

Prevalence of fungemia in a tertiary hospital: Analysis of the last decade

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SUMMARY

Introduction: The prevalence of nosocomial fungemia has increased worldwide, and mortality caused by this disease is high.

Objective: To assess progress in the last decade, and the prevalence and profile of fungal agents isolated in blood cultures performed in a tertiary university hospital.

Method: All the results of blood cultures processed at Hospital das Clínicas, Universidade Federal de Minas Gerais (HC-UFMG), in the time intervals 2001-2003 and 2011-2013 were analyzed retrospectively. For each three-year period, the number of collected blood cultures, the overall positivity rate and the percentage of fungemia were recorded. In addition, all identified fungal species were cataloged. All blood samples were incubated in the BacT/ALERT[®] (bioMérieux) automation system.

Results: In 2001-2003, 34,822 samples were evaluated, with 5,510 (15.8%) positive results. In 2011-2013, the number of blood cultures processed increased to 55,052 samples, with 4,873 (8.9%) positive results. There was an increase in the number of positive cultures for fungi in the analyzed period (2001-2003: 4.16%; 2011-2013: 5.95%; $p < 0.001$). Among the agents, candidemias were predominant, especially those caused by non-*albicans* *Candida* species (2001-2003: 57.64%; 2011-2013: 65.17%; $p < 0.05$). There was also an increase in fungemia caused by other genera (2001-2003: 2.62%; 2011-2013: 4.48%; $p < 0.01$).

Conclusion: There was an increase in the prevalence of fungemia in the last decade at HC-UFMG. Although candidemias have been responsible for most of the cases, there has been an increase in fungemias caused by other species.

Keywords: *Candida*, non-*albicans* *Candida*, fungemia, tertiary health care, prevalence.

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INTRODUCTION

The term fungemia indicates the presence of viable fungi in the bloodstream, confirmed by laboratory tests. Currently, over 80% of infections in the bloodstream, whether caused by fungi or bacteria, are acquired in hospitals or other medical care centers.¹

Mortality among patients with nosocomial fungemia is high, reaching rates as high as 50 to 80%, and has been attributed mainly to absence or inadequacy of initial antifungal therapy.^{2,3}

The prevalence of hospital fungemia has increased in recent decades worldwide.^{3,4} Several studies have shown that the main risk factors for fungemias are prolonged steroid therapy, chemotherapy, malnutrition, malignan-

cy, previous fungal colonization, dialysis, abdominal surgery and immunosuppression.^{5,6} Other determining factors for the increased occurrence of hospital fungemia are the growing use of broad-spectrum antibiotics and invasive technical procedures, such as central venous catheters, mechanical ventilation, parenteral nutrition, and the growing number of organ transplants.^{2,4,7}

The fungus most commonly isolated in the blood of patients with fungemia worldwide is *Candida albicans*.^{2,3,8,9} Pien et al. assessed blood cultures from 1,706 patients in US hospitals, and *C. albicans* was the main fungus found.¹ Costa et al., in a study conducted in São Paulo, Brazil, reported that *C. albicans* was responsible for 50% of the cases of nosocomial fungemia.⁹ According to Cisterna et al.,

in a multicentric study carried out in Spain, *C. albicans* caused 49.08% of the infections, followed by *C. parapsilosis* (20.73%), *C. glabrata* (13.61%) and *C. tropicalis* (10.77%).¹⁰

The incidence of non-*albicans* *Candida* species has shown significant increase in recent decades.³ As susceptibility to antifungal drugs varies among species of *Candida*, it is important to know the prevalence of each of them in hospitals.¹¹ Early distinction between candidemia and blood infection by other fungi is essential for effective therapy.⁴

In Brazil, epidemiological surveys of fungemias in communities and hospitals have become increasingly frequent. Motta et al., in a study conducted at the Faculdade de Medicina, Universidade de São Paulo (FMUSP), Hospital das Clínicas, reported a 4% prevalence of fungemia in blood cultures performed in 2006, with approximately 86% of cases related to species of the *Candida* genus. The species isolated most often were *C. albicans* (52.2%), *C. parapsilosis* (22.1%), *C. tropicalis* (14.7%), *C. glabrata* (6.6%).¹²

In addition to damage to the health of the population, fungemias also cause economic losses to public health systems. Typically, antifungal treatments are long, which increases the length of hospital stay and the cost of drugs.¹³⁻¹⁶ In a recent article, Bloos et al. demonstrated the higher financial cost in patients hospitalized due to candidemia compared with sepsis caused by other agents.¹⁵

Because of the inconvenience caused by this disease, each health care institution should know the profile of agents that cause fungemia in its population. This is vital not only for epidemiological purposes, but especially for care and therapy, serving also as parameter for implementation of preventive measures and hospital infection control.

Our study aims to compare prevalence progression and the profile of fungal species found in blood cultures of patients admitted to the Universidade Federal de Minas Gerais (UFMG), Hospital das Clínicas, Brazil, in the last decade.

METHOD

All the results of blood cultures processed in the sector of Microbiology, Laboratory Medicine Service of Hospital das Clínicas, Universidade Federal de Minas Gerais (HC-UFGM), in 2001-2003 and 2011-2013 were assessed. For each period, the number of samples collected, the overall positivity rate and the percentage of fungemia were recorded. In addition, all identified fungal species were cataloged.

In order to perform the blood cultures at the HC-UFGM, blood samples were taken after skin antisepsis with 0.5% alcoholic chlorhexidine. The number of samples varied according to the physician's discretion. Usually, three samples were collected from different anatom-

ical sites in the case of adult patients, and a single sample for children. For adults, 16 mL to 20 mL of blood were collected from each sample, distributed into two bottles: Green lid (aerobic) and orange lid (anaerobic). For the children, the volume of blood varied from 1 mL to 5 mL, collected in a single jar with a yellow lid.

The blood culture bottles were transported to the Microbiology sector and incubated in a BacT/ALERT® (bio-Mérieux) device. This is an automated system for incubation and identification of microbial growth, based on the colorimetric detection of CO₂ through sensors positioned on the bottom of culture flasks. Growth of microorganisms increases CO₂, showing positivity in the bottle.

The incubation time for antimicrobial growth was based on recommendations by the Clinical and Laboratory Standards Institute (CLSI). Between 2001 and 2003, blood cultures lacking growth of microorganisms within seven days of incubation were considered negative. In case of suspected fungemia reported in the medical test request form, the incubation period would be increased to 30 days.^{16,17} From 2011 to 2013, in turn, according to the recommendations in force, incubation periods were reduced to 5 and 14 days, respectively.^{16,17}

The positive samples were subjected to Gram staining and morphological analysis using optical microscopy. After the identification of yeast-form or mycelial filaments using Gram staining, the samples were seeded on glass slides containing a Sabouraud medium and incubated at 37°C to determine the species. For yeast-form growth, the following tests were performed to identify *Candida*: resistance to cycloheximide, germ tube formation, microculture and physiological fermentation. Whenever *Cryptococcus* was suspected, urea test and inoculation in Niger culture medium were performed. Mycelial fungi were subjected to microculture in Sabouraud's agar. Once the fungi were identified, the results were passed on to the Laboratory Informatics System and electronically released for consultation by the requesting physician and the patient.

For statistical analysis of the data, a two-sided chi-square test of the periods analyzed was adopted. P-value < 0.05 was considered significant.

RESULTS

From January 2001 to December 2003, and from January 2011 to December 2013, 89,874 blood samples were analyzed in the sector of Microbiology, Laboratory Medicine Service of HC-UFGM.

In 2001-2003, 34,822 blood samples were analyzed, of which 5,510 (15.8%) yielded positive results. Among

the latter, fungi were identified in 229 (16.4%) samples. In 2011-2013, 55,052 blood culture bottles were analyzed, with 4,873 (8.9%) positive samples, of which 290 (5.95%) contained fungi (Figure 1).

Among the isolated fungal species, prevalence of candidemias was observed: 97.38% (2001-2003) and 95.52% (2011-2013). The main species isolated in both periods was *Candida albicans*: 39.74% (2001-2003) and 30.34% (2011-2013). There was an increase in the number of positive cultures for fungi in the analyzed period (2001-2003: 4.16%; 2011-2013: 5.95%; $p < 0.001$). There was also an increase in the number of non-*albicans* *Candida* species (2001-2003: 57.64%; 2011-2013: 65.17%; $p < 0.05$) and in non-*Candida* genera (2001-2003: 2.62%; 2011-2013: 4.48%; $p < 0.01$). The fungi isolated in each analyzed period are shown in Table 1.

TABLE 1 Fungal species isolated from blood cultures performed at the Laboratory Medicine Service of Hospital das Clínicas, Universidade Federal de Minas Gerais, Brazil, in 2001-2003 and 2011-2013.

Species	2001-2003		2011-2013		Total	
	N	%	N	%	N	%
<i>Candida albicans</i>	91	39.7	88	30.3	179	34.5
<i>Candida glabrata</i>	1	0.4	7	2.4	8	1.5
<i>Candida guilliermondii</i>	3	1.3	4	1.4	7	1.3
<i>Candida kefyr</i>	0	0	2	0.7	2	0.4
<i>Candida krusei</i>	0	0	9	3.1	9	1.7
<i>Candida parapsilosis</i>	60	26.2	79	27.2	139	26.8
<i>Candida</i> spp.	29	12.7	24	8.3	53	10.2
<i>Candida tropicalis</i>	39	17	64	22.1	103	19.8
<i>Cryptococcus neoformans</i>	3	1.3	7	2.4	10	1.9
<i>Cryptococcus</i> spp.	3	1.3	4	1.4	7	1.3
<i>Fusarium</i> sp.	0	0	2	0.7	2	0.4
Total	229	100	290	100	519	100

N: Absolute number of cases; %: Percentage of cases.

DISCUSSION

Candida spp. is found in the gastrointestinal tract in 20 to 80% of the healthy adult population. These microorganisms can become pathogenic if there are changes in host defense mechanisms. Candidemia has a high incidence in tertiary hospitals, with overall mortality of 60%. Rapid detection of the etiological agent causing the infection is of fundamental importance to allow early adjustment of treatment and hence, the reduction of hospital mortality by fungemia.⁴

The HC-UFGM is a teaching hospital that offers public and general care, health assistance, research services

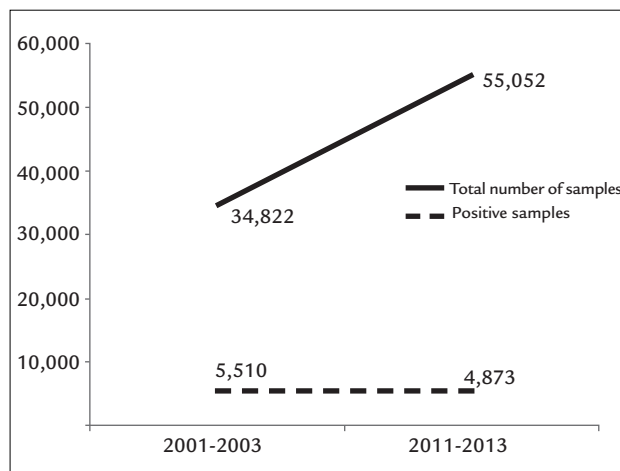


FIGURE 1 Number of total and positive samples of blood cultures, Laboratory Medicine Service of Hospital das Clínicas, Universidade Federal de Minas Gerais, Brazil, in 2001-2003 and 2011-2013.

and further training. It admits patients of the Unified Health System (SUS) only and is located in Belo Horizonte, Minas Gerais, Brazil. The HC-UFGM is a reference hospital in the municipal and state healthcare spheres for medium and high-complexity procedures, mainly in oncohematologic, infectious and parasitic, endocrine-metabolic, and mother and child diseases, as well as transplantation of organs and tissues. The HC-UFGM hospital complex has 511 beds; it includes one central building and seven other buildings for out- and in-patient care comprising all specialties and sub-specialties covered by the SUS system. Each month in these facilities, about 2,300 emergency room visits, 1000 hospitalizations, 24,000 outpatient visits and 155,000 laboratory tests take place.¹⁸

In this study, the authors observed an increased incidence of fungemia in the last decade. This increase was consistent with epidemiological data from other national and international hospitals.^{6,13,19-21} The increased prevalence of fungemia in hospital services may be related to advances in health care, which allowed greater survival of immunosuppressed patients.¹⁹ The frequent use of invasive instruments and materials in hospital care, and the indiscriminate use of broad-spectrum antimicrobials also contributed to the increase in fungemia.^{2,19}

The increase in the percentage of fungemia diagnosed in the last decade can be more significant, considering that false-negative results can occur if the minimum incubation time for growth of fungal species is not observed. At HC-UFGM, only samples reported as suspected fungemia were subjected to longer incubation times, either from 7 to 30 days (2001-2003) or from 5 to 14 days (2011-

2013).¹⁶ Unfortunately, the possibility of fungemia was not always informed on the request form for laboratory testing. In these cases, the samples were discarded early, probably hindering a proper diagnosis of fungemia.

We must also bear in mind that blood culture is a test that has intermediate sensitivity to detect fungal species, and may exhibit a sensitivity of only 50% for candidemias, for example.²² Therefore, the prevalence of fungemia detected in the HC-UFGM, and in other hospitals, can be falsely lower than the actual prevalence of this infection.

The most frequent microorganism in this study was *Candida* spp., which accounted for almost all of the isolated fungi, and *Candida albicans*, which was the most frequent species. These results are in agreement with other studies.^{2,3,8,22}

Even though *C. albicans* was the most prevalent species, our results revealed an increase in the prevalence of other fungal species, especially non-*albicans* *Candida* spp. Other authors have demonstrated high rates of candidemia caused by non-*albicans* species.^{3,23} The reasons for the emergence of other species have not been clarified. However, some risk factors are strongly associated to fungemia by certain species. *C. parapsilosis* has been linked to vascular catheter and parenteral nutrition.^{24,25} *C. tropicalis* was associated with cancer and neutropenia.^{26,27} *C. krusei* has been associated with prior use of azoles, neutropenia and hematologic malignancies.²⁸ *C. glabrata* has been associated with prior use of azoles, and transplants of solid organ and hematopoietic cells.^{2,29-31}

Some *Candida* spp. were isolated between 2011 and 2013: *Candida kefyr*, *Candida krusei* and *Fusarium* sp. Recent isolation of *C. glabrata* can represent a trend of increased incidence of this species in Brazil, already reported in São Paulo between 2006 and 2010,³² whose mortality seems higher than that of *C. albicans*.³

An interesting fact is the increased prevalence of genera other than *Candida*, whose mortality rate is higher than that found in candidemias.⁴ Previous colonization by these fungal species, collagen diseases and dialysis are risk factors associated with fungemia caused by non-*Candida* genera.⁴

Laboratory diagnosis of fungemia is extremely important to establish proper treatment and to reduce patient morbidity and mortality. Knowledge on the most prevalent species responsible for cases of fungemia within the service will allow the attending physician to initiate the most appropriate empiric therapy, as well as the analysis of a possible impact of new anti-fungal drugs introduced to the market on the profile of hospital fungemias.

CONCLUSION

There was an increase in the prevalence of fungemia among patients treated at the HC-UFGM in the last decade. Candidemias represented almost all cases, but there was an increase in fungemias caused by other genera.

RESUMO

Prevalência de fungemia em um hospital terciário: análise da última década

Introdução: a prevalência de fungemia hospitalar tem aumentado em todo o mundo e a mortalidade por essa afecção é elevada.

Objetivo: avaliar a evolução, na última década, da prevalência e do perfil dos agentes fúngicos isolados em hemoculturas realizadas em um hospital universitário terciário.

Método: foram analisados retrospectivamente todos os resultados de hemocultura processados no Hospital das Clínicas da Universidade Federal de Minas Gerais (HC-UFGM), entre os períodos de 2001-2003 e de 2011-2013. Para cada triênio foram registrados o número de hemoculturas coletadas, o percentual de positividade geral e o percentual de fungemia. Também foram catalogadas todas as espécies fúngicas identificadas. Todas as amostras sanguíneas foram incubadas no sistema de automação BacT/ALERT® (bioMérieux).

Resultados: entre 2001-2003, foram avaliadas 34.822 amostras, sendo 5.510 (15,8%) positivas. Entre 2011-2013, o número de hemoculturas processadas aumentou para 55.052 amostras, sendo 4.873 (8,9%) positivas. Observou-se um aumento do número de culturas positivas para fungos no período analisado (2001-2003: 4,16%; 2011-2013: 5,95%; $p < 0,001$). Dentre os agentes, as candidemias foram predominantes, principalmente por espécies de *Candida* não *albicans* (2001-2003: 57,64%; 2011-2013: 65,17%; $p < 0,05$). Houve também aumento da fungemia por outros gêneros (2001-2003: 2,62%; 2011-2013: 4,48%; $p < 0,01$).

Conclusão: houve aumento da prevalência de fungemia na última década no HC-UFGM. Embora as candidemias tenham sido responsáveis pela maioria dos casos, houve aumento de fungemias causadas por outras espécies.

Palavras-chave: *Candida*, *Candida* não *albicans*, fungemia, atenção terciária à saúde, prevalência.

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