Echinocandin Resistance

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1 Introduction

Fungal diseases cause life-threatening illnesses such as meningitis and pneumonias, chronic asthma, other respiratory diseases, and recurrent diseases like oral and vaginal thrush. Invasive fungal infections are a consequence of underlying health problems often associated with immunosuppression [1]. Fungal infections often carry high mortality and successful patient management requires antifungal therapy. Yet, treatment options remain extremely limited due to restricted classes of antifungal agents and by the emergence of prominent antifungal drug resistance. Currently registered antifungal drugs represented by polyenes and azoles, flucytosine, and echinocandins target the cell membrane, nucleic acid biosynthesis, and cell wall, respectively [2]. The latter and most recently approved class, the echinocandins, are now recommended as primary therapy for non-neutropenic patients with invasive candidiasis [3]. It is estimated that 60% of candidemia patients now receive an echinocandin for treatment or prophylaxis [4]. As worldwide use of echinocandins broadens, clinical failures due to resistant organisms are a concern, especially among certain Candida species. The development of echinocandin resistance among most susceptible organisms like Candida albicans is an uncommon event. Yet, there is a disturbing trend of increased resistance among strains of Candida glabrata, which are frequently cross-resistant to azole drugs. Echinocandin resistance is acquired during therapy and its mechanism is firmly established to involve amino acid changes in "hot-spot" regions of the Fks subunits of the target glucan synthase. These changes significantly decrease the sensitivity of the enzyme to drug resulting in higher MIC values and reduced

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Public Health Research Institute, New Jersey Medical School, Rutgers, The State University of New Jersey, 225 Warren Street, New Brunswick, NJ 07103, USA e-mail: perlinds@njms.rutgers.edu pharmacodynamic responses. Biological factors that promote selection of Fks-resistant strains involve complex cellular stress response pathways. The use of broth microdilution assays to assess susceptibility can be problematic with some drug- and species-related variability among clinical microbiology laboratories. Clinical factors promoting resistance include expanding use of echinocandins for therapy and prophylaxis, and localized reservoirs such as those in the gastrointestinal tract or intra-abdominal infections, which can seed emergence of resistant organisms. A basic understanding of the resistance mechanism, along with cellular and clinical factors promoting resistance, will promote better strategies to overcome and prevent echinocandin resistance.

2 Fungal Cell Walls and 1,3-β-D-Glucan

The cell wall of human fungal pathogens is essential for maintaining cell shape and rigidity. It consists primarily of an interwoven mesh of glucans, mannoproteins, and chitin. In yeasts like *Candida albicans*, branched fibrils of 1,3-β-D glucan form a network that acts as a scaffold for other macromolecules [5, 6]. Short 1-6-β-D-glucan chains establish bridges between linear 1,3-β-D glucan and cell wall proteins that coat the external surface of the cell wall. The majority of these proteins are heavily mannosylated through both O- and *N*-glycosidic linkages. Most cell wall proteins are covalently linked to the growing wall structure via 1-6-β-D-glucan. Chitin is found both below the network of $1,3-\beta$ -D glucan and as a linker between glucans. In other pathogenic fungi, including Aspergillus fumigatus and Cryptococcus neoformans, many of the same polysaccharides and mannoproteins are found in the cell wall, but the organization is somewhat different [7, 8] as polymers occur with other linkages between glucose units or sugars (e.g., galactomannan) [9]. When synthesis of a functional cell wall is reduced or eliminated, either by gene disruption or by inhibition with an antifungal inhibitor, cell growth is often adversely impacted leading to lysis and death.

3 Glucan Synthase

The fungal-specific enzyme $1,3-\beta$ -D glucan synthase (GS) is responsible for the biosynthesis of the central cell wall building block 1,3-B-D glucan. The enzyme is a membraneassociated complex that uses UDP-glucose to synthesize a 1,3-β-D glucan polysaccharide product 60 to 80 glucose residues in length. The enzyme has been extensively studied in S. *cerevisiae* [10], although it has also been studied in other yeasts and molds including Neurospora crassa, Aspergillus nidulans, and Aspergillus fumigatus; Schizosaccharomyces pombe; various Candida species; and Cryptococcus neoformans. GS is minimally a heterodimer involving a large integral membrane protein, encoded by FKS genes, that catalyzes the biosynthesis of 1,3-β-D-glucan and Rho, a regulatory GTP-binding protein. The FKS and RHO1 genes are conserved across numerous fungal genera. A high degree of homology among members of the FKS gene family aided cloning of paralogs from C. albicans [11, 12], C. neoformans [13], A. fumigatus [14], Neurospora crassa [15], P. carinii [16], and other fungi [10]. Conservation of FKS extends to the plant kingdom as well, where an FKS homolog is associated with synthesis of plant 1,3- β -D glucan (callose) in cotton and barley [17, 18]. Likewise, RHO1 genes have been identified and characterized in C. albicans [19], C. neoformans [20], and A. fumigatus [14]. Most yeast have three FKS genes, FKS1, FKS2, and FKS3. The FKS1 gene is essential in C. albicans [12, 13] and other Candida spp., while in C. glabrata, FKS1 and FKS2 are functionally redundant [21]. The FKS3 gene is expressed at a very low level relative to the other genes and its role is uncertain [22]. The GS enzyme complex has not been crystallized but it can be studied in an enriched form by a product entrapment technique [23, 24], which has allowed an evaluation of its kinetic properties [25].

4 Glucan Synthase Inhibitors and Echinocandins

There are three structural classes that define natural product inhibitors of 1,3- β -D glucan synthesis [10]. The first class are the lipopeptides including echinocandins, aerothricin lipopeptidolactones, and arborcandins. A second class comprises the glycolipid papulacandins, and a third class, the terpenoids, are represented by enfumafungin, ascosteroside, arundifungin, and ergokonin A. All GS inhibitor classes are noncompetitive with the biosynthetic substrate UDP-glucose. Cells exposed to GS inhibitors distort and lyse due to changes in cell wall glucans [26–28]. Of the three GS inhibitor classes, the echinocandins are best studied. The echinocandins are cyclic hexapeptides with an amide-linked fatty acyl side chain [29]. An early striking feature of this class was the potent activity of echinocandins in animal infection models due to *C. albicans*

[30] and *Pneumocystis jiroveci* [31]. This led to medicinal chemistry efforts at Merck, Eli Lilly, and Fujisawa (Astellas) and the development of current semisynthetic echinocandins caspofungin, anidulafungin, and micafungin, respectively [32]. The US Food and Drug Administration has approved echinocandin drugs for the treatment of esophageal and invasive candidiasis, including candidemia, empirical therapy in febrile neutropenic patients, and prophylaxis in patients undergoing hematopoietic stem cell transplantation (HSCT) [33, 34]. The first in-class drug caspofungin was also approved for salvage therapy for patients with invasive aspergillosis refractory to conventional therapy [35]. Echinocandin drugs show in vitro fungicidal activity against susceptible Candida spp. [36, 37], although they are fungistatic against molds where they alter morphology, cell wall composition, and organization [38, 39]. The echinocandins are largely inactive against invasive Zygomycetes, Cryptococcus spp., or Fusarium spp. As echinocandin drugs have a distinct mechanism of action specific for glucan synthase, they are highly effective against yeasts with reduced susceptibility to azoles, such as C. glabrata and C. krusei [40-42]; they are also active against some Candida biofilms [43-46]. The echinocandins have an excellent therapeutic index with a low potential for renal or hepatic toxicity or serious drug-drug interactions [47, 48].

5 Antifungal Spectrum and Breakpoints

The CLSI and EUCAST have established standardized microbroth dilution susceptibility tests for Candida and echinocandins, which show uniformly potent activity against most Candida species including C. albicans, C. glabrata, Candida tropicalis, and Candida krusei [49, 50]. The C. parapsilosis complex (Candida parapsilosis sensu stricto, C. orthopsilosis, and C. metapsilosis) and C. guilliermondii are notable exceptions displaying higher echinocandin antifungal MIC values relative to other highly susceptible Candida species [51-56]. Intrinsic reduced susceptibility has an unclear clinical significance, as patients infected with these strains are successfully treated with echinocandin drugs [57], although clinical response may vary with patient population [58–60]. The effect of echinocandins on filamentous fungi in vitro is less prominent with molds like A. fumigatus and other Aspergillus spp., showing reduced growth and altered hyphae morphology [39]. The multidrug-resistant pathogen Aspergillus lentulus is largely unresponsive to echinocandin action [61]. For A. fumigatus, the echinocandin-induced change in cell wall morphology correlates with exposure of masked epitopes (e.g., 1,3- β -D glucan), which promote a robust immune response contributing to in vivo efficacy [62]. Echinocandins show similar in vitro behavior with black molds such as Alternaria spp., and hyalohyphomycetes such as Scedosporium apiospermum [63]. In contrast, *Rhizopus oryzae* and other zygomycetes are

Antifungal agent	MIC breakpoint (mg/L)										
	Candida albicans		Candida	Candida glabrata		Candida krusei		Candida parapsilosis		Candida tropicalis	
	S	R	S	R	S	R	S	R	S	R	
Anidulafungin											
EUCAST	0.03	0.03	0.06	0.06	0.06	0.06	0.002	4	0.06	0.06	
CLSI	0.25	0.5	0.12	0.25	0.25	0.5	2	4	0.25	0.5	
Caspofungin											
EUCAST	ND ^b	ND	ND	ND	ND	ND	ND	ND	ND	ND	
CLSI	0.25	0.5	0.12	0.25	0.25	0.5	2	4	0.25	0.5	
Micafungin											
EUCAST	0.016	0.016	0.03	0.03	IEc	IE	0.002	2	IE	IE	
CLSI	0.25	0.5	0.06	0.125	0.25	0.5	2	4	0.25	0.5	

Table 29.1 EUCAST and CLSI antifungal breakpoints for major Candida species^a

^aAdapted from Arendrup et al. [72]

^bND: Not determined due to significant inter-laboratory variation in MIC ranges

°IE: Insufficient evidence (IE) due to small number of cases

largely unaffected by caspofungin [64]. Micafungin is active against mycelial forms of Histoplasma capsulatum, Blastomyces dermatitidis, and Coccidioides immitis but it is less active against yeast-like forms [65]. Like Aspergillus species, dermatophytes Trichophyton rubrum and Microsporum canis show diminished growth and malformed hyphae in response to echinocandins [66]. Finally, the neurotropic pathogen Cryptococcus neoformans is unresponsive to echinocandins [67, 68]. However, in vitro susceptibility can be overcome by addition of the calcineurin inhibitor FK506 [69]. Epidemiologic cutoff values (ECVs) have been determined for echinocandins against the most clinically important yeasts and molds from numerous global surveillance studies verifying the potent behavior of these drugs [70, 71]. The CLSI and EUCAST have also established species- and drug-specific clinical breakpoints (CBP) for echinocandin drugs based on extensive pharmacokinetic, microbiological, enzyme kinetic, and clinical response data [72, 73] (Table 29.1). See section on "Standardized Testing for Resistance."

6 Epidemiology of Echinocandin Resistance

Candida species isolates resistant to echinocandin drugs were first reported in 2005 [74]. Their frequency remains relatively low at less than 2–3% with *C. albicans* and most other *Candida* species [75–78]. Yet, consistent with the broader application of echinocandin therapy, high MIC clinical isolates associated with clinical failures are more commonly reported [22, 25, 79–89]. Despite these reports, echinocandin resistance among most *Candida* species has been largely unchanged in the past decade [90]. However, this is not the case for *C. glabrata*, where echinocandin resistance is rising and there is serious cause for concern since many isolates also display azole resistance [91–93], which

greatly limits therapy. The SENTRY Antimicrobial Surveillance Program reported echinocandin resistance of 8.0-9.3% among bloodstream isolates (BSI) of C. glabrata from 2006 to 2010 [92]. In a study of C. glabrata bloodstream isolates from Duke hospital spanning 10 years, echinocandin resistance of C. glabrata rose from 2 to 3% in 2001–2006 to more than 13 % in 2009–2010 [91]. Resistance is not uniform, as a study involving 1380 isolates of C. glabrata collected between 2008 and 2013 from four US cities showed that 3.1-3.6% of the isolates were resistant to the echinocandin drugs [93]. This is consistent with rates of 3.6 and 5.7% from anidulafungin and caspofungin, respectively, obtained from regional data of Candida non-albicans strains at US medical centers over a 6-year period (2006–2011) [90]. Yet, echinocandin resistance among C. glabrata has also coincided with a nearly parallel rise in azole resistance resulting in multidrug-resistant strains (Fig. 29.1). In a recent study covering 1032 isolates, nearly all isolates containing an FKS mutation were resistant to at least one echinocandin and 36 % were also resistant to fluconazole [93]. The expanding use of echinocandin and azole prophylaxis in many healthcare centers has prompted an epidemiologic shift with C. glabrata emerging as the most dominant fungal bloodstream pathogen [94, 95]. The development of echinocandin resistance typically occurs after prolonged therapy (3-4 weeks or longer) [87]. Yet, it has been observed to emerge shortly after the start of therapy [88, 96]. Echinocandin resistance in molds is rarely encountered but it has been reported for A. fumigatus [97] and the inherently multidrug-resistant A. lentulus [61].

7 Mechanism of Acquired Resistance

Echinocandin resistance resulting in clinical failures due to high MIC isolates involves modification of the catalytic subunit of glucan synthase, which is encoded by genes *FKS1* **Fig. 29.1** Rise in antifungal resistance of *Candida* glabrata to azole (fluconazole) and echinocandin (anidulafungin, caspofungin, and micafungin) drugs from 2001 to 2010. Adapted from Alexander et al. [91]



Table 29.2 Amino acid substitutions in hot-spot regions of Fks subunits of glucan synthase associated with reduced echinocandin susceptibility^{a,b}

			FKS1p)	FKS2p				
	Hot spot 1			Hot spot 2		Hot spot 1		Hot spot	
	AA	-							
	Pos								
C. albicans	641	FLTLSLRDP	1357	D <mark>W</mark> IR <mark>R</mark> YTL					
C. dubliniensis	641	FLTL <mark>S</mark> LRDP	1357	DWIRRYTL					
C. glabrata	625	F L I L S LRDP	1340	D <mark>WV</mark> RRYTL	659	FLILSLRDP	1374	DWIRRYTL	
C. kefyr	54*	LTL <mark>S</mark> LRDP	769*	DWVRRYTL					
C. krusei	655	<pre>FLILSIRDP</pre>	1364	DWIR <mark>R</mark> YTL					
C. lusitaniae	634*	FLTL <mark>S</mark> LRDP	* *	DWIRRYTL					
C. tropicalis	76*	<u>F</u> LT <u>LS</u> LRDP	792*	DWIRRYTL					
C. parapsilosis	652	FLTLSLRDA	1369	DWIRRYTL					
C. metapsilosis	104*	FLTLSLRD <u>A</u>	821*	DWIRRYTL					
C. orthopsilosis	39*	FLTLSLRD <u>A</u>	756*	DW <u>V</u> RRYTL					
C. guilliermondii	632	FMALSLRDP	1347	DWIRRYTL					
C. lipolytica	662	FL <mark>I</mark> LSLRDP	1387	DWIRR <mark>CV</mark> L					
S. cerevisiae	639	FLVLSLRDP	1353	DWVRRYTL	658	FLILSLRDP	1372	DWVRRYTL	

^aAdapted from Arendrup and Perlin [98]

^bRed: Strong resistance, difficult to treat; yellow: weak resistance, can be overcome with dosing; blue: natural polymorphism, elevated MIC but treatable; green: no effect on susceptibility

and/or *FKS2*. Echinocandin drugs are not substrates for multidrug transporters like azole drugs [42], and other cellular mechanisms conferring azole resistance do not affect echinocandin susceptibility. This has led to the recommendation of echinocandins as preferred therapy for infections involving azole-resistant strains of *Candida*. Echinocandin resistance is well characterized and known to be conferred by restricted mutations in two highly conserved "hot-spot" regions of the *FKS* genes [34] (Table 29.2). These *fks* mutations result in amino acid substitutions that induce elevated MIC values from 20- to 100-fold and reduced sensitivity of glucan synthase (IC₅₀) to drug by 50- to 3000-fold [22, 25, 99]. These less susceptible *fks* mutant strains respond poorly to echinocandin drugs in pharmacodynamic models of infection

[100-103], and the manifestation of characteristic *fks* mutations is associated with reduced clinical response [104–106]. The presence of an FKS mutation was found to be the only independent risk factor associated with echinocandin failure among C. glabrata isolates in a study of patients with invasive candidiasis [105]. The FKS resistance mechanism has been observed in many Candida species including C. albicans, C. glabrata, C. tropicalis, C. krusei, C. kefyr, and C. lusitaniae [96, 107, 108]. In all Candida species, except C. glabrata, mutations occur within two "hot-spot" regions of FKS1, encoding residues Phe641-Pro649 and Arg1361 (Table 29.2). In C. albicans, amino acid substitutions at Ser645 and Phe641 are the most abundant (Table 29.2). In C. glabrata, mutations occur in the homologous hot-spot regions of FKS1 and FKS2 [22, 99], although mutations are observed within FKS2 at twice the frequency of FKS1 [22, 34, 109]. Amino acid substitutions at Fks1 positions F625 and S629 and Fks2 positions F659 and S663 are most prominent inducing elevated MIC values (Table 29.2) [98]. In some cases, nonsense mutations and deletions are observed in FKS1 or FKS2 in C. glabrata [22, 98, 112]. Mutations in FKS1 or FKS2 can significantly alter the relative expression of their genes [21, 22], which can influence susceptibility. In C. glabrata, FKS2 expression is calcineurin dependent and downregulated by FK506 [111], and echinocandin resistance conferred by mutations in FKS2 are mitigated with FK506 [21]. A third hot-spot modification W695 (S. cerevisiae) was recently identified by in vitro selection [112], but it is not associated with clinical failures.

8 Biofilms

Biofilms also play a factor in resistance. They are one of the most important microbial communities encountered in nature, and they are well established to contribute to antifungal drug resistance [113]. It has been shown for echinocandin drugs that the extensive production of β -glucan within the extracellular glucan matrix helps sequester drugs by decreasing their concentration at the cell membrane surface [114]. Decreasing glucan productions, either by genetic or chemical means, increases the susceptibility to antifungal agents [115]. Genetic factors that regulate glucan formation promoting drug-sequestering biofilms include Rlm, Smi1, and glucan synthase Fks1 [115].

9 Acquired Resistance and Microbial Fitness

It is a well-established microbial paradigm that drug resistance often carries a fitness cost for microorganisms. The most prominent amino acid substitutions (e.g., Ser645 in *C. albicans*) in hot-spot regions of Fks subunits have been shown to decrease the catalytic efficiency for glucan biosynthesis [22, 25]. This reduced capacity for glucan production results in compensatory changes that alter cell wall morphology [116], which can reduce the fitness of such mutants. In *C. albicans*, reduced fitness has been observed for *fks* mutants in animal models [21, 22, 116]. The *fks* mutant strains compete weakly with their wild-type equivalents [116]. This reduced competition may account for the observation that resistance is with acquired during therapy and patient-patient transmission is not observed.

10 Cellular Stress and Drug Tolerance

The inhibition of glucan synthase following exposure of cells to an echinocandin drug induces significant cellular stress. In response, fungi activate a wide range of adaptive mechanisms that promote survival by helping protect against cell stress [117, 118]. These stress adaptation responses often result in drug-tolerant cells with elevated in vitro MIC values to echinocandins. Yet, they are not typically associated with clinical failures [119-121], as drug-exposed cells are less robust because glucan synthase is inhibited. Cell wall stress is sensed by receptors such Mtl2 and Wsc1, which induce stress tolerance involving cell wall integrity, protein kinace C (PKC), calcineurin-Crz1, and HOG [122, 123] interacting pathways. Hsp90 is an important protein that helps induce tolerance through its major client proteins calcineurin, along with its effector Crz1 [124-126]. Genetic or chemical impairment of Hsp90 function diminishes the ability of C. albicans and C. glabrata to develop tolerance in the presence of caspofungin [126, 127].

Chitin and glucans comprise the major structural components of the fungal cell wall and there is a prominent biosynthetic interdependence for both constituents [128]. Therefore, it is not surprising that echinocandin exposure results in compensatory increases in chitin synthesis to strengthen the cell wall and resistant drug action. Cell wall mutants with higher basal chitin contents are less susceptible to caspofungin [122, 123, 129, 130] and they confer reduced pharmacodynamics responses in animal model [131]. Paradoxical growth at very high drug levels has also been linked to prominent compensatory responses in chitin biosynthesis [132, 133]. Finally, defects in sphingolipid biosynthesis can differentially alter in drug-dependent fashion responses to echinocandin drugs. This mixed susceptibility phenotype is linked to interactions of the aliphatic tail of echinocandins and membrane sphingolipids [134, 135].

In general, tolerance pathways are insufficient to result in clinical drug failure. Yet, they are important for stabilizing cells in the presence of drug, and may account for stasis behavior of cells exposed to echinocandin drugs in animal model systems [102]. Even though these cells are not sufficiently resistant to induce therapeutic failures, they are poised to develop higher level resistance, as the drug-tolerant state allows cells sufficient time to overcome drug action by forming stable *FKS* mutations. It is not entirely clear how this ultimately occurs, although it may involve defects in DNA repair. Genome plasticity, observed widely in *C. albicans* and *C. glabrata* in response to azole drugs [136, 137], may also emerge as a factor for echinocandin drugs [138].

11 Mechanisms of Inherent Reduced Susceptibility

Candida parapsilosis complex (C. parapsilosis sensu stricto, Candida orthopsilosis, and Candida metapsilosis) and C. guilliermondii are intrinsically less susceptible in vitro to echinocandin drugs (MIC 0.5-8 µg/mL) relative to other highly susceptible Candida species [70, 95, 139], which prompted the CLSI to adopt higher breakpoints [73]. The clinical significance of this reduced susceptibility is unclear since patients can be successfully treated with echinocandins at standard dosages [54-56]; however, clinical efficacy may vary with patient population [58–60]. The underlying molecular mechanism appears to be naturally occurring polymorphisms in FKS hot-spot regions, which confer reduced sensitivity of glucan synthase to drug [140]. In C. parapsilosis complex, a highly conserved Pro660 is converted to alanine at the distal edge of hot-spot 1. Enzyme kinetic inhibition studies demonstrated that glucan synthase from the C. parapsilosis group were 10- to 50-fold less to echinocandin drugs than from enzymes obtained from highly susceptible species like C. albicans [140]. Furthermore, an engineered lab strain and clinical isolates of C. albicans and C. glabrata strains containing amino acid substitutions at this position display comparable decreases in target enzyme sensitivity and increased MIC values [140]. An additional I1359V polymorphism is observed in hot-spot 2 of C. orthopsilosis and S. cerevisiae, which confers higher MIC values. C. guilliermondii shows several additional amino acid polymorphisms in HS1 [140], although their relative contribution to overall insensitivity is unclear.

Cryptococcus neoformans is inherently resistant to echinocandin drugs even though 1,3 glucan synthase is essential and appears fully inhibited by echinocandin drugs in vitro [141]. It has been suggested that capsular melanin may help protect but capsule-deficient strains are also unresponsive to drug [142]. Finally, *Aspergillus lentulus*, a sibling species of *A. fumigatus*, is inherently resistant to a wide range of antifungal drugs including the echinocandins. The mechanism of this resistant is unclear but appears to be independent of *FKS* mutations [143].

12 Serum Effects on Drug Action

The echinocandin drugs are highly serum protein bound (>98%), which reduces their relative in vitro efficacy causing a shift in MIC [144–146]. The magnitude of the shift depends on the specific drugs with anidulafungin and micafungin showing a larger relative shift than caspofungin. A consequence of this shift in efficacy is that serum alters the relative fungicidal properties of the drugs, often resulting in fungistatic behavior against certain *Candida* species [147, 148]. The serum effects are more pronounced with mutant strains carrying *FKS* mutations [149].

13 Standardized Testing for Resistance

The Clinical and Laboratory Standards Institute (CLSI) and the European Committee on Antimicrobial Susceptibility Testing (EUCAST) have established comparable standards for broth microdilution (BMD) antifungal susceptibility testing of echinocandins against *Candida* species [53, 150, 151]. The objective for susceptibility testing is to establish an in vitro assessment that differentiates infecting strains as either susceptible or likely to respond to therapy or as resistant with an enhanced probability to fail therapy. In the case of echinocandin drugs, it is essential to capture high MIC strains containing FKS mutations. Initially, the CLSI used clinical and microbiological data to establish a preliminary common clinical breakpoint (CBP) for all three echinocandins against Candida spp. [120]. However, resistant strains with FKS mutations were often misclassified by this CBP [25, 152]. In response, the CLSI revised the CBP based on pharmacokinetic, microbiological, enzyme kinetic, and clinical data and established new species- and drug-specific breakpoints that better accounted for strains containing FKS mutations [73] (Table 29.3). However, the lower CBPs presented a clinical microbiology testing challenge, as BMD testing using either CLSI and EUCAST failed to promote consistent inter-laboratory test results without major errors (misclassifying wild-type strains as resistant or fkscontaining mutants as susceptible) between laboratory groups [153–154]. Disturbingly, there were wide modal ranges encountered with C. glabrata and caspofungin [153-155]. Consistent MIC results were obtained for micafungin and anidulafungin, and it was suggested that they could serve as testing surrogates for the class to assess resistance [98, 156, 157]. EUCAST has now established species-specific clinical breakpoints for micafungin against C. albicans, C. glabrata, and C. parapsilosis [72], and they have established breakpoints for anidulafungin to accommodate use of these compounds in some clinical situations [72, 158]. EUCAST has not set caspofungin breakpoints and does not currently

			MIC (µg/mL)	MIC (µg/mL)		ECV (µg/mL) ^a		
Antifungal agent tested	Species	No. of isolates	Range	Mode	≥95%	≥97.5%	≥99%	
Anidulafungin	C. albicans	8210	0.008-2	0.03	0.06	0.12	0.12	
	C. glabrata	2680	0.008–4	0.06	0.12	0.12	0.25	
	C. parapsilosis	3976	0.008-8	2	4	8	8	
	C. tropicalis	2042	0.008-2	0.03	0.12	0.12	0.12	
	C. krusei	322	0.008-2	0.06	0.12	0.25	0.25	
	C. lusitaniae	234	0.008-1	0.25	1	1	1	
	C. guilliermondii	222	0.03–4	1	4	8	8	
	C. dubliniensis	131	0.015–4	0.03	0.12	0.12	0.12	
Micafungin	C. albicans	7874	0.008–4	0.015	0.03	0.03	0.03	
	C. glabrata	3102	0.008–4	0.015	0.03	0.03	0.03	
	C. parapsilosis	3484	0.015–4	1	2	4	4	
	C. tropicalis	1605	0.008-8	0.015	0.06	0.06	0.12	
	C. kruse	617	0.015-1	0.06	0.25	0.25	0.25	
	C. lusitaniae	258	0.008-≥16	0.25	0.5	0.5	1	
	C. guilliermondii	234	0.015-8	0.5	2	2	4	
	C. dubliniensis	117	0.008-8	0.06	0.12	0.12	0.12	

Table 29.3 Anidulafungin and micafungin ECVs for eight species of Candida*

^aAdapted from Pfaller et al. [49]

Calculated ECVs comprising ≥95%, ≥97.5%, or ≥99% of the statistically modeled MIC population

recommend caspofungin MIC testing for clinical decision making involving echinocandin drugs [72]. Epidemiological cutoff values (ECV or ECOFF), which define the upper limit of the wild-type MIC population in the absence of a known resistance mechanism [49, 159], have been defined for anidulafungin and micafungin against common *Candida* species (Table 29.3). The ECV does not replace the BP, but it provides additional information for clinical decision making when a BP is not available. Although the designation of NWT does not allow a clinician to determine whether a particular isolate will respond to a particular antifungal agent, it does allow for a more informed decision based on how wildtype organisms would likely respond to therapy.

Rather than seeking testing surrogates or special conditions for BMD to distinguish wild-type strains from resistant isolates containing an FKS hot-spot mutation, it has been suggested that direct molecular testing for resistance mutations may provide a reliable alternative [160]. Direct DNA sequencing or real-time probing with allele-specific molecular probes provides an easy and unequivocal assessment of the resistance potential. The presence of an FKS mutation is the most important independent risk factor in predicting echinocandin therapeutic responses among patients with invasive candidiasis [104, 105, 110], which is well supported by extensive pharmacodynamics, MIC, and biochemical data [161, 162]. One criticism of this approach is that molecular testing requires specific knowledge of known resistance mechanisms and an unknown mechanism would not be detected. Yet, this probability is sufficiently remote given the large body of current data. Molecular testing to directly identify mutant strains containing FKS mutations would eliminate the current controversy surrounding some susceptibility testing, which prevents an accurate determination of resistance.

14 Paradoxical Growth Effects

The "paradoxical effect" refers to the unusual behavior of echinocandin drugs in susceptibility testing assays to show strong growth inhibition at low and moderate levels of drugs and then loss of inhibition at supra high drug concentrations, well in excess of the MIC. First described by Stevens and colleagues, it is a commonly observed property of echinocandin drugs [163]. This behavior is largely conditional as paradoxical strains show normal susceptibility properties following culture. The mechanism responsible for paradoxical growth is unclear, but is unrelated to mutations in FKS [124, 164]. It is not due to antifungal degradation or instability. The drug-induced growth behavior is more consistent with adaptive stress responses, which can lead to reduced susceptibility. In one instance, a paradoxical C. albicans strain showed a 900% increase in chitin content [133]. Consistent with changes in cell wall composition, remodeling is observed [165, 166]. The paradoxical effect is eliminated by serum, chitin synthase inhibitor nikkomycin Z, and calcineurin pathway inhibitors [167], and in C. albicans mutants that lack phosphatidylinositol-(4,5)-bisphosphate 5'-phosphatase [167]. Paradoxical behavior has been observed in a murine model of pulmonary aspergillosis [168] and in a patient with pulmonary aspergillosis [169]. Paradoxical growth in response to caspofungin in Candida species does not confer

survival advantage in a Drosophila or moth model of candidiasis [165, 170]. The clinical significance of the paradoxical growth remains unclear, as the drug levels necessary to induce it exceed normal human dosing levels.

15 Risk Factors for Resistance Emergence

The gastrointestinal (GI) tract is colonized with Candida species, often at very high burdens [171–178], which are in the form of a complex microbial biofilm [179]. Typically, drug penetration varies across the biofilm and drug concentrations in the glucan matrix are irregular [114]. This creates a drug exposure environment that can select for resistant variants, which may desorb from the biofilm and cause systemic infections. As biofilms are difficult to eradicate, they can form a resistance reservoir that seeds resistant infections. Similarly, intra-abdominal candidiasis occurs in 40% or more of patients following repeated gastrointestinal surgery, GI perforation, or necrotizing pancreatitis [180]. The high burden of Candida in this protected space with poor drug penetration creates a strong selection for resistant variants. Prophylaxis is another potential source for resistance. Prior and repeated exposure to echinocandin drugs is a risk factor development of resistance. As the FKS resistance mechanism is a prominent risk factor for therapeutic failure [105], resistance emergence is directly linked prior to exposure [106, 181, 182]. Antifungal prophylaxis with an azole or echinocandin class drug is standard prevention in many clinical settings with immunosuppressed patients at high risk for development of invasive fungal infections. Echinocandin drugs have been used because they have favorable pharmacokinetics and safety profile, and they are active against azole-resistant yeasts and molds. Both micafungin and caspofungin have been successfully applied for this purpose in adults [183-186] and children [187]. Meta-analyses have confirmed that echinocandin prophylaxis reduces the incidence of invasive fungal infections greater than fluconazole or itraconazole [188, 189]. Micafungin is FDA approved for prophylaxis of Candida infections in patients undergoing hematopoietic SCT or expected to be neutropenic for at least 10 days [190] and the European Society of Clinical Microbiology and Infectious Diseases guidelines also recommend micafungin for prophylaxis against Candida infections in allogeneic HSCT adult and pediatric patients, as well as in pediatric patients with acute myeloid and recurrent leukemia [191]. A consequence of the expanding use of echinocandins for prophylaxis is that patient drug exposure is on the rise, which has implication for inducing higher rates of echinocandin drug resistance, especially among resistance-prone organisms like C. glabrata. Even more concerning is the high prevalence of multidrug-resistant C. glabrata isolates crossresistant to both azole- and echinocandin-class drugs [91, 192-196]. The coevolution of azole and echinocandin multidrug resistance among *C. glabrata* is an alarming trend [91]. Breakthrough infections involving *C. albicans* are also reported in patients following transplantation who received micafungin prophylaxis [197]. It is not surprising that broadening patient exposure to echinocandin drugs would promote development of resistance. Echinocandin prophylaxis may continue to fuel an increase in the frequency of isolates that are resistant to multiple classes of antifungal drugs. Furthermore, prior antifungal exposure, especially with fluconazole, leads to genomic instability, which increases azole resistance [138] and may potentially predispose for enhanced mutations leading to *FKS*-mediated drug resistance.

16 Conclusions

Echinocandin resistance among Candida species is low but significant, especially among C. glabrata where high-frequency resistance is often associated with azole resistance resulting in multidrug-resistant strains. Characteristic mutations in hot-spot regions of FKS genes encoding glucan synthase remain the most significant factor responsible for resistant isolates that are refractory to therapy. However, in response to echinocandin action, cellular stress response pathways induce drug-adapted persister states, which can ultimately facilitate development of stable FKS-resistant genotypes. Host factors that promote resistance include biofilm formation within the gastrointestinal tract and intraabdominal candidiasis. The widespread use of echinocandin prophylaxis needs to be monitored for its effects on promoting enhanced drug exposure and resistance emergence. Effective antibiotic stewardship is required, especially in certain settings where resistance is prominent. Finally, new drug- and speciesspecific breakpoints have resulted in testing challenges, which may require drug surrogates for the class, but it may be more prudent to transition to sequence-based evaluation of FKS genotypes as the new gold standard for resistance assessment for all echinocandin drugs.

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