Antifungals: resistant mechanisms, drug susceptibility testing, and emerging drug resistant fungi

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Disclosures

- Received research funds from IMMY diagnostics, Vela diagnostics
- Advisor, CLSI (Clinical and Laboratory Standards Institute) Antifungal Susceptibility Subcommittee



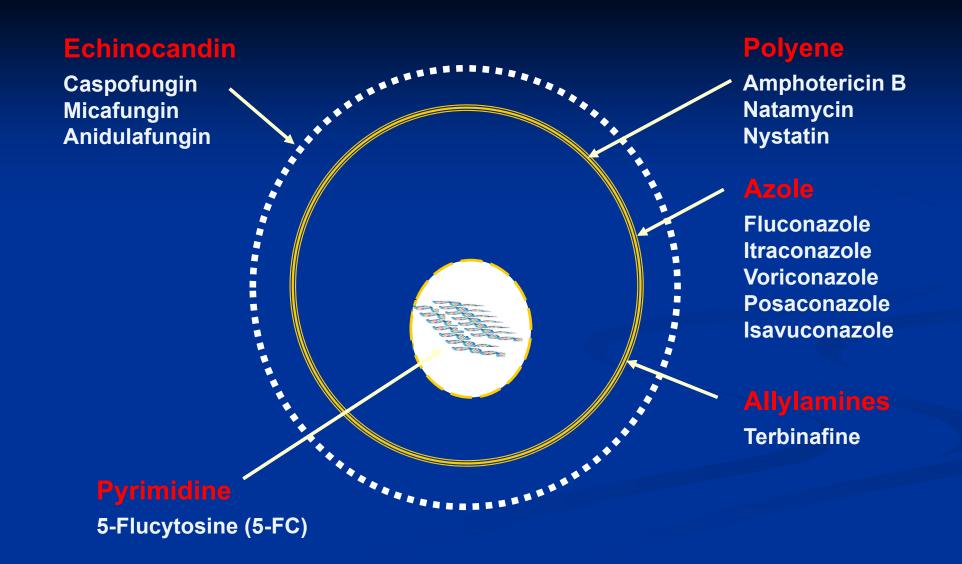
Objectives

- To review the common antifungal drug resistant mechanisms
- To describe antifungal drug susceptibility testing methods
- To recognize emerging antifungal drug resistant organisms



COMMON ANTIFUNGAL DRUG RESISTANT MECHANISMS





Antifungal drug resistance

- Primary resistance (intrinsic resistance; inherent resistance)
 - Found naturally without prior drug exposure.
 - Underlying resistant mechanism is inherent and not acquired during therapy.
- Acquired resistance
 - Drug selection pressure
 - Horizontal transmission between patients (rarely)

Arendrup MC J Infect Dis 2017; 216:S445



Amphotericin B

- Fungicidal
- Mechanisms of action: binding to ergosterol to destroy fungal cell membrane
- Resistance (No CLSI breakpoints; ECV of 2 µg/ml for *Candida* sp, and 2 or 4 µg/ml for *Aspergillus* sp)
 - High intrinsic reduced-susceptibility found in *C. krusei, C. auris, Trichosporon* sp, *A. terreus, L. prolificans, Fusarium* sp
 - Acquired resistance rare
- Mechanisms of resistance
 - Reduced content of ergosterol in the cell membrane
 - Biofilms



Azoles (Fluconazole \rightarrow Voriconazole, Itraconazole \rightarrow Posaconazole, Isavuconazole)

- Fungistatic (e.g. Fluconazole)
- Mechanisms of action
 - Inhibiting ergosterol biosynthesis by interfering with the action of lanosterol 14αdemethylase (encoded by *ERG11, Cyp51A*)
- Resistance
 - Fluconazole has no activity against Aspergillus sp, Fusarium sp, Mucorales
 - Intrinsic resistance to fluconazole: C. krusei, C. guilliermondii
 - Acquired resistance to fluconazole seen in C. albicans, C. tropicalis, C. glabrata



Mechanisms of azoles resistance

- Drug-target modification (genetic modification of the target *Erg11, Cyp51A*)
- Increase the target abundance (mutation in UPC2 leads overexpression of ERG11)
- Upregulation of efflux transporter genes (drug efflux pumps) resulting reduction of intracellular drug concentration
 - C. albicans (MDR1)
 - C. glabrata (CDR1, CDR2)
 - C. krusei (ABC1)
- Modification of other ergosterol biosynthesis pathway (ERG3)
- Biofilms



Echinocandins (Caspofungin, Micafungin, Anidulafungin)

- Fungicidal for Candida sp, but fungistatic for Aspergillus sp.
- Mechanisms of action
 - Targeting 1,3-beta-D-glucan synthase (bind to Fksp major subunit) to irreversibly inhibit fungal cell wall synthesis
- Resistance
 - No activities for C. neoformans, Fusarium sp, Mucorales
 - Intrinsic resistance seen in C. parapsilosis species complex, C. guilliermondii
 - Acquired resistance found in C. glabrata, C. albicans, C. tropicalis
- Mechanisms of resistance
 - FKS1 gene mutations (C. albicans)
 - FKS1 and FKS2 mutations (C. glabrata)



ANTIFUNGAL DRUG SUSCEPTIBILITY TESTING (AFST)



Antifungal susceptibility testing (AFST)

 To measure the lowest concentration of a drug that inhibits the growth of the organism, so called Minimum Inhibitory Concentration (MIC)



Why we perform susceptibility testing?

- To reliably estimate antimicrobial activities against pathogens
- To correlate with in vivo activity and to predict likelihood of outcome of therapy
- To survey/monitor resistance development
- To provide spectrum of activity of newly developed agents



Important factors about antifungal susceptibility testing (AFST)

- Host factors often more important than AST results in determining clinical outcomes
 - Underline conditions
 - PK and PD (Pharmacokinetics and pharmacodynamics)
- In vitro AST results do not 100% predict successful treatment
- Resistance described in vitro often predicts clinical failure



IDSA Candidiasis guideline

- Recent surveillance studies suggest that triazole resistance among *C. glabrata* isolates has increased to a degree that is it difficult to rely upon these agents for therapy in the absence of susceptibility testing
- A similar trend has begun to emerge for a smaller proportion of *C. glabrata* isolates and the echinocandins
- Because of these trends, susceptibility testing is increasingly used to guide the management of candidemia and invasive candidiasis

Pappas et al. Clin Infect Dis 2015 doi: 10.1093/cid/civ933.



IDSA Candidiasis guideline

- Recommended to test for azole susceptibility for all bloodstream and other clinically relevant *Candida* isolates
 - Those from sterile sites
 - Non-sterile may be clinically relevant
 - Neutropenic patients
- Consider echinocandin susceptibility testing
 - C. glabrata or C. parapsilosis infections
 - Prior echinocandin exposure

Pappas et al. Clin Infect Dis 2016.



Who sets susceptibility testing standards in the US?

• Clinical and Laboratory Standards Institute (CLSI)

	Method	Standards
Yeast	M27-A4 (broth dilution)	MCO Ord
	M44-A3 (disk diffusion)	M60 2 nd
Mold (filamentous fungi)	M38-A3 (broth dilution)	NACA Ord
	M51-A (disk diffusion)	M61 2 nd



Antifungal drug susceptibility testing methods

- Visual measurement (CLSI broth microdilution method M27-A4, M38-A3)
- Disk diffusion (CLSI M44-A3, M51A)
- Colorimetrical (YeastOne Sensititre)
- Gradient diffusion (E-test)
- Automated (Vitek 2, bioMerieux)
- Optical density (EUCAST)

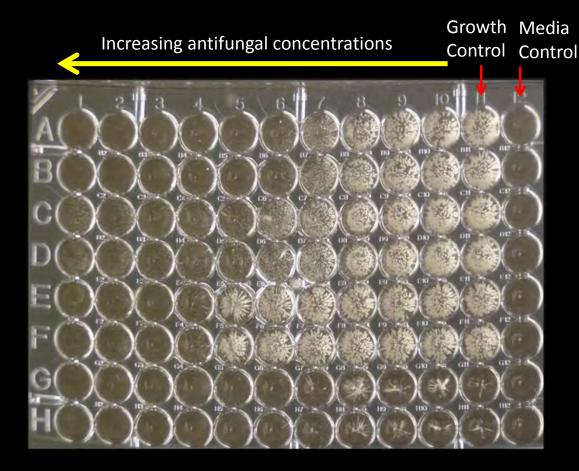


Test methods used in the Clinical Labs in the USA as per recent CAP survey (2015 – 2018)

Test methods	Participant response (%)			
	2016	2017	2018	2019
Vitek 2	152 (42%)	172 (42%)	198 (44%)	214 (46%)
YeastOne colorimetric microdilution	155 (42%)	162 (40%)	171 (38%)	182 (39%)
Gradient diffusion strips (e.g. E-test, MTS)	28 (8%)	36 (9%)	36 (8%)	34 (7%)
Broth microdilution	21 (6%)	21 (5)	26 (6%)	19 (4%)
Disk Diffusion	6 (2%)	6 (1%)	8 (2%)	9 (2%)
Other	4 (1%)	8 (2%)	8 (2%)	8 (2%)
Total	366	405	447	466



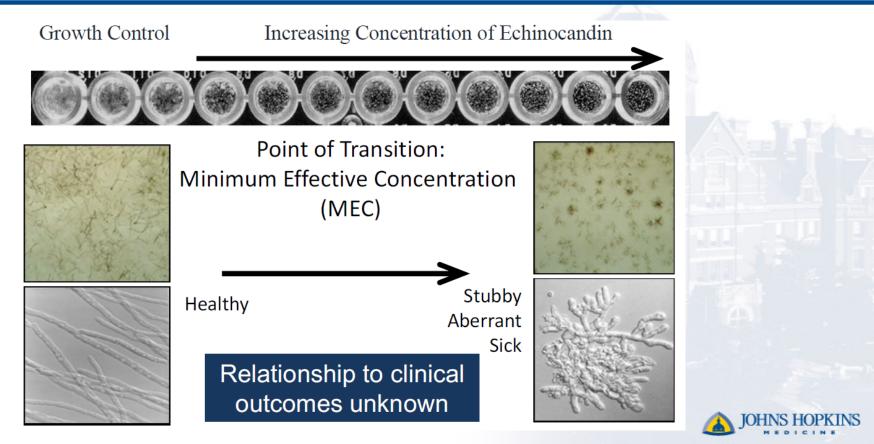
CLSI broth microdilution susceptibility testing



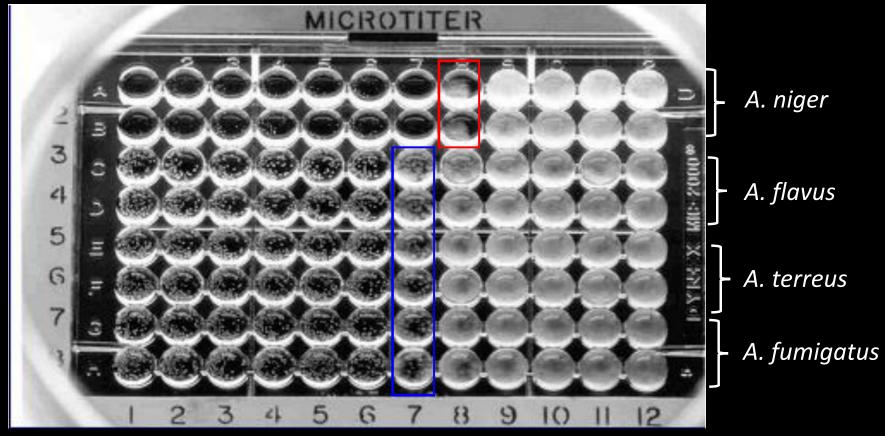
Endpoints

- Amphotericin B: 100% inhibition
- Azoles: 50% inhibition for yeast; 100% inhibition for mold
- Echinocandins: 50% inhibition for yeast; MEC for mold

Echinocandins: MEC (Minimum Effective Concentration) for filamentous fungi



Minimal Effective Concentration (MEC) of Caspofungin (CLSI M38-A3)

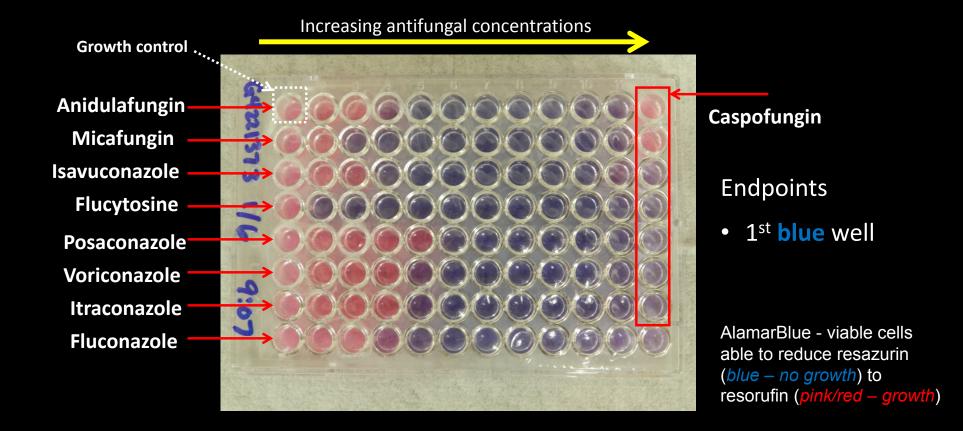


Limitations for broth microdilution

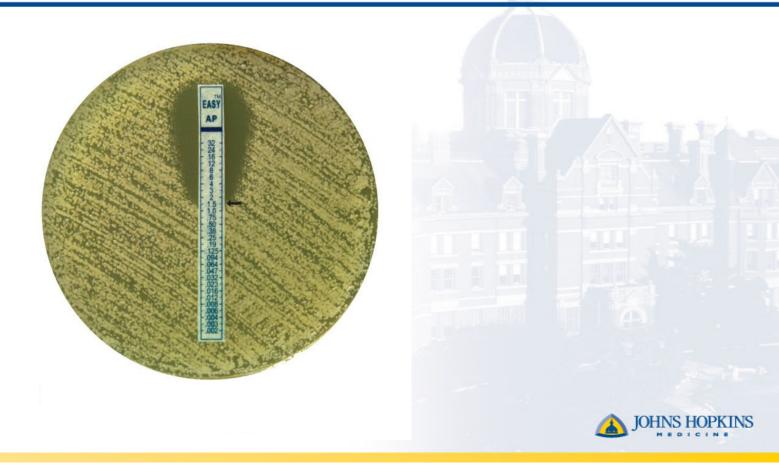
- Time consuming to prepare in house
- Availability of pure antifungal compounds
- Lack of standards in materials/reagents used
- Human errors



YeastOne Colorimetric assay (Thermo Scientific TREK Diagnostics)



E-test (agar diffusion method with strips containing predefined antifungal concentration gradient): Amphotericin B



Clinical Breakpoint (CBPs)

- Threshold MIC value established to classify microbes as susceptible or resistant to a drug based on
 - MIC distribution
 - Pharmacokinetics/pharmacodynamics
 - Clinical outcomes
- In the US, CBPs are set up by
 - Clinical and Laboratory Standards Institute (CLSI) antifungal subcommittee
 - CLSI M60 2nd (yeast); CLSI M61 2nd (mold)
 - FDA



CLSI M60 2nd (2020): CBPs for *In vitro* susceptibility testing of *Candida* sp. and selected **azoles** after 24h incubation

		MIC Breakpoints and Interpretive Categories, μg/mL			tive
Antifungal Agent ^a	Species	S	I ^b	SDD ^c	R
Fluconazole ^{2,c}	C. albicans	≤ 2	_	4	≥ 8
	C. glabrata ^g	_	_	<32	>64
	C. krusei ^h	_	—	—	_
	C. parapsilosis ^e	≤ 2	—	4	≥ 8
	C. tropicalis	≤2	_	4	<u>≥</u> 8
Voriconazole ^{3,d}	C. albicans	≤0.12	0.25-0.5	_	≥1
	C. glabrata ⁱ	—	—	—	—
	C. krusei	≤ 0.5	1	—	≥ 2
	C. parapsilosis ^e	≤0.12	0.25-0.5	_	≥ 1
	C. tropicalis	≤0.12	0.25-0.5		≥ 1



CLSI M60 2nd (2020): CBPs for *In vitro* susceptibility testing of *Candida* sp. and selected **echinocandins** after 24h incubation

		MIC Range (µg/mL)		
Antifungal Agent	Species	S	I ^a	R
	C. albicans	≤0.25	0.5	≥ 1
	C. glabrata	≤ 0.12	0.25	≥0.5
Anidulafungin ^b	C. tropicalis	≤0.25	0.5	≥1
0	C. krusei	≤0.25	0.5	≥ 1
	C. parapsilosis	≤ 2	4	≥ 8
	C. guilliermondii	≤ 2	4	≥ 8
	C. albicans	≤ 0.25	0.5	≥1
	C. glabrata	≤ 0.12	0.25	≥0.5
Caspofungin ^{b,c}	C. tropicalis	≤0.25	0.5	≥1
1 0	C. krusei	≤0.25	0.5	≥1
	C. parapsilosis	≤ 2	4	≥ 8
	C. guilliermondii	≤ 2	4	≥ 8
	C. albicans	≤0.25	0.5	≥1
	C. glabrata	≤0.06	0.12	≥0.25
Micafungin ^b	C. tropicalis	≤0.25	0.5	≥ 1
_	C. krusei	≤0.25	0.5	≥1
	C. parapsilosis	≤ 2	4	≥ 8
	C. guilliermondii	≤ 2	4	≥ 8



CLSI M61 2nd (2020): CBPs for Aspergillus fumigatus

		MIC Breakpoints and		ıd
		Interpretive Categories, µg/mL		ug/mL
Antifungal Agent	Species	S	Ι	R
Voriconazole ^a	A. fumigatus	≤0.5	1	≥2

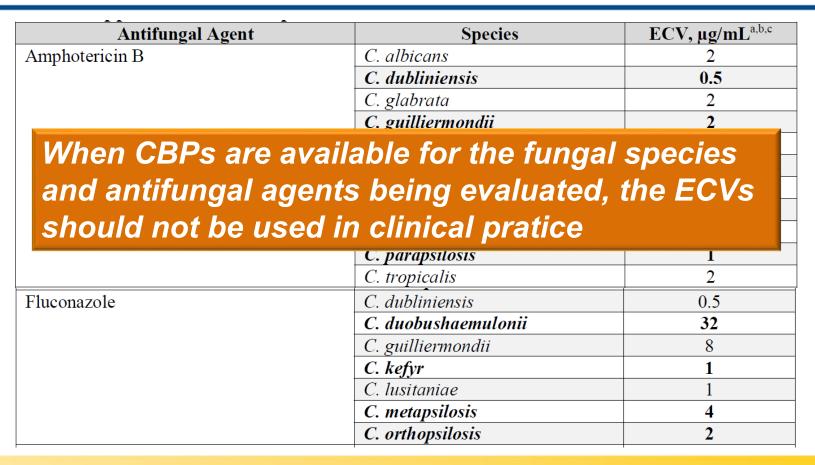


Epidemiology cut-off value (ECV)

- ECV are MIC values that separate organism into those with wild-type (WT) and non-wild-type (NWT) population based on in vitro MIC data only.
 - WT: MIC result is consistent with no acquired mutational resistance
 - NWT: MIC result is consistent with either acquired or mutational resistance
 - Method: CLSI M57 (Principles and procedures for the development of ECVs for AFST)
 - Standards: CLSI M59 3rd ED (ECVs for AFST)
- ECVs do not predict clinical outcome to therapy as clinical breakpoints do



CLSI M59 3rd ED (2020): ECVs for In vitro susceptibility testing of *Candida* sp. with no CBPs



OHNS HOPKINS

CLSI M59 3rd ED (2020): ECVs for In vitro susceptibility testing of *Candida* sp. with no CBPs

- C. parapsilosis complex (C. parapsilosis, C. orthopsilosis, C. metapsilosis)
 - If the clinical labs are now able to identify the subspecies within the complex (mostly by MALDI), then *C. parapsilosis* CBP should not be applied to *C. orthorpsilosis* and *C. metapsilosis* (CLSI M60Ed2). Instead, ECVs should be applied (CLSI M59Ed3)
 - However, identification of sub-species is not possible, *C. parapsilosis* CBP may still be applied since the prevalence of *C. orthropsilosis* or *C. metapsilosis* is still very low (may vary in different institutions).
- More antifungal agent ECVs are added to Candida species without CBP (CLSI M59Ed3)



CLSI M59 3rd ED (2020): ECVs for In vitro susceptibility testing of *Cryptococcus* spp.



- No CBP for *Cryptococcus*, but ECVs are available but interpretation based on genotype and has recent nomenclature change (CLSI M59Ed3)
- If the lab identified *C. neoformans*, it may still apply ECVs established for *C. neoformans* VNI since VNI is the most common molecular genotype of *C. neoformans* (may include a note when report).
- If the lab identified *C. gattii*, it may be challenging in deciding which ECVs to apply since most clinical labs don't have capacity to get down to genotype VGI vs VGII (new name *C. deuterogattii*)

Antifungal Agent	Species (Genotype)	ECV, μg/mL ^{b,c}
Amphotericin B	C. gattii (VGI)	0.5
	C. deuterogattii (formerly C. gattii) (VGII)	1
	C. neoformans (VNI)	0.5
Fluconazole	C. gattii (VGI)	16
	C. deuterogattii (formerly C. gattii) (VGII)	32
	C. neoformans (VNI)	8
Flucytosine	C. gattii (VGI)	4
	C. deuterogattii (formerly C. gattii) (VGII)	32
	C. neoformans (VNI)	8
Itraconazole	C. gattii (VGI)	0.5
	C. deuterogattii (formerly C. gattii) (VGII)	1
	C. neoformans (VNI)	0.25
Posaconazole	C. neoformans (VNI)	0.25
Voriconazole	C. gattii (VGI)	0.5
	C. deuterogattii (formerly C. gattii) (VGII)	0.5
	C montaining $(VNII)$	
	C. neoformans (VNI)	0.25
C. neoformans	C. neoformans (VNI, C. deneoformans (VN C. gattii (VGI) C. deuterogattii (VGII	VNII) NV)
C. neoformans C. gattii —	C. neoformans (VNI, C. deneoformans (VN C. gattii (VGI)	∨NII) JI∨)) I)

Hagen F. et al. Fungal Gen Biololgy 2015; 78:16

CLSI M59 3rd ED(2020): ECVs for In vitro susceptibility testing of Aspergillus spp.

Antifungal Agent	Species	ECV (µg/mL) ^{‡§}
	A. flavus	4
	A. fumigatus	2
Amphotericin B	A. niger	2
	A. terreus	4
	A. versicolor	2
	A. flavus	0.5
Caspofungin [¶]	A. fumigatus	0.5
Casporungin	A. niger	0.25
	A. terreus	0.12
	A. flavus	1
Isavuconazole	A. fumigatus	1
Isavuconazore	A. niger	4
	A. terreus	1
	A. flavus	1
Itraconazole	A. fumigatus	1
111 aconazore	A. niger	4
	A. terreus	2
	A. flavus	0.5
Posaconazole	A. niger	2
	A. terreus	1
	A. flavus	2
Voriconazole	A. fumigatus	1
V OI ICOIIAZOIC	A. niger	2
	A. terreus	2



New CAP Checklist Questions

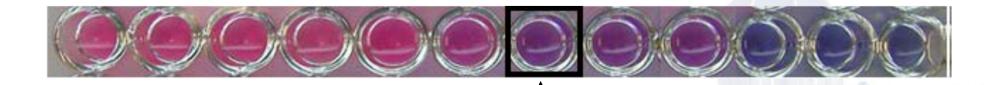
Question	Description	
MIC.42700	Inconsistent Antifungal Susceptibility Reports (PHASE I)	
	A policy addresses unusual or inconsistent antifungal testing	5
	results.	the second
	Results from testing of patient isolates should be reviewed,	
	and unusual or inconsistent results should be investigated.	
	Each laboratory should have a policy for confirming unusual or	
	inconsistent results. For yeasts and moulds, the time of	
	endpoint reading (particularly for the echinocandins) and the	
	effect of trailing growth (particularly for the azoles and	HIT IS AN AN
	flucytosine) can be significant factors impacting susceptibility	
	results. In some cases, it may be necessary to repeat	
	susceptibility testing and/or identification procedures to	
	confirm initial results. This may involve using alternative	
	testing methods or sending the isolate to a reference lab.	
		JOHNS HOPK

Unusual or inconsistent AFST results

- Candida albicans R to all azoles
- Candida albicans R to echinocandins
- Candida glabrata S to azoles but R to echinocandins



Trailing growth seenin YeastOne Sensititre plate

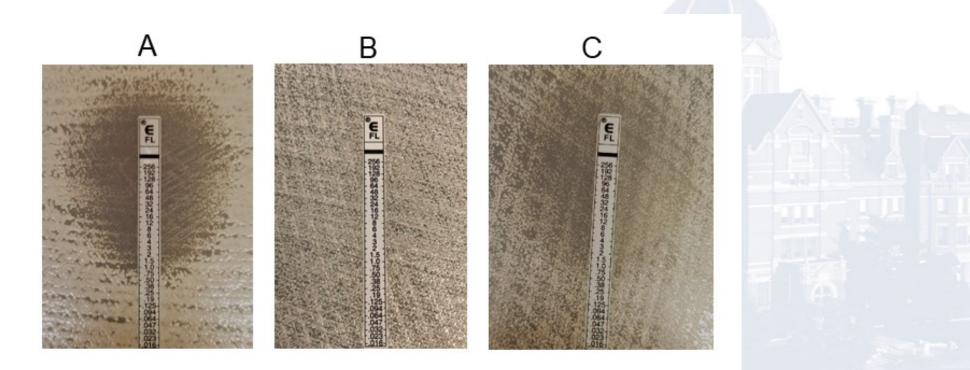


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- A slight color change persists above the MIC and it is often identical for several or all drug concentrations above the MIC.
- The MIC should be read as the first well showing a less intense color change compared to the more positive growth wells of the lower concentrations.
- Commonly seen in azoles
- Overcall R or false R



Trailing growth seen in E-Test



Lockhart S. Clin Micro Newsletter 2019



Should the clinical lab be testing Caspofungin for *Candida* spp.?

- Significant inter-laboratory variability
- Overall resistant
- EUCAST does not recommend testing it
- Anidulafungin & Micafungin can serve as surrogate markers for caspofungin



Espinel-Ingroff et al. AAC 2013; 57:5836. Pfaller et al. JCM 2014; 52:3223

Intrinsic resistance (IR) organisms (CLIS M59 3rd Ed)

- Candida krusei: IR to fluconazole
- Cryptococcus neoformans/C. gattii: IR to echinocandins (anidulafungin, caspofungin, micafungin)
- Aspergillus spp.: IR to fluconazole, flucytosine
- These antifungal drugs should not be tested and need to be reported as R



EMERGING ANTIFUNGAL DRUG RESISTANT ORGANISMS



Candida auris: Here's what you should know about the superbug fungus spreading worldwide

There are three main reasons we should be worried about C. auris infection

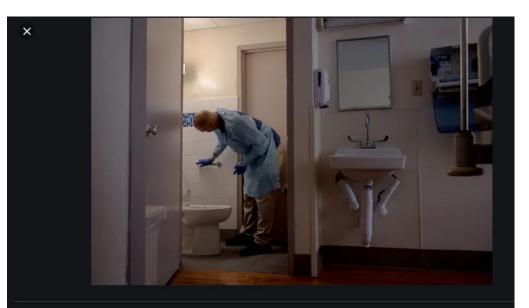


BY **BAILEY KING** PhillyVoice Staff



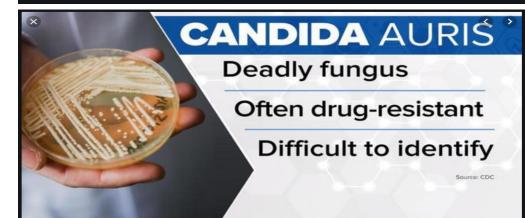


SOURCE/CENTERS FOR DISEASE CONTROL AND PREVENTION



The New York Times

Candida Auris: The Fungus Nobody Wants to Talk About - The ...



Multi-drug resistant Candida auris

- Resistant ≥2 antifungal classes
- Cause invasive infections with high mortality

 Up to 60% patients infected with C. auris died (CID,2017;64:134)
- Spread in healthcare settings and cause healthcare-associated outbreaks
 - Colonize patients' skin and other body sites indefinitely; patients can continue to be colonized with *C. auris* despite daily chlorhexidine bathing (Antimicrob Resist Infect Control 2016;5:35)
 - Persist in the healthcare environment for very long time (JCM 2017 Jul 26)
- Difficulty to identify (close to C. haemulonii, C. duobushaemulonii)



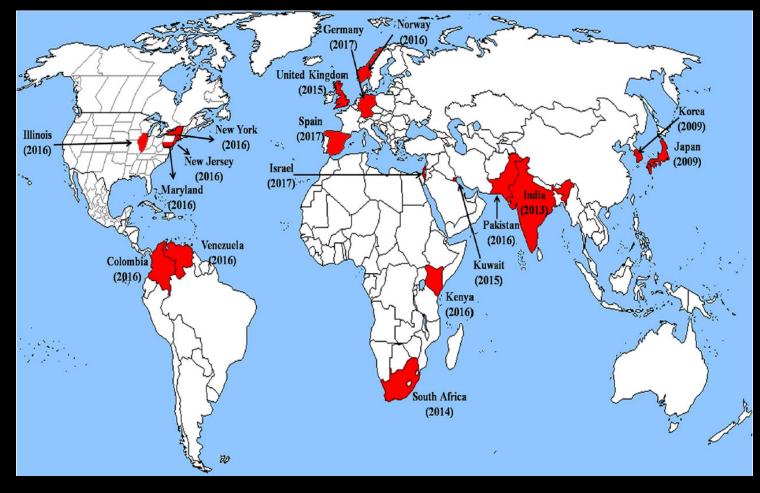
C. auris (Auris: Latin word for ear): history and epidemiology

- First recognized in 2009 from an ear canal specimen in Japan; reservoir is unknown.
 Earliest strain actually dated back to 1996 from a BSI in a child in Korean
- No single strain for the widespread is identified; whole-genome sequencing revealed four distinct geographical clades
 - South Asia (Clade I), East Asia (Clade II), Africa (Clade III), South America (Clade IV)
- A nosocomial pathogen
 - First outbreak in healthcare settings was reported in an ICU in UK in 2015
 - In a span of only 7 years, it has become widespread across a dozen countries causing a severe healthcare-associated invasive fungal infections
 - The transmission of *C. auris* in one hospital outbreak was found to be linked to reusable axillary temperature probes, indicating that this emerging pathogen can persist in the environment and be transmitted in health care settings (N. Eng J. Med 2018; 379:1322)

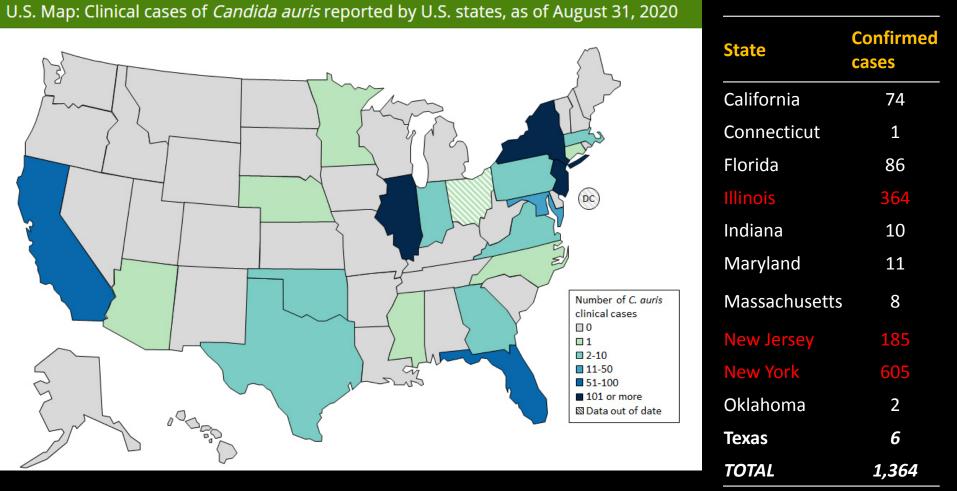
Bradley SF. JAMA 2019; Antimicrob Resist Infect Control 2016;5:35; Microbiol Immunol 2009;53:41



A global map depicting rapid emergence of multidrug-resistant clinical *Candida auris* strains in 33 countries across 5 continents



Bradley SF. JAMA 2019 Chowdhary A et al. PLOS Pathogens 2017 13(5): e1006290



Most US isolates are closely related to isolates from South Asia (India, Pakistan), East Asia (South Korea, Japan), South African and South America (Venezuela) by whole-genome sequencing analysis (www.cdc.gov/fungal/diseases/candidiasis/)

C. auris antifungal resistance

- Tentative MIC breakpoints (µg/mL) (CDC recommendation)
 - Fluconazole: ≥32
 - − Amphotericin B: \geq 2
 - Caspofungin: ≥2
 - Micafungin and
 Anidulafungin: ≥4

54 isolates from 3 continents		123 isolates in India	
MIC Range	MIC ₉₀	MIC Range	MIC ₉₀
0.38 - 4	2	0.125 - 8	2
4 - 256	256	4 - ≥64	≥64
0.03 - 16	8	0.03 - 16	4
0.06 - 1	1	0.016 - 8	0.125
0.06 - 4	2	0.015 - 8	0.25
0.125 - 16	1	0.015 - 8	0.5
0.03 - 16	1		
0.125 - 128	0.5		
	continents MIC Range 0.38 - 4 4 - 256 0.03 - 16 0.06 - 1 0.06 - 4 0.125 - 16 0.03 - 16	continents MIC Range MIC ₉₀ 0.38 - 4 2 4 - 256 256 0.03 - 16 8 0.06 - 1 1 0.06 - 4 2 0.125 - 16 1 0.03 - 16 1	continentsMIC $_{90}$ MIC RangeMIC Range0.125 - 80.38 - 420.125 - 84 - 2562564 - ≥640.03 - 1680.03 - 160.06 - 110.016 - 80.06 - 420.015 - 80.125 - 1610.015 - 80.03 - 161

Lockhart et al. CID 2017; 64:134 Arendrup MC et al. Antimicrob. Agents Chemother. 2017;61:e00485-17



C. auris antifungal resistance

- ~80% R to fluconazole; >50% R to voriconazole; 30-40% R to amphotericin B; 1-7% R to echinocandins; ~40% R to 2 antifungal classes; R to all three classes has been observed.
- Resistant mechanisms
 - Erg11 mutation (Y132F, K143R)
 - ABC, CDR1 efflux transporter (deletion of CDR1 abrogates resistance)
 - FKS1 HS1 (S639F)
- Most *C. auris* isolates susceptible to echinocandins, but acquired R to echinocandin could become more common
 - In one patient, R to echinocandin drugs developed while being treated with echinocandins

CID 2017; 64:134; Perlin DS Lancet Infect Dis. 2017 July 31); MMWR 2017; 66:514; Rybak et al. AAC 2019 63(4)



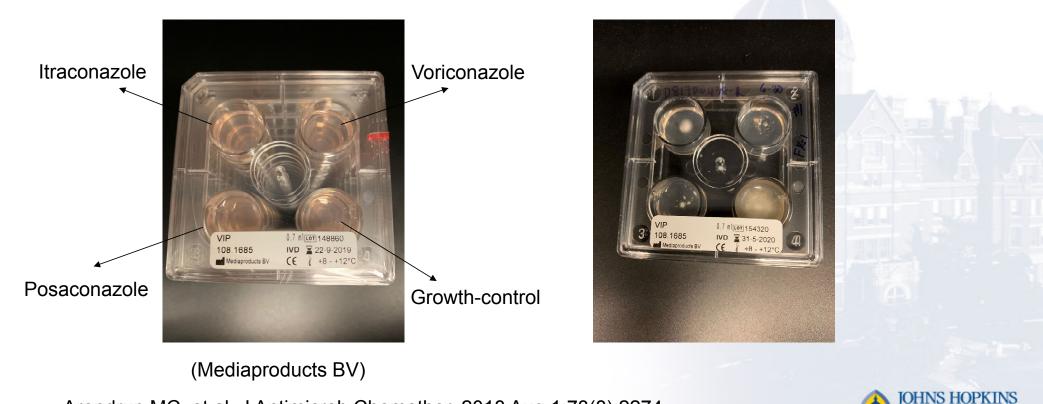
Azole resistant *Aspergillus*

- Surveillance
 - In the US, 1.4% (19/1356) *A. fumigatus* isolates from 2015 to 2017 showed elevated MIC against azoles, of them 5 harbored *Cyp51A* mutations (TR34/L98H)
 - In the Netherlands, 11% (508/4496) of *A. fumigatus* isolates from 2013 2018 showed azole resiatance; resistant rate increased from 7.6% in 2013 to 14.7% in 2018.
- Source
 - Prolong azole therapy
 - Environmental derived azole resistant ones (due to agricultural usage of azoles)
- The actual azole R *Aspergillus* in the US may be underestimated since most clinical labs do not routinely testing azoles in *Aspergillus* clinical isolates

Berkow EL, et al. AAC 2018, 26;62(5):e02240-17; Lestrade PPA et al. Emerg Infect Dis. 2020 Jul;26(7):1447-1455.



VIP (<u>V</u>oriconazole-<u>I</u>traconazole-<u>P</u>osaconazole) plate: screen azole resistant *A. fumigatus* ≤ 48h



Arendrup MC, et al. J Antimicrob Chemother. 2018 Aug 1;73(8):2274.

Summary

- Amphotericin B resistance is still rare; echinocandin resistance is mostly associated with target mutation and is emerging. Azole resistant mechanism is multimodal and more commonly seen.
- Clinical labs should embrace AFST capacity and utilize clinical breakpoints and ECVs to aid clinicians to choose appropriate antifungal drugs for treatment
- Clinical labs should be aware of emerging antifungal drug resistant fungal pathogens and be able to detect them

