confidence interval.

**Table 2.** Antifungal activities of the nine tested drugs against *C. auris* (*n* = 22) and NACS (*n* = 52).

Species	Drug	Number (and Cumulative Percentage) of <i>Candida</i> spp. Strains with MIC (µg/mL)															
		≤0.015	0.03	0.06	0.125	0.25	0.50	1	2	4	8	16	32	64	≥128		
<i>C. auris</i> <i>n</i> = 22	ANF	6 (27)	3 (41)	6 (68)	6 (95)	1 (100)											
	MCF	6 (27)	5 (50)	4 (68)	7 (100)												
	CAS	7 (32)	6 (59)	6 (86)	3 (100)												
	5-FC			19 (86)	3 (100)												
	PSC	3 (14)	4 (32)	1 (36)	5 (59)	8 (95)	1 (100)										
	VRC	1 (5)	2 (14)	3 (28)	2 (37)	4 (55)	7 (87)	3 (100)									
	ISV		1 (5)	4 (22)	2 (31)	8 (67)	7 (100)										
	FLC										2 (9)			1 (14)	5 (38)	5 (55)	9 (100)
	AMB						4 (18)	1 (100)	5 (41)	12 (96)	1 (100)						
<i>C. albicans</i> <i>n</i> = 19	ANF	16 (84)	2 (95)														
	MCF	18 (95)															
	CAS	5 (26)	11 (79)	2 (95)													
	5-FC	16 (84)			1 (89)												
	PSC	13 (68)	5 (95)														
	VRC	18 (95)															
	ISV	6 (32)	9 (79)	3 (95)				1 (100)									
	FLC				2 (11)	9 (58)	6 (89)	1 (100)						1 (100)			
	AMB						11 (58)	8 (100)									
<i>C. tropicalis</i> <i>n</i> = 16	ANF	4 (25)	2 (13)	2 (50)	7 (94)	1 (100)											
	MCF	6 (38)	10 (100)														
	CAS	1 (6)	13 (88)	2 (100)													
	5-FC	13 (81)			1 (87)										2 (100)		
	PSC				9 (56)	6 (94)	1 (100)										
	VRC		3 (19)		10 (81)	2 (75)	1 (100)										
	ISV				8 (50)	7 (94)	1 (100)										
	FLC						1 (6)	9 (63)	5 (94)								
	AMB						1 (6)	13 (88)	1 (94)	1 (100)				1 (100)			

Table 2. Cont.

Species	Drug	Number (and Cumulative Percentage) of <i>Candida</i> spp. Strains with MIC (µg/mL)														
		≤0.015	0.03	0.06	0.125	0.25	0.50	1	2	4	8	16	32	64	≥128	
<i>C. parapsilosis</i> n = 16	ANF			1 (6)	1 (13)		3 (31)	<b>8 (75)</b>	<u>3 (100)</u>							
	MCF			1 (6)		1 (13)	4 (38)	<b>7 (81)</b>	<u>3 (100)</u>							
	CAS			2 (13)	1 (19)	<b>8 (69)</b>	3 (88)	<u>2 (100)</u>								
	5-FC			<u>15 (94)</u>				1 (100)								
	PSC	4 (25)	<b>6 (63)</b>	4 (88)	<u>1 (94)</u>		1 (100)									
	VRC	7 (44)	<b>1 (50)</b>	4 (75)	<u>1 (81)</u>	1 (100)	<u>1 (94)</u>	1 (100)								
	ISV		6 (38)	7 (81)	<u>2 (94)</u>	1 (100)	<u>1 (100)</u>									
	FLC						4 (25)	3 (44)	<b>4 (69)</b>	2 (81)	<u>2 (94)</u>			1 (100)		
	AMB					2 (13)	<b>12 (88)</b>	<u>2 (100)</u>								
	5-FC			1 (100)												
<i>C. glabrata</i> n = 1	ANF	1 (100)														
	MCF	1 (100)														
	CAS			1 (100)												
	PSC							1 (100)								
	VRC						1 (100)									
	ISV						1 (100)									
	FLC												1 (100)			
AMB								1 (100)								

ANF: anidulafungin; MCF: micafungin; CAS: caspofungin; 5-FC: 5-fluorocytosine; PSC: posaconazole; VRC: voriconazole; ISV: isavuconazole; FLC: fluconazole; AMB: amphotericin B. MIC<sub>50s</sub> (minimal inhibitory concentration at which ≥50% of the strains are inhibited) and MIC<sub>90s</sub> (minimal inhibitory concentration at which ≥90% of the strains are inhibited) are depicted in bold letters and are underlined, respectively.

A total of 9 (12.2%) patients did not receive antifungal treatment, 2 in the *C. auris* group and 7 in the NACS group. Among the 65 (87.8%) patients who received antifungal treatments, 48.6% received CAS, 29.7% FLC, 5.4% AMB and 4.1% a combined antifungal therapy. The use of CAS as a fungemia treatment was higher in patients with *C. auris* (86.4% vs. 32.7%). The average time to start antifungal therapy was 3.6 days. A total of 63 (85.1%) patients received adequate antifungal therapy, without significant differences between the 2 groups (*C. auris*, 86.4% vs. NACS, 84.6%) ( $p$ -value 1.1; OR 0.27–4.8). The crude mortality at 30 and 90 days of candidemia was up to 37.8% and 40.5%, respectively. However, there was no difference in mortality at either 30 or 90 days between the group with candidemia caused by *C. auris* and the group with candidemia caused by NACS (31.8% vs. 40.4%; OR 0.6; 95% CI 0.24–1.97 and 36.4% vs. 42.3%; 0.77; 0.27–2.1, respectively).

#### 4. Discussion

An epidemiologic study conducted between 2008 and 2010 on candidemia in Latin America found that in Colombia, *C. parapsilosis* (38.5%) and *C. albicans* (36.7%) predominated, followed by *C. tropicalis* (17.4%), *C. glabrata* (4.6%) and *C. guilliermondii* (1.8%) [1]. However, in the last years, the situation has changed with the emergence of *C. auris* throughout the country [20,29]. Despite cases of infections by *C. auris* being described in several cities, most of the cases have been concentrated on the Atlantic coast [16,29].

This prospective study was performed during 2016–2017 at 12 health institutions in Valledupar, Cesar, a city of 450,000 inhabitants located in northern Colombia (approximately 180 km from the Atlantic coast). Seventy-four patients with candidemia were enrolled. *C. auris* was responsible for 29.7% of the fungemia, leading *C. auris* to be the most prevalent species in the area. The demographic and clinical characteristics of patients with *Candida* spp fungemia were similar to those reported in other studies [1,14]. There was no difference in risk factors for the development of fungemia in the group with *C. auris* vs. NACS, except for the previous use of FLC, which was higher in the *C. auris* group. This could be explained by the selection pressure exerted by this azole drug. In line with previous studies, the risk factors were not different from those associated with invasive infection due to NACS [30].

In the present study, 12.2% of patients did not receive antifungal treatment. The most likely reason was a late diagnosis as most of these patients died very early after the diagnosis of fungemia. CAS was the main antifungal given as the primary treatment. The *C. auris* group received CAS more frequently than the NACS fungemia group, probably as a consequence of the FLC prophylaxis that was more frequent in this group.

The literature has reported that mortality rates due to invasive *C. auris* infections range from 30% to 72% [31–34]. However, as in our study, in the multicentre retrospective case-control study of Simon et al., *C. auris* bloodstream infections were not associated with increased 30 day or 90 day mortality [35]. Indeed, in our study, mortality due to candidemia was similar between *C. auris* and NACS. An appropriate antifungal therapy in both groups may have contributed to finding no differences in outcomes. Similarly, comparative animal models, including murine and invertebrate models, indicated that *C. auris* was less virulent than *C. albicans*. Based on these models [36], it has been suggested that the decrease in virulence is likely related to the inability of *C. auris* to develop hyphae or pseudohyphae in mammals, which play a fundamental role in tissue invasion [37].

Although this study was conducted before the COVID-19 outbreak, we previously reported 20 cases of fungemia in hospitalised patients with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection across 4 institutions in the northern region of Colombia between June and September 2020 [38]. Of these, six patients had fungemia caused by *C. auris*. In this case series, the time to develop fungemia was similar to that reported in the current article (17.7 vs. 18.4 days). However, the mortality rate for this group of patients was higher (60% vs. 37.8%). Similarly, Villanueva-Lozano found a mortality rate of 83.3% (5/6) among patients with fungemia during an outbreak of *C. auris* infection in a COVID-19 hospital in Mexico [39] and Chowdhary found a mortality rate of 60% in patients with candidemia caused by *C. auris* who were critically ill with coronavirus disease

and admitted to an intensive care unit between April and July 2020 in New Delhi, India [40]. Therefore, it is expected that patients with fungemia caused by *C. auris* and SARS-CoV-2 infection may have a higher mortality rate.

Although *C. auris* has been described as a multidrug-resistant pathogen capable of generating resistance to the three most important classes of available antifungal drugs, there are geographical differences in the pattern of resistance [29]. A study in the USA found that more than 90% of *C. auris* isolates were resistant to FLC, more than 60% were resistant to AMB, 3.9% were resistant to echinocandins and 3 isolates were found to be pan-resistant [41]. In contrast, a report from India found that 40% of the isolates displayed a high MIC to CAS [42]. Our strains were characterised by a high percentage of resistance to azoles and AMB, placing echinocandins as the only available option for the treatment of these infections at the moment.

However, the emergence of echinocandin resistance in *C. auris* has become a major concern in several countries and the larger use of this drug to control nosocomial outbreaks could complicate patient management in the future [43]. This risk of multi-resistance explains why it is important to know the resistance levels of circulating clinical isolates and, therefore, to have validated susceptibility assays [44].

Although the mortality of *C. auris* candidemia was not higher than for other *Candida* species, this species can cause intra-hospital outbreaks. This is why limiting the transmission of this microorganism, adherence to infection control programs (contact isolation and hand washing) and adequate cleaning and environmental disinfection protocols should be emphasised.

**Author Contributions:** C.A.A.-M., P.L.P. and J.Y.R. participated in the conception and design of the study. J.Y.R. participated in the clinical management of the patients. S.M.-L. and C.M.P.-G. participated in the microorganism identification process. A.C.-G., E.R., C.P. and P.L.P. performed the molecular identification and susceptibility tests. C.A.A.-M. performed the statistical analyses. This manuscript was initially drafted by J.Y.R., G.J.R. and C.A.A.-M. and then revised by the other authors of this study. All authors have read and agreed to the published version of the manuscript.

**Funding:** This research received no external funding.

**Institutional Review Board Statement:** The research protocol was submitted and approved by the ethics committees of each participating institution (CE: CI-2021-07-04; CA: CI-2022-02-07; CV: CI-2022-06-05; CC: CI-2022-05-11; CELD and CSI: CI-2022-03-22; ICVC: CI-2022-03-29; CACC and CM: CI-2022-03-28).

**Informed Consent Statement:** As it was a retrospective study considered to be without risk, an exception was made for obtaining informed consent from all institutions.

**Data Availability Statement:** Not applicable.

**Conflicts of Interest:** PLP received grants from Astellas, Basilea, MSD and Pfizer and speaker's fees from Gilead, Basilea, Pfizer and MSD. All other authors declare no competing interest. All authors have submitted the ICMJE Form for Disclosure of Potential Conflicts of Interest.

## References

1. Nucci, M.; Queiroz-Telles, F.; Alvarado-Matute, T.; Tiraboschi, I.N.; Cortes, J.; Zurita, J.; Guzman-Blanco, M.; Santolaya, M.E.; Thompson, L.; Sifuentes-Osornio, J.; et al. Epidemiology of candidemia in Latin America: A laboratory-based survey. *PLoS ONE* **2013**, *8*, e59373. [CrossRef] [PubMed]
2. da Matta, D.A.; Souza, A.C.R.; Colombo, A.L. Revisiting Species Distribution and Antifungal Susceptibility of *Candida* Bloodstream Isolates from Latin American Medical Centers. *J. Fungi* **2017**, *3*, 24. [CrossRef] [PubMed]
3. Thatchanamoorthy, N.; Rukumani Devi, V.; Chandramathi, S.; Tay, S.T. *Candida auris*: A Mini Review on Epidemiology in Healthcare Facilities in Asia. *J. Fungi* **2022**, *8*, 1126. Available online: <https://www.mdpi.com/2309-608X/8/11/1126/htm> (accessed on 10 March 2023). [CrossRef] [PubMed]
4. Sabino, R.; Veríssimo, C.; Pereira, Á.A.; Antunes, F. *Candida auris*, An Agent of Hospital-Associated Outbreaks: Which Challenging Issues Do We Need to Have in Mind? *Microorganisms* **2020**, *8*, 181. Available online: <https://www.mdpi.com/2076-2607/8/2/181/htm> (accessed on 10 March 2023). [CrossRef] [PubMed]

5. Garcia-Bustos, V.; Cabanero-Navalon, M.D.; Ruiz-Saurí, A.; Ruiz-Gaitán, A.C.; Salavert, M.; Tormo, M.; Pemán, J. What Do We Know about *Candida auris*? State of the Art, Knowledge Gaps, and Future Directions. *Microorganisms* **2021**, *9*, 2177. Available online: <https://pubmed.ncbi.nlm.nih.gov/34683498/> (accessed on 13 June 2022). [CrossRef]
6. Jeffery-Smith, A.; Taori, S.K.; Schelenz, S.; Jeffery, K.; Johnson, E.M.; Borman, A.; *Candida auris* Incident Management Team; Manuel, R.; Brown, C.S. *Candida auris*: A Review of the Literature. *Clin. Microbiol. Rev.* **2018**, *31*, e00029-17. Available online: <https://pubmed.ncbi.nlm.nih.gov/29142078/> (accessed on 13 June 2022). [CrossRef]
7. Day, A.M.; McNiff, M.M.; da Silva Dantas, A.; Gow, N.A.R.; Quinn, J. Hog1 Regulates Stress Tolerance and Virulence in the Emerging Fungal Pathogen *Candida auris*. *mSphere* **2018**, *3*, e00506-18. [CrossRef]
8. Ku, T.S.N.; Walraven, C.J.; Lee, S.A. *Candida auris*: Disinfectants and implications for infection control. *Front. Microbiol.* **2018**, *9*, 726. [CrossRef]
9. Chakrabarti, A.; Sood, P. On the emergence, spread and resistance of *Candida auris*: Host, pathogen and environmental tipping points. *J. Med. Microbiol.* **2021**, *70*, 1318. [CrossRef]
10. Horton, M.V.; Johnson, C.J.; Kernien, J.F.; Patel, T.D.; Lam, B.C.; Cheong, J.Z.A.; Meudt, J.J.; Shanmuganayagam, D.; Kalan, L.R.; Nett, J.E. *Candida auris* Forms High-Burden Biofilms in Skin Niche Conditions and on Porcine Skin. *mSphere* **2020**, *5*, e00910-19. [CrossRef]
11. Satoh, K.; Makimura, K.; Hasumi, Y.; Nishiyama, Y.; Uchida, K.; Yamaguchi, H. *Candida auris* sp. nov., a novel ascomycetous yeast isolated from the external ear canal of an inpatient in a Japanese hospital. *Microbiol. Immunol.* **2009**, *53*, 41–44. [CrossRef]
12. Lee, W.G.; Shin, J.H.; Uh, Y.; Kang, M.G.; Kim, S.H.; Park, K.H.; Jang, H.C. First three reported cases of nosocomial fungemia caused by *Candida auris*. *J. Clin. Microbiol.* **2011**, *49*, 3139–3142. [CrossRef]
13. Du, H.; Bing, J.; Hu, T.; Ennis, C.L.; Nobile, C.J.; Huang, G. *Candida auris*: Epidemiology, biology, antifungal resistance, and virulence. *PLoS Pathog.* **2020**, *16*, e1008921. Available online: <https://pubmed.ncbi.nlm.nih.gov/33091071/> (accessed on 13 June 2022). [CrossRef]
14. Calvo, B.; Melo, A.S.A.; Perozo-Mena, A.; Hernandez, M.; Francisco, E.C.; Hagen, F.; Meis, J.F.; Colombo, A.L. First report of *Candida auris* in America: Clinical and microbiological aspects of 18 episodes of candidemia. *J. Infect.* **2016**, *73*, 369–374. [CrossRef]
15. Parra-Giraldo, C.M.; Valderrama, S.L.; Cortes-Fraile, G.; Garzon, J.R.; Ariza, B.E.; Morio, F.; Meis, J.F.; Lopes Colombo, A. First report of sporadic cases of *Candida auris* in Colombia. *Int. J. Infect. Dis.* **2018**, *69*, 63–67. [CrossRef]
16. Morales-Lopez, S.E.; Parra-Giraldo, C.M.; Ceballos-Garzon, A.; Martinez, H.P.; Rodriguez, G.J.; Alvarez-Moreno, C.A.; Le Pape, P. Invasive Infections with Multidrug-Resistant Yeast *Candida auris*, Colombia. *Emerg. Infect. Dis.* **2017**, *23*, 162–164. [CrossRef]
17. Escandón, P.; Cáceres, D.H.; Lizarazo, D.; Lockhart, S.R.; Lyman, M.; Duarte, C. Laboratory-Based Surveillance of *Candida auris* in Colombia, 2016–2020. *Mycoses* **2022**, *65*, 222–225. Available online: <https://pubmed.ncbi.nlm.nih.gov/34731508/> (accessed on 13 June 2022). [CrossRef]
18. Fisher, M.C.; Alastruey-Izquierdo, A.; Berman, J.; Bicanic, T.; Bignell, E.M.; Bowyer, P.; Brüggemann, R.; Garber, G.; Cornely, O.A.; Gurr, S.J.; et al. Tackling the emerging threat of antifungal resistance to human health. *Nat. Rev. Microbiol.* **2022**, *20*, 557–571. Available online: <https://pubmed.ncbi.nlm.nih.gov/35352028/> (accessed on 4 February 2023). [CrossRef]
19. Jacobs, S.E.; Jacobs, J.L.; Dennis, E.K.; Taimur, S.; Rana, M.; Patel, D.; Gitman, M. *Candida auris* Pan-Drug-Resistant to Four Classes of Antifungal Agents. *Antimicrob Agents Chemother* **2022**, *66*, e00053-22. [CrossRef]
20. Ceballos-Garzon, A.; Peñuela, A.; Valderrama-Beltrán, S.; Vargas-Casanova, Y.; Ariza, B.; Parra-Giraldo, C. Emergence, and circulation of azole-resistant *C. albicans*, *C. auris* and *C. parapsilosis* bloodstream isolates carrying Y132F, K143R or T220L Erg11p substitutions in Colombia. *Front. Cell. Infect. Microbiol.* **2023**, *13*, 309. [CrossRef]
21. Chen, J.; Tian, S.; Han, X.; Chu, Y.; Wang, Q.; Zhou, B.; Shang, H. Is the superbug fungus really so scary? A systematic review and meta-analysis of global epidemiology and mortality of *Candida auris*. *BMC Infect. Dis.* **2020**, *20*, 827. Available online: <https://pubmed.ncbi.nlm.nih.gov/33176724/> (accessed on 13 June 2022). [CrossRef] [PubMed]
22. Kean, R.; Brown, J.; Gulmez, D.; Ware, A.; Ramage, G. *Candida auris*: A Decade of Understanding of an Enigmatic Pathogenic Yeast. *J. Fungi* **2020**, *6*, 30. Available online: <https://pubmed.ncbi.nlm.nih.gov/32110970/> (accessed on 13 June 2022). [CrossRef] [PubMed]
23. Rudramurthy, S.M.; Chakrabarti, A.; Paul, R.A.; Sood, P.; Kaur, H.; Capoor, M.R.; Kindo, A.J.; Marak, R.S.K.; Arora, A.; Sardana, R.; et al. *Candida auris* candidaemia in Indian ICUs: Analysis of risk factors. *J. Antimicrob. Chemother.* **2017**, *72*, 1794–1801. Available online: <https://pubmed.ncbi.nlm.nih.gov/28333181/> (accessed on 13 June 2022). [CrossRef]
24. Ceballos-Garzon, A.; Amado, D.; Vélez, N.; Jiménez-A, M.J.; Rodríguez, C.; Parra-Giraldo, C.M. Development and Validation of an in-House Library of Colombian *Candida auris* Strains with MALDI-TOF MS to Improve Yeast Identification. *J. Fungi* **2020**, *6*, 72. Available online: <https://pubmed.ncbi.nlm.nih.gov/32471074/> (accessed on 10 March 2023).
25. Theill, L.; Dudiuk, C.; Morales-Lopez, S.; Berrio, I.; Rodríguez, J.Y.; Marin, A.; Gamarra, S.; Garcia-Effron, G. Single-tube classical PCR for *Candida auris* and *Candida haemulonii* identification. *Rev. Iberoam. Micol.* **2018**, *35*, 110–112. Available online: <https://pubmed.ncbi.nlm.nih.gov/29685376/> (accessed on 4 February 2023). [CrossRef] [PubMed]
26. Pfaller, M.A.; Chaturvedi, V.; Diekema, D.J.; Ghannoum, M.A.; Holliday, N.M.; Killian, S.B.; Knapp, C.; Messer, S.A.; Miskou, A.; Ramani, R. Comparison of the Sensititre YeastOne colorimetric antifungal panel with CLSI microdilution for antifungal susceptibility testing of the echinocandins against *Candida* spp., using new clinical breakpoints and epidemiological cutoff values. *Diagn. Microbiol. Infect. Dis.* **2012**, *73*, 365–368. [CrossRef]

27. CLSI Document M27-A3; Reference Method for Broth Dilution Antifungal Susceptibility Testing of Yeasts; Approved Standard. 3rd ed. Clinical and Laboratory Standards Institute: Wayne, PA, USA, 2008.
28. Centers for Disease Control and Prevention. *Candida auris*. In *Antifungal Susceptibility Testing*; CDC: Atlanta, GA, USA, 2018. Available online: <https://www.cdc.gov/fungal/candida-auris/c-auris-antifungal.html> (accessed on 1 December 2022).
29. Escandon, P.; Chow, N.A.; Caceres, D.H.; Gade, L.; Berkow, E.L.; Armstrong, P.; Rivera, S.; Misas, E.; Duarte, C.; Moulton-Meissner, H.; et al. Molecular Epidemiology of *Candida auris* in Colombia Reveals a Highly Related, Countrywide Colonization with Regional Patterns in Amphotericin B Resistance. *Clin. Infect. Dis.* **2019**, *68*, 15–21. [CrossRef]
30. Sarma, S.; Upadhyay, S. Current perspective on emergence, diagnosis and drug resistance in *Candida auris*. *Infect. Drug Resist.* **2017**, *10*, 155–165. Available online: <https://pubmed.ncbi.nlm.nih.gov/28652784/> (accessed on 4 February 2023). [CrossRef]
31. Al-Rashdi, A.; Al-Maani, A.; Al-Wahaibi, A.; Alqayoudhi, A.; Al-Jardani, A.; Al-Abri, S. Characteristics, Risk Factors, and Survival Analysis of *Candida auris* Cases: Results of One-Year National Surveillance Data from Oman. *J. Fungi* **2021**, *7*, 31. Available online: <https://pubmed.ncbi.nlm.nih.gov/33430221/> (accessed on 4 February 2023). [CrossRef]
32. Cortegiani, A.; Misseri, G.; Fasciana, T.; Giammanco, A.; Giarratano, A.; Chowdhary, A. Epidemiology, Clinical Characteristics, Resistance, and Treatment of Infections by *Candida auris*. *J. Intensive Care* **2018**, *6*, 69. Available online: <https://pubmed.ncbi.nlm.nih.gov/30397481/> (accessed on 4 February 2023). [CrossRef]
33. Lockhart, S.R.; Etienne, K.A.; Vallabhaneni, S.; Farooqi, J.; Chowdhary, A.; Govender, N.P.; Colombo, A.L.; Calvo, B.; Cuomo, C.A.; Desjardins, C.A.; et al. Simultaneous Emergence of Multidrug-Resistant *Candida auris* on 3 Continents Confirmed by Whole-Genome Sequencing and Epidemiological Analyses. *Clin. Infect. Dis.* **2017**, *64*, 134–140. Available online: <https://pubmed.ncbi.nlm.nih.gov/27988485/> (accessed on 4 February 2023). [CrossRef]
34. Osei Sekyere, J. *Candida auris*: A systematic review and meta-analysis of current updates on an emerging multidrug-resistant pathogen. *Microbiologyopen* **2018**, *7*, e00578. Available online: <https://pubmed.ncbi.nlm.nih.gov/29345117/> (accessed on 4 February 2023). [CrossRef]
35. Simon, S.P.; Li, R.; Silver, M.; Andrade, J.; Tharian, B.; Fu, L.; Villanueva, D.; Gonzalez Abascal, D.; Mayer, A.; Truong, T.; et al. Comparative Outcomes of *Candida auris* Bloodstream Infections: A Multicenter Retrospective Case-control Study. *Clin. Infect. Dis.* **2022**, *76*, e1436–e1443. Available online: <https://pubmed.ncbi.nlm.nih.gov/36062367/> (accessed on 4 February 2023). [CrossRef]
36. Forgács, L.; Borman, A.M.; Prépost, E.; Tóth, Z.; Kardos, G.; Kovács, R.; Szekely, A.; Nagy, F.; Kovacs, I.; Majoros, L. Comparison of in vivo pathogenicity of four *Candida auris* clades in a neutropenic bloodstream infection murine model. *Emerg. Microbes Infect.* **2020**, *9*, 1160–1169. Available online: <https://www.tandfonline.com/doi/abs/10.1080/22221751.2020.1771218> (accessed on 10 March 2023). [CrossRef]
37. Yue, H.; Bing, J.; Zheng, Q.; Zhang, Y.; Hu, T.; Du, H.; Wang, H.; Huang, G. Filamentation in *Candida auris*, an emerging fungal pathogen of humans: Passage through the mammalian body induces a heritable phenotypic switch. *Emerg. Microbes Infect.* **2018**, *7*, 1–13. Available online: <https://www.tandfonline.com/doi/abs/10.1038/s41426-018-0187-x> (accessed on 10 March 2023). [CrossRef]
38. Rodriguez, J.Y.; Le Pape, P.; Lopez, O.; Esquea, K.; Labiosa, A.L.; Alvarez-Moreno, C. *Candida auris*: A Latent Threat to Critically Ill Patients with Coronavirus Disease 2019. *Clin. Infect. Dis.* **2021**, *73*, e2836–e2837. Available online: <https://europepmc.org/articles/PMC7665436> (accessed on 10 March 2023). [CrossRef]
39. Villanueva-Lozano, H.; Treviño-Rangel, R.D.J.; González, G.M.; Ramírez-Elizondo, M.T.; Lara-Medrano, R.; Aleman-Bocanegra, M.C.; Guajardo-Lara, C.E.; Gaona-Chávez, N.; Castilleja-Leal, F.; Torre-Amione, G.; et al. Outbreak of *Candida auris* infection in a COVID-19 hospital in Mexico. *Clin. Microbiol. Infect.* **2021**, *27*, 813–816. [CrossRef]
40. Chowdhary, A.; Tarai, B.; Singh, A.; Sharma, A. Multidrug-Resistant *Candida auris* Infections in Critically Ill Coronavirus Disease Patients, India, April–July 2020. *Emerg. Infect. Dis.* **2020**, *26*, 2694–2696. Available online: <https://pubmed.ncbi.nlm.nih.gov/32852265/> (accessed on 10 March 2023). [CrossRef]
41. Ostrowsky, B.; Greenko, J.; Adams, E.; Quinn, M.; O'Brien, B.; Chaturvedi, V.; Berkow, E.; Vallabhaneni, S.; Forsberg, K.; Chaturvedi, S.; et al. *Candida auris* Isolates Resistant to Three Classes of Antifungal Medications—New York, 2019. *Morb. Mortal. Wkly. Rep.* **2020**, *69*, 6. [CrossRef] [PubMed]
42. Chowdhary, A.; Anil Kumar, V.; Sharma, C.; Prakash, A.; Agarwal, K.; Babu, R.; Dinesh, K.R.; Karim, S.; Singh, S.K.; Hagen, F.; et al. Multidrug-resistant endemic clonal strain of *Candida auris* in India. *Eur. J. Clin. Microbiol. Infect. Dis.* **2014**, *33*, 919–926. Available online: <https://link.springer.com/article/10.1007/s10096-013-2027-1> (accessed on 8 February 2023). [CrossRef]
43. Sharma, D.; Paul, R.; Chakrabarti, A.; Bhattacharya, S.; Soman, R.; Shankarnarayan, S.; Hagen, F.; Meis, J.F. Caspofungin resistance in *Candida auris* due to mutations in Fks1 with adjunctive role of chitin and key cell wall stress response pathway genes. *bioRxiv* **2020**. [CrossRef]
44. Ceballos-Garzon, A.; Garcia-Effron, G.; Cordoba, S.; Rodriguez, J.Y.; Alvarez-Moreno, C.; Le Pape, P.; Parra-Giraldo, C.M.; Morales-López, S. Head-to-head comparison of CLSI, EUCAST, Etest and VITEK<sup>®</sup>2 results for *Candida auris* susceptibility testing. *Int. J. Antimicrob. Agents* **2022**, *59*, 106558. Available online: <https://pubmed.ncbi.nlm.nih.gov/35227828/> (accessed on 4 February 2023). [CrossRef] [PubMed]

**Disclaimer/Publisher's Note:** The statements, opinions and data contained in all publications are solely those of the individual author(s) and contributor(s) and not of MDPI and/or the editor(s). MDPI and/or the editor(s) disclaim responsibility for any injury to people or property resulting from any ideas, methods, instructions or products referred to in the content.