

A 10-Year Survey of Antifungal Susceptibility of Candidemia Isolates from Intensive Care Unit Patients in Greece[▽]

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This study retrospectively reviews the susceptibility of 135 baseline ICU candidemia isolates (from 1997 to 2007) to nine antifungals as determined by the AFST-EUCAST microdilution method and identifies the most frequent causative agents of confirmed point-source candidemia outbreaks in local intensive care units. A minority of common and rare *Candida* species displayed decreased susceptibility to all antifungals.

Candidemia occurs in up to 10% of the patients in intensive care units (ICU), leading to prolonged use of mechanical ventilation and duration of hospital stay (5, 7, 8, 14, 16). As the choice of antifungals for treating candidemia in critically ill patients depends on knowledge of the local epidemiology, this study reviews the susceptibility of ICU candidemia strains isolated from 1997 to 2007 and archived in the UOA/HCPF1 collection (World Data Centre of Microorganisms, WDCM929 [http://wdcm.nig.ac.jp/hpcc.html]).

A total of 135 baseline *Candida* species from patients with ICU-acquired candidemia were tested. Isolates were identified to the species level by morphological and biochemical analyses and, where necessary, by sequencing of the internal transcribed spacer region and/or the D1/D2 domain, as described previously (4, 11, 12). MICs of pure compounds were determined with the AFST-EUCAST reference procedure (22). We tested standard compounds of fluconazole (Pfizer, Sandwich, Kent, UK), amphotericin B (Sigma, St. Louis, MO), itraconazole (Janssen, Beerse, Belgium), and flucytosine (Sigma, St. Louis, MO), voriconazole (Pfizer, Sandwich, Kent, United Kingdom), caspofungin (AS MedCare, Lexington, KY), and posaconazole (Schering Plough Research Institute, Kenilworth, NJ). Newer echinocandins, micafungin (Astellas Pharma Inc., Osaka, Japan) and anidulafungin (Pfizer, Groton, CT), were also tested. Quality control was performed for every testing occasion by using the designated quality control strains, *C. krusei* ATCC 6258 and *C. parapsilosis* ATCC 22019, which were within the MIC range given for the EUCAST-evaluated drugs (22). The quality control MIC range for anidulafungin was 0.25 to 0.5 µg/ml and 0.5 to 1 µg/ml for micafungin and caspofungin. For ATCC 22019, anidulafungin and micafungin MIC ranges were 0.5 to 2 µg/ml, and the MIC range for caspofungin was 2 to 4 µg/ml.

Inclusion of outbreak strains in this study may have constrained the accuracy of *Candida* species ranking order, but it

revealed the predominant causes of point source ICU outbreaks of candidemia during the study period. In that respect, *C. lusitanae* (1), *C. krusei*, and *C. parapsilosis* (data not shown) were causes of verified ICU outbreaks according to the related case definitions and descriptive and molecular epidemiology parameters. Therefore, a higher percentage (66.6%) of non-*C. albicans* isolates is reported than the 42% recorded for Europe in the SENTRY study (17). Despite *C. parapsilosis* being reported as the most frequent non-*C. albicans* ICU candidemia agent in Europe, geographical and possibly time-associated variations are highlighted in certain European studies in which *C. glabrata* is the second most frequently reported ICU candidemia agent (10, 13). Temporal variation was shown in a recent study in which, in 2002, *C. parapsilosis* reached 51% of the total isolation rates yet without reference to ICU outbreaks, while the absence of *C. tropicalis* was noted in the years 2000 to 2001 (3). Our study indicates that in the past 10 years, *C. glabrata* and *C. tropicalis* were always isolated at fluctuating isolation rates, whereas *C. dubliniensis*, known for its potential to acquire fluconazole resistance (2, 20), comprised 8.8% of the *C. albicans* isolates and was isolated only in the first 5 years of the study.

Our results show that drugs traditionally used in the ICU as standard treatments, such as fluconazole and amphotericin B, generally demonstrate low MICs, in agreement with previous observations (9, 19). However, high amphotericin B MICs were noted in two outbreak-unrelated *C. krusei* strains, the outbreak strain *C. lusitanae* (MIC, 4 µg/ml) (Table 1), and 3/10 *C. tropicalis* strains (MIC, 2 µg/ml). Interestingly, high flucytosine MICs (32 µg/ml) were recorded for 6/45 *C. albicans* isolates (MIC, ≥64 µg/ml), compared with 3/19 *C. krusei* isolates, 2/18 *C. lusitanae* isolates, 1/2 *C. rugosa* isolates, and a *C. intermedia* isolate. Besides *C. glabrata* and *C. krusei* resistance to fluconazole, only 4/18 outbreak-unrelated *C. parapsilosis* isolates demonstrated resistance (MIC, 32 µg/ml) (21). Nonetheless, fluconazole resistance associated with ICU fluconazole usage over the study period was not documented. Overall, posaconazole and voriconazole displayed potent antifungal activities against most ICU candidemia isolates. Yet, voriconazole resistance was identified in 8/19 *C. krusei* and 2/10 *C. tropicalis* isolates in accordance with their increased itracon-

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TABLE 1. Antifungal agent MIC ranges against 135 baseline ICU candidemia isolates and median MIC₉₀s of 121 *Candida* species isolated at high frequency

EUCAST microdilution MIC (μg/ml) for:																													
Drug ^a	<i>C. albicans</i> (n = 45)		<i>C. krusei</i> (n = 19) ^b		<i>C. parapsilosis</i> (sensu lato) (n = 18) ^c		<i>C. lusitanae</i> (n = 18) ^d		<i>C. glabrata</i> (sensu lato) (n = 11) ^e		<i>C. tropicalis</i> (n = 10)		<i>C. dublini- ensis</i> (n = 4)		<i>C. guilliermondii</i> (n = 3)		<i>C. rugosa</i> (n = 2)		<i>C. pelliculosa</i> (n = 1)		<i>C. kefyr</i> (n = 1)		<i>C. lipolytica</i> (n = 1)		<i>C. pulcherrima</i> (n = 1)		<i>C. intermedia</i> (n = 1)		
	Range		Range		Range		Range		Range		Range		Range		Range		Range		Range		Range		Range		Range		Range		
	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀	MIC ₉₀
ANID	0.03–0.5	0.12	0.03–0.5	0.06	0.5–4	2	0.03–0.5	1	0.03–0.12	0.12	0.03–8	0.25	0.03–1	0.5–2	0.12–2	1.0	16	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03
MICA	0.03–0.5	0.06	0.03–8	0.12	0.5–4	2	0.03–2	0.12	0.03–0.12	0.25	0.03–8	0.12	0.03–0.5	0.25–2	0.03–8	0.5	4	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03
CASP	0.12–1	0.12	0.5–8	0.12	0.5–8	2	0.25–8	1	0.25–4	1	0.12–4	0.25	0.25–2	1–16	0.03–8	0.5	8	0.12	0.12	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03
FC	0.12–32	2	0.12–64	2	0.12–0.5	0.5	0.12–64	2	0.12–0.5	0.25	0.12–0.5	0.5	0.12–1	0.12–0.25	0.03–0.25	0.03	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	64
POS	0.06–4	0.12	0.06–2	0.5	0.03–2	0.12	0.03–0.5	0.03	0.12–4	0.5	0.25–8	0.5	0.03–4	0.06–0.5	0.03–0.25	0.12	0.25	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.25
VO	0.01–2	0.03	0.12–16	1	0.06–2	0.06	0.12–4	0.5	0.12–4	0.5	0.06–16	0.12	0.03–4	0.06–0.5	0.01–0.5	0.12	4	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.03	0.06
IT	0.03–16	0.5	0.12–8	0.12	0.03–8	0.12	0.12–4	0.25	0.5–8	0.5	0.12–16	0.12	0.06–8	1–4	0.03–1	0.25	0.25	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.06
FL	0.12–32	0.5	32–64	32	1.0–32	4	0.12–4	4	8.0–64	16	0.5–4	2	0.12–0.25	0.25–4	2.0–16	2.0	2.0	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	1
AB	0.12–4	0.12	0.25–4	0.25	0.12–4	0.25	0.12–4	4	0.5–4	0.5	0.25–2	0.25	0.06–1	0.5–2	0.12–1	0.25	0.5	1	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.12	0.5

^a ANID, anidulafungin; MICA, micafungin; CASP, caspofungin; FC, fluconazole; POS, posaconazole; VO, voriconazole; IT, itraconazole; FL, fluconazole; AB, amphotericin B.

^b Includes tests for 1/10 outbreak strains.

^c Differential MICs for *C. orthopsilosis* and *C. metapsilosis* were incorporated; includes tests for 1/13 outbreak strains.

^d Includes tests for 1/8 outbreak strains.

^e Differential MICs for *C. brucensis* and *C. nivariensis* within the *C. glabrata* clade were incorporated.

azole MICs. Elevated posaconazole MICs were recorded for the same *C. albicans*, *C. krusei*, *C. glabrata*, and *C. tropicalis* strains, which displayed high itraconazole MICs and resistance to fluconazole and voriconazole, indicating azole cross-resistance.

Elevated echinocandin MICs were recorded for some *C. parapsilosis*, *C. guilliermondii*, and *C. rugosa* isolates, for a single *C. kefyr* isolate, and for some *C. tropicalis* isolates. The overall MIC₉₀ for anidulafungin and caspofungin against the six most frequently found *Candida* isolates was 0.5 µg/ml and for micafungin was 0.03 µg/ml, whereas 92% of the 121 isolates frequently found (Table 1) were inhibited at anidulafungin and caspofungin MICs of ≤2 µg/ml, as recorded previously (6, 19). Overall, the MIC₉₀ of echinocandins against *C. parapsilosis* (2 µg/ml) was conspicuously higher (18) than those recorded for the common *Candida* species, such as *C. albicans* (MIC range, 0.06 to 0.25 µg/ml), *C. glabrata* (MIC range, 0.12 to 1 µg/ml), and *C. tropicalis* (MIC range, 0.12 to 0.25 µg/ml).

The majority of *Candida* species demonstrated limited species-related and echinocandin-specific trailing and paradoxical (14) (Eagle) phenomena. With anidulafungin, *C. albicans* trailing was limited to 7/45 isolates. Paradoxical phenomena occurred in only 10/45 *C. albicans* isolates, 1/18 *C. lusitanae* isolates, and 1/4 *C. dubliniensis* isolates, while these phenomena occurred with micafungin (15) in only 4/45 *C. albicans* and only 1/4 *C. dubliniensis* isolates.

Data for the relative isolation rates and the recognition of outbreak-associated *Candida* species, including isolate susceptibility to nine established and recently licensed antifungal agents from Greek ICU, are reviewed for the first time. Indigenous ICU isolate susceptibility trends show that decreased susceptibility to all antifungals has a sporadic distribution among all isolates.

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