

Validation of 24-Hour Fluconazole MIC Readings versus the CLSI 48-Hour Broth Microdilution Reference Method: Results from a Global *Candida* Antifungal Surveillance Program[∇]

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We performed 24- and 48-h MIC determinations and disk diffusion testing of fluconazole against more than 11,000 clinical isolates of *Candida* species. By using the reference MIC breakpoints, the categorical agreement between the 24-h and reference 48-h broth microdilution results ranged from 93.8% (all *Candida* species) to 94.9% (all *Candida* species minus *Candida krusei*), with only 0.1% very major errors (VME). The essential agreement (within 2 log₂ dilutions) between the 24-h and 48-h results was 99.6%. The categorical agreement between the 24-h disk diffusion results and the 24-h MIC results, using the previously established breakpoints, was 94.4%, with 0.1% VME. Both the MIC and the disk diffusion results obtained for fluconazole after only 24 h of incubation may be used to determine the susceptibilities of *Candida* spp. to this widely used antifungal agent.

Recent studies examining the clinical utility of “real-time” antifungal susceptibility testing in the treatment of candidemia have shown that when such testing is available on site, physicians find the results helpful and not infrequently alter therapy on the basis of results (2, 15, 17, 18). Collins et al. (7) found that susceptibility testing of *Candida glabrata* isolates results in lower overall treatment costs, based on de-escalation of therapy from an expensive echinocandin to fluconazole, for patients with documented fungemia. Thus, it would appear that routine antifungal susceptibility testing can serve as an adjunct in the treatment of candidemia in the same way that antibacterial testing aids in the treatment of bacterial infections (13, 19, 40).

In light of the need to provide clinicians with useful information sooner rather than later (14, 22) and to avoid the potentially confounding effects of trailing growth on 48-h fluconazole MICs (1, 37, 39), the Clinical and Laboratory Standards Institute (CLSI; formerly NCCLS) Subcommittee for Antifungal Testing sought to determine if reading the broth microdilution (BMD) fluconazole MIC at 24 h would produce valid results when they were interpreted using the existing (48-h) breakpoints (25). Reanalysis of the MIC data set that was used to create the original CLSI M27 48-h fluconazole susceptibility breakpoints for *Candida* spp. (32, 38) showed good correlation between 24-h and 48-h MICs. Furthermore, when the 48-h fluconazole breakpoints were applied to MICs read at 24 h, the earlier reading predicted therapeutic outcomes as accurately as the 48-h MICs: 82% success for those episodes in which the 24-h MIC was ≤ 8 $\mu\text{g/ml}$ (susceptible [S] isolates), 55% success for those episodes in which the MIC was

16 to 32 $\mu\text{g/ml}$ (susceptible dose-dependent [SDD] isolates), and 39% success for those episodes in which the MIC was ≥ 64 $\mu\text{g/ml}$ (resistant [R] isolates) (25). Based on these results, the CLSI Subcommittee has included the option to read fluconazole MICs for *Candida* species after a 24-h incubation, using the original interpretive breakpoints, in CLSI documents M27-A3 and M27-S3 (5, 6).

Although earlier studies of 24-h fluconazole readings in different data sets and with a variety of methods versus the 48-h reference BMD method have shown similar results and relevance (8, 11, 12, 28, 31, 35, 41), further evaluation of this concept in other data sets is warranted (25). The purpose of the present study was to provide further documentation of the correlation between 24-h and 48-h fluconazole BMD MICs by assessing the essential agreement (EA; calculated as the percent agreement within ± 2 log₂ dilutions of the reference MIC) as well as the absolute categorical agreement (CA) and error rates obtained with a large data set of 24- and 48-h MIC results compiled in the course of global surveillance studies (27, 29, 30, 33, 34, 36). We also provide a reassessment of the fluconazole disk diffusion zone diameters as they relate to the 24-h fluconazole MIC results.

MATERIALS AND METHODS

Study design. A total of 11,654 clinical isolates of *Candida* spp. isolated from blood and other normally sterile body fluids via a global network of 105 sentinel hospital sites between January 2001 and December 2006 were included in the study. All isolates were saved on agar slants and were sent to the University of Iowa College of Medicine (Iowa City) for storage and further characterization by reference identification methods and susceptibility testing against fluconazole by BMD and disk diffusion methods (4–6, 16, 23).

Organism identification. All *Candida* species isolates were identified at the participating institutions by the routine method used in each laboratory. Upon receipt at the University of Iowa, the isolates were subcultured onto potato dextrose agar (Remel, Lenexa, KS) and CHROMagar *Candida* medium (Hardy Laboratories, Santa Maria, CA) to ensure viability and purity. Confirmation of species identification was performed with Vitek and API products (bioMérieux, St. Louis, MO) as recommended by the manufacturer or by conventional meth-

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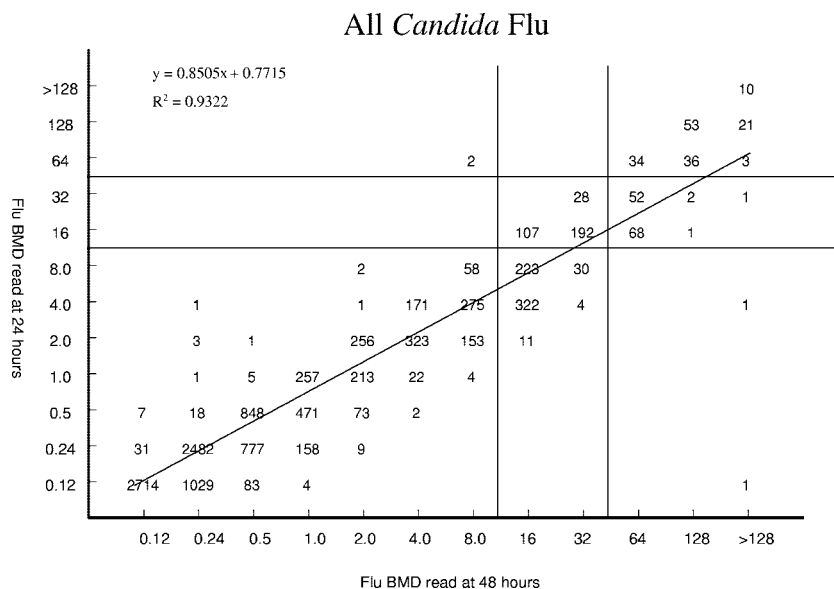


FIG. 1. Comparison of fluconazole (Flu) BMD MICs at 24 and 48 h for 11,654 *Candida* species isolates. The horizontal and vertical lines indicate the interpretive MIC breakpoints.

ods as required (16). Isolates were stored as water suspensions until they were used.

Susceptibility testing. Reference antifungal susceptibility testing of all 11,654 isolates was performed by BMD exactly as described in CLSI document M27-A3 (5). Fluconazole reference powder was obtained from Pfizer Pharmaceuticals (Groton, CT). Frozen BMD panels containing serial twofold dilutions of fluconazole (range, 0.12 to 128 $\mu\text{g/ml}$) in RPMI 1640 medium buffered to pH 7.0 with 0.165 M morpholinepropanesulfonic acid buffer were thawed and inoculated with an organism suspension adjusted to attain a final inoculum concentration of $1.5 \times 10^3 \pm 1.0 \times 10^3$ cells/ml. The panels were incubated in air at 35°C and observed for the presence or absence of growth at 24 and 48 h. The fluconazole MIC was read as the lowest concentration that produced a prominent decrease in turbidity (a ca. 50% reduction in growth) relative to that of the drug-free control (5).

Disk diffusion testing of fluconazole was performed on 11,237 of the isolates as described in NCCLS document M44-A (23). Fluconazole disks (25 μg) were obtained from Becton Dickinson (Sparks, MD). For disk diffusion testing, 150-mm-diameter plates containing Mueller-Hinton agar (Difco Laboratories) supplemented with 2% glucose and methylene blue (0.5 $\mu\text{g/ml}$) at a depth of 4.0 mm were used. The agar surface was inoculated by using a swab dipped in a cell suspension adjusted to the turbidity of a 0.5 McFarland standard. The plates were incubated in air at 35°C and read at 24 h. Zone diameter end points were read at 80% growth inhibition by using the BIOMIC image analysis plate reader system (version 5.9; Giles Scientific, Santa Barbara, CA). MIC interpretive criteria for fluconazole were those published by Pfaller et al. (32) and the CLSI (5, 6) and were as follows: S, MIC of ≤ 8 $\mu\text{g/ml}$; SDD, MIC of 16 to 32 $\mu\text{g/ml}$; R, MIC of ≥ 64 $\mu\text{g/ml}$. The interpretive criteria for the fluconazole disk test were those published by Pfaller et al. (32) and the NCCLS/CLSI (4, 23): S, zone diameter of ≥ 19 mm; SDD, zone diameter of 15 to 18 mm; R, zone diameter of ≤ 14 mm.

QC. Quality control (QC) was performed for BMD in accordance with CLSI documents M27-A3 and M27-S3 (6) by using *Candida krusei* ATCC 6258 and *Candida parapsilosis* ATCC 22019. QC determinations made on each day of testing were within the 24- and 48-h control limits described by the CLSI (6). QC for disk diffusion testing was performed by using *Candida albicans* ATCC 90028 and *C. parapsilosis* ATCC 22019 (4, 23).

Analysis of results. The MIC results obtained for fluconazole after 24 h of incubation were compared with those obtained after 48 h of incubation by using regression statistics and a scattergram (Fig. 1). Both on-scale and off-scale results were included in the analysis. As with previous studies (31, 35), high off-scale MIC results were converted to the next highest concentration, and low off-scale MIC results were left unchanged. Discrepancies among MIC end points (24-h versus 48-h results) of more than 2 dilutions (two wells) were used to calculate the EA. The CLSI interpretive breakpoints for fluconazole were used to obtain

CA percentages between the MICs determined after 24 h of incubation and the reference 48-h BMD results. Very major errors (VME) were identified when the reference MIC indicated R and the 24-h MIC indicated S. Major errors (ME) were identified when the isolate was classified as R at 24 h of incubation and as S at 48 h. Minor errors were identified when the result of one of the readings (at 24 h or at 48 h) was either S or R and that of the other was SDD.

In a similar fashion, the diameters of the zones of inhibition (in millimeters) surrounding the fluconazole disks at 24 h of incubation were plotted against their respective BMD MICs read at 24 h (Fig. 2). The method of least squares was used to calculate a regression line for each comparison. The interpretive breakpoints defined by the CLSI (4, 6) were used to determine the CA between the disk diffusion and 24-h BMD results for fluconazole. Error rates were calculated as described above using the BMD MIC as the reference test.

RESULTS AND DISCUSSION

Table 1 summarizes the in vitro susceptibilities of 11,654 isolates of *Candida* spp. (14 species) to fluconazole as determined by the CLSI BMD method and read at 24 and 48 h. The MIC results were typical of each species of *Candida* (28, 32), with the lowest MICs at both 24 and 48 h observed for *C. albicans* and the highest MICs observed for *C. glabrata* and *C. krusei*. In general, the MICs read at 24 h of incubation were twofold lower than those read at 48 h.

The overall EA between the 24-h and 48-h MIC readings was 99.6% (91.7% were within ± 1 dilution). Figure 1 illustrates the high degree of correlation between the two MIC readings ($R^2 = 0.9322$). Of the 44 discrepancies noted between the two readings, the MICs read after 24 h of incubation were lower than those obtained at 48 h in 38 instances (86.4%). Among the various species, the greatest numbers of discrepancies were observed with *C. albicans* (11 discrepancies), *C. glabrata* (21 discrepancies), and *Candida tropicalis* (6 discrepancies), species noted by others to exhibit the trailing phenomenon following incubation for 48 h (1, 24).

Regarding the individual species of *Candida*, the EA between the 24-h and 48-h BMD MICs was $>98\%$ for each of the 14 species included in the survey (Table 1). Given the CLSI

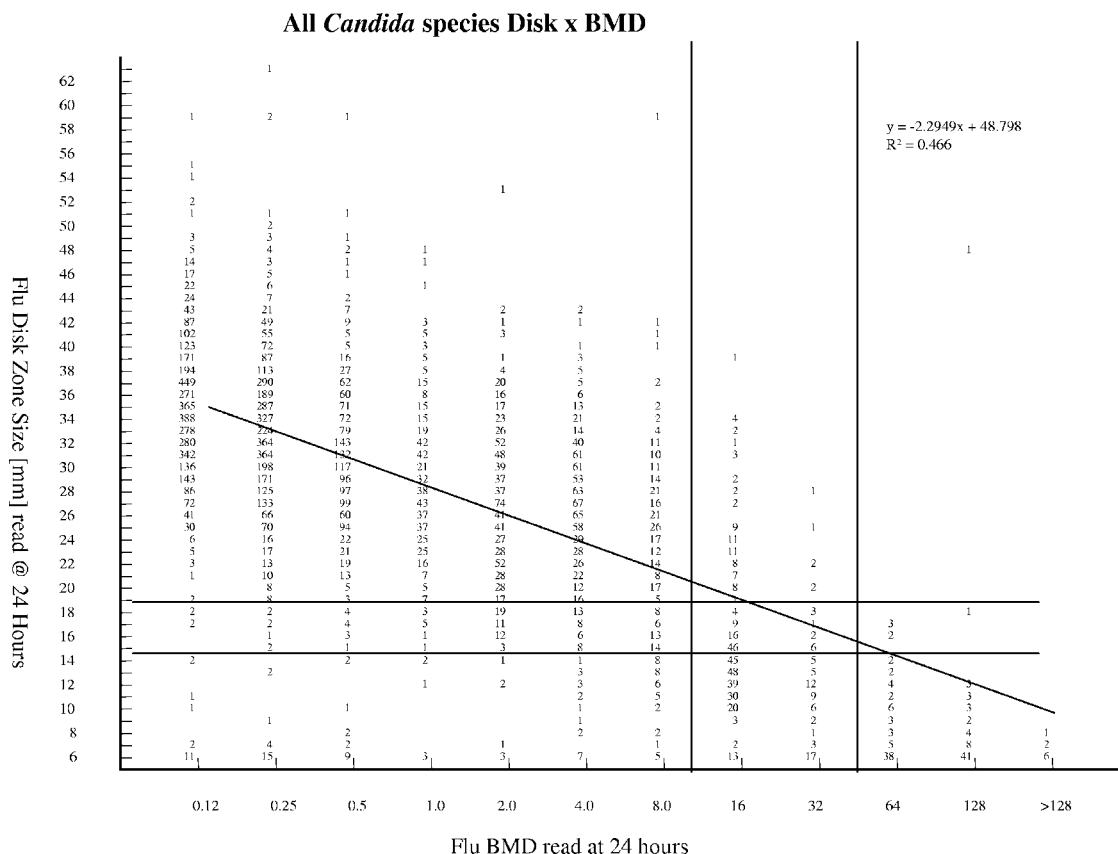


FIG. 2. Comparison of fluconazole (Flu) disk diffusion zone diameters at 24 h and BMD MICs at 24 h for 11,237 *Candida* species isolates. The horizontal and vertical lines indicate the interpretive zone diameter and MIC breakpoints, respectively.

recommendation that *C. krusei* should be considered to be intrinsically resistant to fluconazole and thus should not be tested against this agent (5), we also determined the EA for all isolates minus *C. krusei*; 99.6% of these results were within ± 2 dilution of one another.

The CA between the 24-h and 48-h fluconazole MICs is shown in Table 2. Excellent CA was observed for all comparisons with the exception of *C. glabrata* and *C. krusei*. The overall CA between the 24- and 48-h results was 93.8% when all isolates were included and 94.9% when the *C. krusei* results were omitted. Importantly, there were only two VME (false-susceptible results) and two ME (false-resistant results) in the entire 11,654-isolate comparison.

Although the absolute CA for *C. glabrata* was only 67.5%, virtually all of the errors (99.4%) were minor; they were predominantly the result of isolates determined to be S at 24 h and SDD at 48 h (96.2% of all minor errors). This is not surprising given the tendency of fluconazole MICs for *C. glabrata* to fall close to the susceptible breakpoint: 59% of MICs determined at 24 h and 78% of MICs determined at 48 h fell between 4 and 16 $\mu\text{g/ml}$ (data not shown). It should be noted, however, that 86% of the 156 *C. glabrata* isolates that were classified as R at the 48-h MIC determination were also R at the 24-h reading, and only 2 isolates were S at the 24-h reading and R at the 48-h reading (1.2% of all R *C. glabrata* isolates and 0.1% of all *C. glabrata* isolates tested). This accuracy in detecting fluconazole resistance among *C. glabrata* isolates is comparable or superior

to that observed with the FDA-approved commercial products Sensititre YeastOne (TREK) and Vitek 2 yeast antifungal test (bioMerieux) (31, 35). In view of the shift of results from SDD (at 48 h) to S (at 24 h) for this species, the CLSI Subcommittee has cautioned physicians and laboratorians to be aware that when an isolate is identified as *C. glabrata* and the 24-h or 48-h fluconazole MIC is $\leq 32 \mu\text{g/ml}$, patients should receive a maximum dosage of fluconazole (e.g., 12 mg/kg of body weight/day) (5, 6, 25, 26).

Disk diffusion testing of fluconazole has now been established as a simple and inexpensive qualitative method for determining the susceptibility of *Candida* to this agent, with results available within 24 h (30, 32, 34). Previously, the zone diameter breakpoints for fluconazole disk diffusion testing were derived by comparing the zone diameters read at 24 h with the BMD results at 48 h by using the error rate bounded method (21), whereby the number of discrepancies between the zone diameter and MIC categories was minimized (32). This process resulted in zone diameter breakpoints of ≥ 19 mm (S), 15 to 18 mm (SDD), and ≤ 14 mm (R), with an overall CA between the disk diffusion test results and the 48-h MIC test results of 92.8% (2,949 isolates) and very few VME (0.1%) or ME (0.4%) (32). Figure 2 shows the correlation between the fluconazole disk zone diameters read at 24 h and the BMD MIC results read at 24 h for 11,237 *Candida* isolates. By using the MIC and zone diameter breakpoints developed previously (32), the overall CA was 94.4%, with 0.1% VME and 1.1%

TABLE 1. Susceptibilities of 11,654 isolates of *Candida* spp. to fluconazole as determined by CLSI BMD methods and read after 24 and 48 h of incubation

Species	No. of isolates tested	Incubation time (h)	MIC ($\mu\text{g/ml}$) ^a			EA (%) ^b
			Range	50%	90%	
<i>C. albicans</i>	6,320	24	0.12->128	0.12	0.25	99.8
		48	0.12->128	0.25	0.5	
<i>C. parapsilosis</i>	1,664	24	0.12-64	0.5	2	99.8
		48	0.12->128	0.5	2	
<i>C. glabrata</i>	1,628	24	0.25->128	4	16	98.7
		48	0.25->128	8	32	
<i>C. tropicalis</i>	1,286	24	0.12-32	0.25	1	99.5
		48	0.12-64	0.5	2	
<i>C. krusei</i>	316	24	0.25-64	16	32	99.7
		48	0.25->128	32	64	
<i>C. guilliermondii</i>	142	24	0.5-32	2	4	100.0
		48	0.5-32	4	8	
<i>C. lusitanae</i>	139	24	0.12-64	0.5	1	99.3
		48	0.12-64	0.5	1	
<i>C. kefyr</i>	58	24	0.12-1	0.25	0.5	100.0
		48	0.12-2	0.25	1	
<i>C. pelliculosa</i>	34	24	0.5-8	2	4	100.0
		48	1-8	4	8	
Miscellaneous <i>Candida</i> spp. ^c	67	24	0.12-16	2	8	98.5
		48	0.12-64	2	8	
All <i>Candida</i> spp.	11,654	24	0.12->128	0.25	4	99.6
		48	0.12->128	0.25	16	
All <i>Candida</i> spp. minus <i>C. krusei</i>	11,338	24	0.12->128	0.25	4	99.6
		48	0.12->128	0.25	8	

^a 50% and 90%, MICs encompassing 50% and 90% of isolates tested, respectively.

^b Between 24- and 48-h BMD MICs.

^c Including *C. famata* (20 isolates), *C. rugosa* (14 isolates), *C. dubliniensis* (13 isolates), *C. lipolytica* (12 isolates), and *C. zeylanoides* (8 isolates).

ME. Thus, the disk diffusion test for fluconazole performs comparably to the 24-h MIC test without necessitating a change in interpretive criteria.

The findings of the present study confirm and extend the results of previous studies regarding the feasibility, accuracy, and clinical utility of 24-h fluconazole MIC readings (8, 11, 12, 28, 31, 35, 41). Indeed, if the 24-h MIC reading were considered to be a "new test," its performance relative to the 48-h reference BMD test would be considered superior to those reported for the fluconazole disk diffusion test, the Etest, the YeastOne colorimetric method, and the Vitek 2 yeast antifungal test (9, 10, 20, 35). An earlier multicenter study by Espinel-Ingroff et al. (11) not only documented excellent EA and CA for the comparison of 24- versus 48-h fluconazole MICs but also found a high degree (98%) of interlaboratory reproducibility among the six participating laboratories.

Clearly, the determination of fluconazole MICs after only 24 h of incubation would provide potentially important results in a more clinically useful time frame. Furthermore, previous investigations have shown that the 24-h fluconazole MIC end point correlated better than the 48-h end point with sterol quantification (1) and with treatment outcome both clinically (37) and in a murine model of invasive candidiasis (39). These findings suggest that fluconazole results for isolates of *Candida* spp. with significant trailing (e.g., *C. albicans*, *C. glabrata*, and *C. tropicalis* in the present study) should be interpreted on the basis of the lower MIC observed at the earlier (24-h) time point.

In addition to the data provided by Ostrosky-Zeichner et al.

(25), the clinical validity of 24-h fluconazole MICs was also addressed in a recent study by Baddley et al. (3), in which the authors demonstrated the association between patient characteristics, MICs for *Candida*, fluconazole pharmacodynamics, and mortality among hospitalized patients with candidemia. These investigators confirmed our findings that fluconazole MICs read after 24 and 48 h of incubation were very similar (Spearman's rank correlation coefficient, 0.91). Furthermore, classification and regression tree (CART) analysis was used to identify breakpoints for survival of 11.5 for a fluconazole AUC (area under the concentration-time curve)-to-MIC ratio and of 64 $\mu\text{g/ml}$ for MICs read after either 24 or 48 h of incubation. For 24-h MICs, 74% (57/77) of patients survived when the AUC/MIC ratio or MICs were above or below these thresholds, respectively (i.e., >11.5 or <64 $\mu\text{g/ml}$). Conversely, only 42.9% (3/7) of patients survived when either the AUC/MIC ratio was less than 11.5 or the MIC exceeded 64 $\mu\text{g/ml}$. Similar results were evident for 48-h MICs. Thus, regardless of the timing of MIC end point determination, infection with a fluconazole-resistant isolate was associated with increased mortality. Furthermore, these studies suggest that a clinician-controlled variable, fluconazole dose, may impact individual patient survival (3). In addition to host factors, the fluconazole dose and MICs may be helpful in managing and optimizing outcomes for patients with candidemia (3).

The simplicity and flexibility of disk diffusion testing makes it a very appealing method for use in the clinical laboratory. Although previous studies have already established the correlation between the 24-h fluconazole disk zone diameter and

TABLE 2. Categorical agreement between 24-h and 48-h CLSI BMD fluconazole MICs for 11,654 isolates of *Candida* spp.

Species (no. of isolates tested)	Incubation time (h)	% of isolates ^a that tested:			CA (%)	% of errors		
		S	SDD	R		VME	ME	Minor errors
<i>C. albicans</i> (6,320)	24	99.5	0.4	0.1	99.9	0.0	0.0	0.1
	48	99.4	0.5	0.1				
<i>C. parapsilosis</i> (1,664)	24	97.5	2.3	0.2	98.2	0.0	0.0	1.8
	48	96.1	3.3	0.6				
<i>C. glabrata</i> (1,628)	24	84.8	6.9	8.3	67.5	0.1	0.1	32.3
	48	53.7	36.7	9.6				
<i>C. tropicalis</i> (1,286)	24	99.6	0.4	0.0	99.5	0.0	0.0	0.5
	48	99.1	0.8	0.1				
<i>C. krusei</i> (316)	24	14.9	81.3	3.8	56.6	0.0	0.3	43.1
	48	1.6	65.5	32.9				
<i>C. guilliermondii</i> (142)	24	97.2	2.8	0.0	95.8	0.0	0.0	4.2
	48	93.0	7.0	0.0				
<i>C. lusitaniae</i> (139)	24	97.1	2.2	0.7	99.3	0.0	0.0	0.7
	48	97.1	1.4	1.5				
<i>C. kefyr</i> (58)	24	100.0	0.0	0.0	100.0	0.0	0.0	0.0
	48	100.0	0.0	0.0				
<i>C. pelliculosa</i> (34)	24	100.0	0.0	0.0	100.0	0.0	0.0	0.0
	48	100.0	0.0	0.0				
Miscellaneous <i>Candida</i> spp. ^b (67)	24	91.0	9.0	0.0	97.0	0.0	0.0	3.0
	48	89.6	9.0	1.4				
All <i>Candida</i> spp. (11,654)	24	94.8	3.9	1.3	93.8	0.02	0.02	6.16
	48	89.7	7.9	2.4				
All <i>Candida</i> spp. minus <i>C. krusei</i> (11,338)	24	97.0	1.7	1.3	94.9	0.02	0.01	5.07
	48	92.2	6.2	1.6				

^a Isolates were classified as S at a MIC of ≤ 8 $\mu\text{g/ml}$, as SDD at a MIC of 16 to 32 $\mu\text{g/ml}$, and as R at a MIC of ≥ 64 $\mu\text{g/ml}$.

^b Including *C. famata* (20 isolates), *C. rugosa* (14 isolates), *C. dubliniensis* (13 isolates), *C. lipolytica* (12 isolates), and *C. zeylanoides* (8 isolates).

48-h MICs (32), the data presented here demonstrate even better agreement between the zone diameters and 24-h fluconazole MICs and establish the continuing validity of the published zone interpretive criteria.

In summary, the MICs of fluconazole can be determined after 24 h of incubation for all species of *Candida* by using the CLSI BMD method. The high degree of accuracy of the 24-h reading compared to the 48-h reference method compared favorably to those reported previously for the FDA-approved methods Sensititre YeastOne and Vitek 2 yeast antifungal test. Both the 24-h MIC test and the 24-h disk diffusion test reliably identify fluconazole resistance among *Candida* spp. by using the previously established interpretive breakpoints. The availability of fluconazole susceptibility results within a 24-h time frame will be an important step in optimizing antifungal therapy for candidiasis.

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