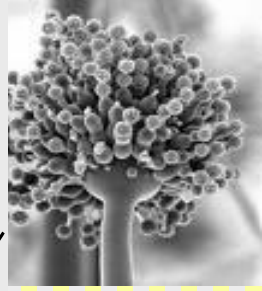




*XIX. Türk Klinik Mikrobiyoloji ve
İnfeksiyon Hastalıkları Kongresi*

28-31 Mart 2018, Antalya



Antifungal Direnç Ne Kadar Sorun?

Prof. Dr. Sevtap Arıkan Akdağlı, FESCMID, FECMM
Hacettepe Üniversitesi Tıp Fakültesi
Tıbbi Mikrobiyoloji Anabilim Dalı

★
Temel bilgiler

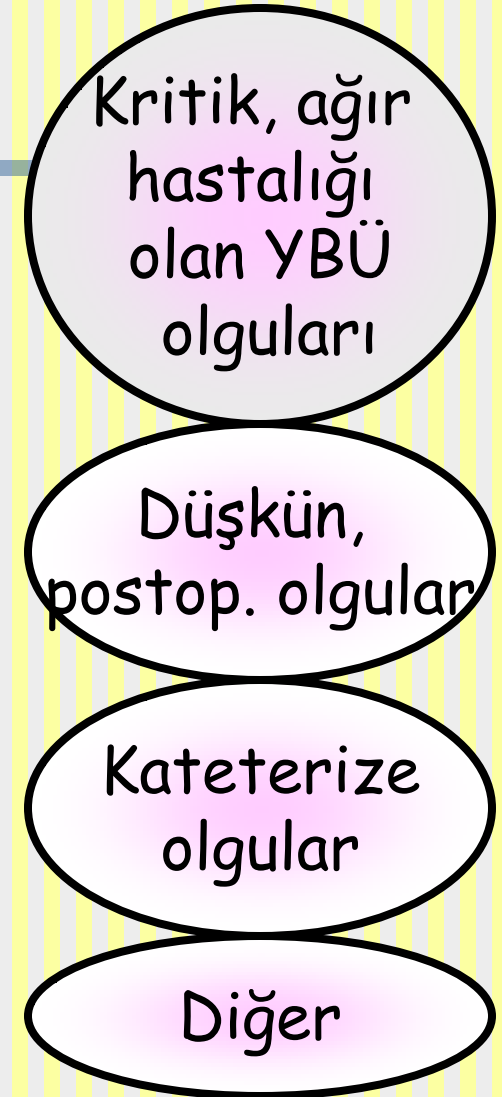
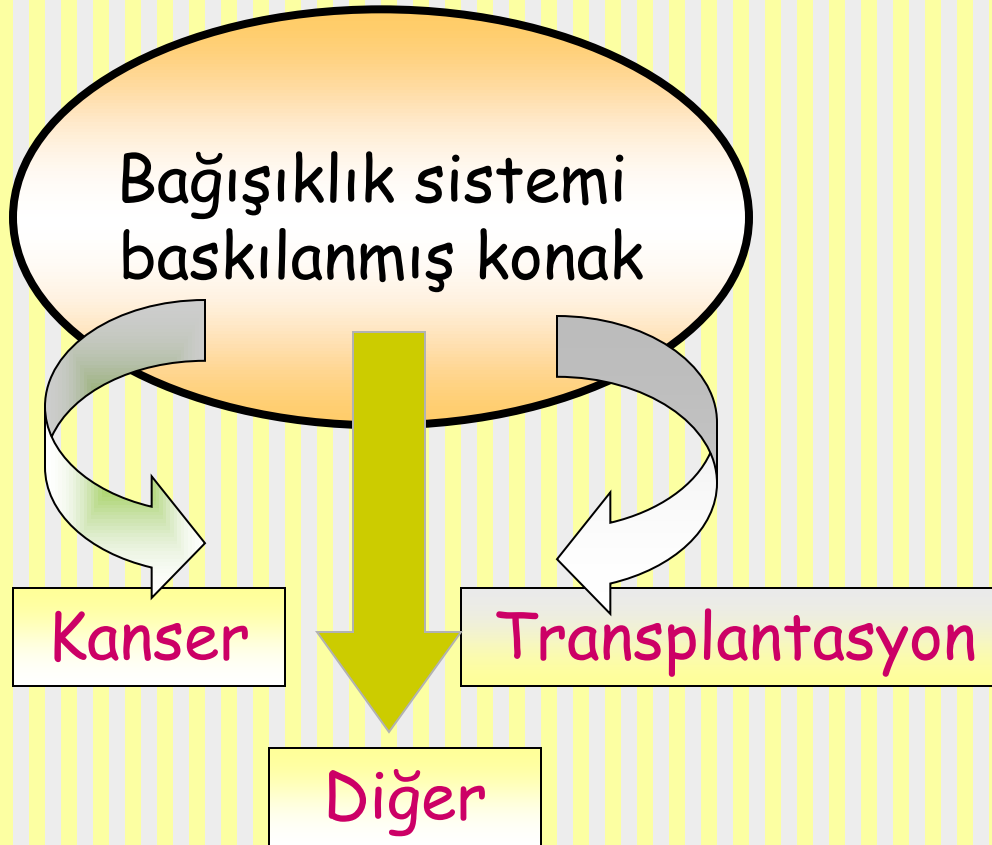
Sunum Planı

YÖNTEM ve
DEĞERLENDİRME

DİRENÇ VARLIĞI /
ORANI

ÖNERİLER

Fırsatçı mantar enfeksiyonları: KONAK



"Özel konak" koşulları, antifungal **direnç** riskini beraberinde getirir

Antifungal profilaksi

Tekrarlayan fungal enf. -
tekrarlayan antifungal tdv.

Yabancı cisim -biyofilm oluşumu

DİRENÇ

Özel konakta Hangi mantar?

SIK İZOLE EDİLEN ETKENLER

- Yanık
- YBÜ
- Yabancı cisim
- DM
- Uç yaş grupları
- Anti-TNF ve diğer immünsupresan tedaviler
- SOT
- HIV
- Hematoonkolojik malignansi
- KİT, PKHT
-

CANDIDA
ASPERGILLUS

DİĞER

Mucorales

C. neoformans

Fusarium

P. jirovecii

Scedosporium

Dematisiyöz

küfler.....

1950'lerden günümüze: Değişen antifungal ilaç spektrumu

1950'ler

Grizeofulvin
İlk azol

1960'lar

Amfoterisin B
Mikonazol
Klotrimazol
Flusitozin

1970'ler

Ekonazol
Mikonazol (IV)

1980'ler

Ketokonazol
(po)

1990'lar

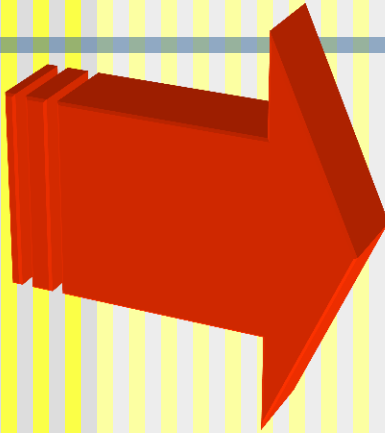
Flukonazol
Itrakonazol
Terbinafin
Lipid Amfo
bileşikleri

2000'ler

Kaspofungin
Vorikonazol

Mikafungin
Anidulafungin
Posakonazol

.....



Candida Aspergillus

SUŞLARINDA

ANTİFUNGAL DİRENÇ

Candida

Ekinokandin

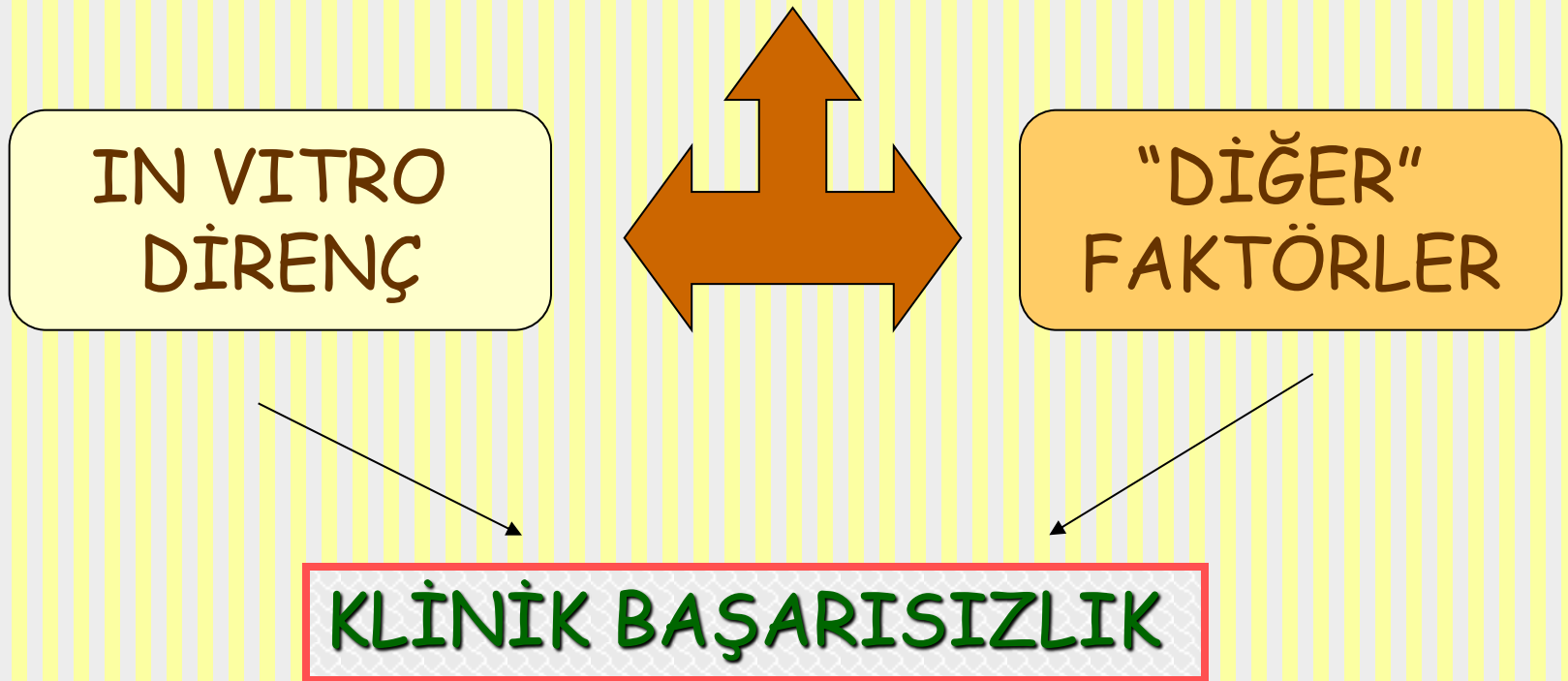
Multipl ilaç direnci

Aspergillus

Azol

Ekinokandin

Mantar Enfeksiyonlarında Klinik Başarısızlığın Nedenleri



"Diğer" faktörler...

MANTAR

- Direnç
- Hücre tipi
- İnokulum miktarı

İLAÇ

- Yanlış doz
- Fungistatik etki
- Yetersiz emilim, dağılım ve metabolizma
- İlaç etkileşimleri

- İmmün yanıt
- Yabancı cisim
- Enfeksiyon bölgesi
 - Apse
- Tedaviye uyumsuzluk

KONAK

Antifungal direncin getirdiği sorunlar

- Tedavi başarısına olumsuz etki
 - Çapraz direnç olasılığı
- Tedavi seçeneklerinin kısıtlanması
(Genel durumu, ilaç etkileşimleri, biyoyararlanım, mevcut formülasyon,... ve direnç ...)

Primer ve Sekonder direnç

Direnç türü	Örnek
PRİMER (DOĞAL)  Tür tanımlaması	<i>C. krusei</i> -flukonazol <i>C. glabrata</i> -fukonazol <i>C. norvegensis</i> -flukonazol <i>C. lusitaniae</i> -amfoterisin B <i>C. krusei</i> -flusitozin Aspergillus-flukonazol Mucorales-vorikonazol
SEKONDER (EDİNİLMİŞ)	<i>C. albicans</i> (-orofaringiyal kandidoz-HIV)-flukonazol <i>C. glabrata</i> -ekinokandin Aspergillus-ITC/VCZ/POS ...

TABLE 3. Intrinsic susceptibility pattern for *Candida* species

EUCAST breakpoints have been established for the common *Candida* species allowing classification of wild-type isolates into S, I and R categories. For the uncommon *Candida* species, breakpoints have not been established. For these species, an 'X' denotes that the MICs for the antifungal compound are elevated compared with those for *Candida albicans*

	AMB	ECS	FCO	Comments
Common <i>Candida</i> species				
<i>C. albicans</i>	S	S	S	
<i>C. dubliniensis</i>	S	S	S	Closely related to <i>C. albicans</i> ; fluconazole resistance acquired more easily
<i>C. glabrata</i>	S	S	I	Efflux pumps often induced during azole therapy
<i>C. krusei</i>	S	S	R	
<i>C. parapsilosis</i>	S	I	S	
<i>C. tropicalis</i>	S	S	S	
Uncommon <i>Candida</i> species				
<i>C. lusitanae</i>	X			
<i>C. fermentati</i>		X		
<i>C. guilliermondii</i>		X	X	
<i>C. metapsilosis</i>		X		Closely related to <i>C. parapsilosis</i>
<i>C. orthopsilosis</i>		X		Closely related to <i>C. parapsilosis</i>
<i>C. cifferrii</i>			X	
<i>C. inconspicua</i>			X	
<i>C. humicola</i>			X	
<i>C. lambica</i>			X	
<i>C. lipolytica</i>			X	
<i>C. norvegensis</i>			X	
<i>C. palmiophila</i>			X	
<i>C. rugosa</i>			X	
<i>C. valida</i>			X	
<i>S. cerevisiae</i> ^a			X	Closely related to <i>C. glabrata</i>

ECS: echinocandins; FCO: fluconazole.

^aAnamorphic state is *C. robusta*.

TABLE 2. Intrinsic resistance (R) and variable (V) susceptibility in *Aspergillus*

	AMB	Azoles	Echinocandins
<i>Aspergillus</i> section <i>fumigati</i>			
<i>A. fumigatiaffinis</i>	R	R	
<i>A. lentulus</i>	R	R	V
<i>N. pseudofischeri</i>	V	R	
<i>A. viridinutans</i>	R	R	
<i>N. udagawae</i>	R	R (vor)	
<i>A. terreus</i> (and <i>A. alabamensis</i>)	R		
<i>A. flavus</i>	R	R	
<i>A. versicolor</i> (and <i>A. sydowi</i>)	R	V	
<i>A. calidoustus</i>		R	V
<i>A. alilaceus</i>	V		V

Data compiled from [12–15].

Vor, Voriconazole.

Sunum Planı

Temel bilgiler



YÖNTEM ve
DEĞERLENDİRME



DİRENÇ VARLIĞI /
ORANI



ÖNERİLER

Direncin in vitro saptanması: Antifungal duyarlılık testleri-I

CLSI M27-A3

Candida, Cryptococcus

CLSI M38-A2

Aspergillus, Rhizopus, Fusarium, P. boydii,
S. schenckii-küf formu, dermatofitler

CLSI M44-A2

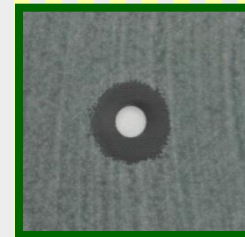
Candida

CLSI M60

M27 ve M44; Kalite kontrol değerleri

CLSI M51-A

Dermatofit dışı küfler



Direncin in vitro saptanması: Antifungal duyarlılık testleri-II

EUCAST E.Dis 7.3.1

Candida

15 Ocak 2017



EUCAST E.Dis 9.3.1

Konidyum oluşturan küfler

15 Ocak 2017

DIN-58940-84

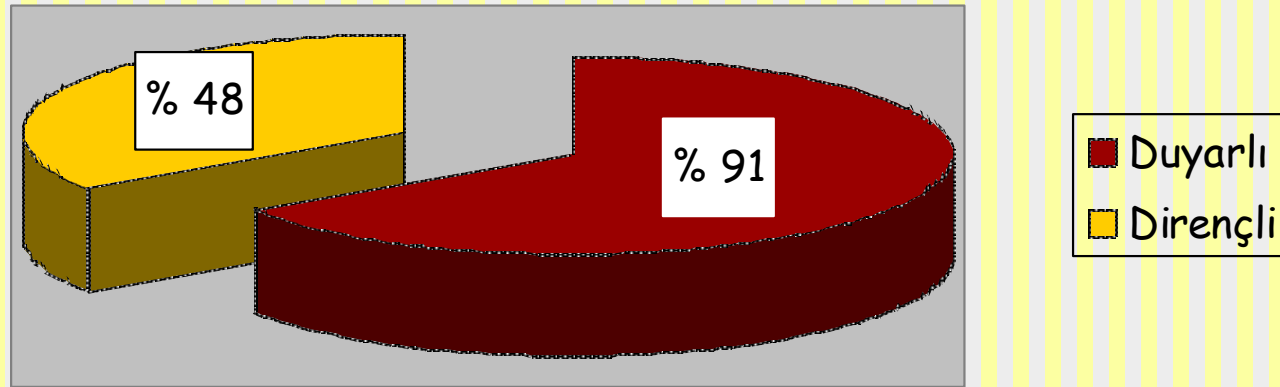
Duyarlılık testlerinin rutin amaçlı uygulanma endikasyonları

(Candida)

- İnvazif enfeksiyon; steril bölgeden izolasyon
- Direncin görülebildiği türler (örn. *C. glabrata* - flukonazol)
- (Beklenmeyen) klinik yanıtsızlık
- Tedaviye bağlı sekonder direnç gelişimi riski nedeniyle izlem için
- Epidemiyolojik veri (her merkez - periyodik)

Ancak... Duyarlılık testi sonucu klinik yanıtı ne kadar yansıtıyor ?

Meta
analiz



Candida - flukonazol

Candida - itrakonazol

Candida - ketokonazol

C. neoformans - flukonazol

Histoplasma - flukonazol

Duyarlılık testlerinin rutinde uygulanması -KÜFler

(Aspergillus)

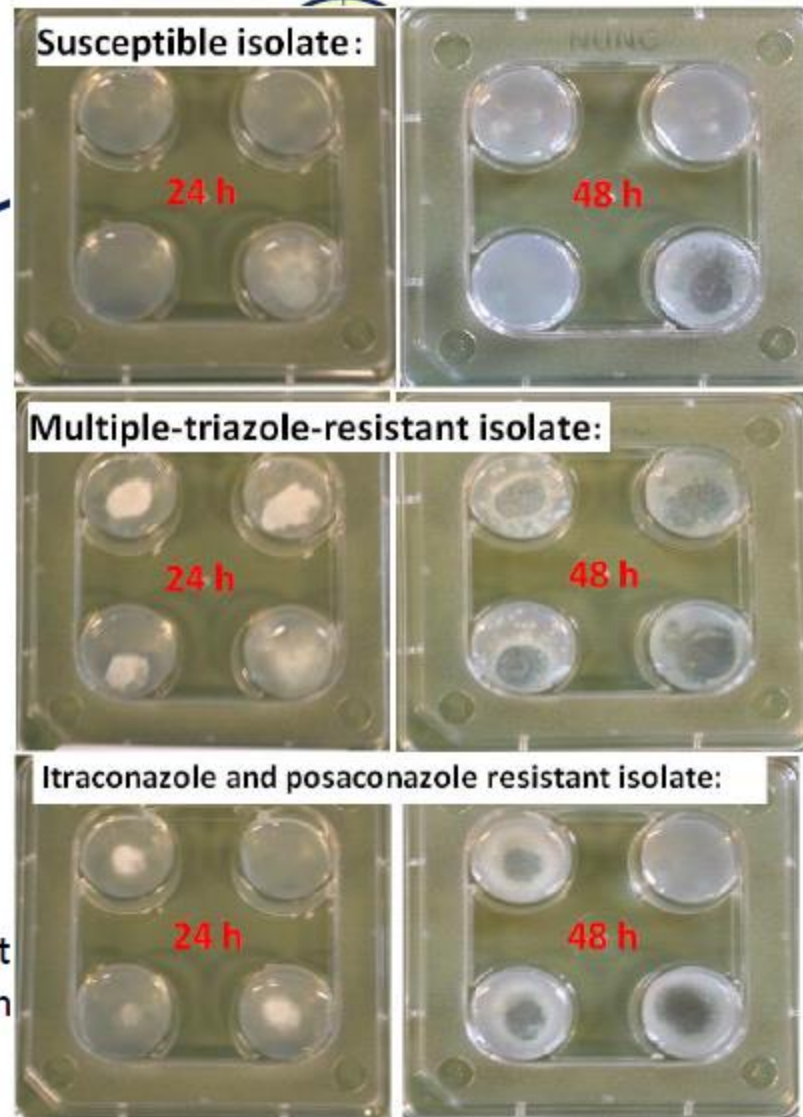
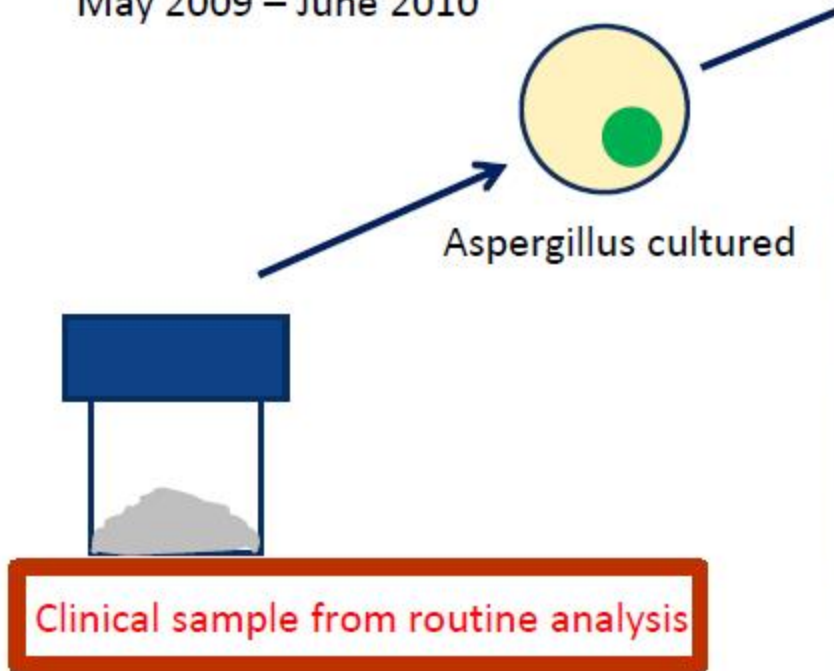
❑ CLSI M38-A2

❑ EUCAST E.Dis 9.3.1

15 Ocak 2017

Surveillance-method

Prospective, multicenter surveillance
May 2009 – June 2010



• Compleet
questionn

www.umcn.nl

• Cyp-sequencing

Courtesy of Paul Verweij & Jan van der Linden

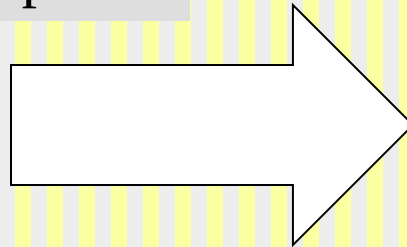
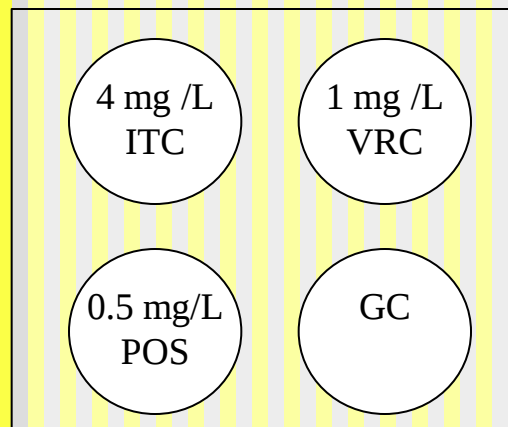
J.W.M. van der Linden, M.C. Arendrup,
A. Warris, K. Lagrou, H. Pelloux, P.M. Hauser,
E. Chrysanthou, E. Mellado, S.E. Kidd,
A.M. Tortorano, E. Dannaoui, P. Gaustad,
J.W. Baddley, A. Uekötter, C. Lass-Flörl,
N. Klimko, C.B. Moore, D.W. Denning,
A.C. Pasqualotto, C. Kibbler, S. Arian-Akdagli,
D. Andes, J. Meletiadis, L. Naumiuk,
M. Nucci, W.J.G. Melchers, P.E. Verweij

Prospective Multicenter International Surveillance of Azole Resistance in *Aspergillus fumigatus*

To investigate azole resistance in clinical *Aspergillus* isolates, we conducted prospective multicenter international surveillance. A total of 3,788 *Aspergillus* isolates were screened in 22 centers from 19 countries. Azole-resistant *A. fumigatus* was more frequently found (3.2% prevalence) than previously acknowledged, causing resistant invasive and noninvasive aspergillosis and severely compromising clinical use of azoles.

Multicenter validation of the 4-well plates: Agar screening method for azole R in *A.fumigatus* ¹

- 40 WT
- 39 *cyp51A* mutant
- Simulated mixed samples



Inter-plate agreement
Inter-observer agreement
EA
CA
Sens
Spec

In-house vs. commercial plates, three lab.s, two observers, scoring (0-3)

Multicenter validation of the 4-well plates: Agar screening method for azole R in A.fumigatus ₂

	% Range
Inter-observer agreement	80-100
Qualitative agreement (NG vs. G)	87-100
Sensitivity - overall	97-100
Sensitivity - simulated WT-mutant	83-100
Specificity - overall	95-100

High inter-plate agreement

Rutin Lab. kullanımı için pratik ve güvenilir

Practice Guidelines for the Diagnosis and Management of Aspergillosis: 2016 Update by the Infectious Diseases Society of America

Thomas F. Patterson,^{1,a} George R. Thompson III,² David W. Denning,³ Jay A. Fishman,⁴ Susan Hadley,^{5,b} Raoul Herbrecht,⁶ Dimitrios P. Kontoyiannis,⁷ Kieren A. Marr,⁸ Vicki A. Morrison,⁹ M. Hong Nguyen,¹⁰ Brahm H. Segal,¹¹ William J. Steinbach,¹² David A. Stevens,¹³ Thomas J. Walsh,¹⁴ John R. Wingard,¹⁵ Jo-Anne H. Young,¹⁶ and John E. Bennett^{17,a}

Clinical Infectious Diseases® 2016;63(4):433–42

When Should Antifungal Susceptibility Testing Be Performed, and How Should Results Be Interpreted and Affect Management?

Recommendation.

24. Routine antifungal susceptibility testing (AFST) of isolates recovered during initial infection is not recommended. AFST of *Aspergillus* isolates using a reference method is reserved for patients suspected to have an azole-resistant isolate or who are unresponsive to antifungal agents, or for epidemiological purposes (strong recommendation; moderate-quality evidence).

Diagnosis and management of *Aspergillus* diseases: executive summary of the 2017 ESCMID-ECMM-ERS guideline

A.J. Ullmann^{1, 62, 63}, J.M. Aguado^{2, 62, 63}, S. Arikan-Akdoglu^{3, 62, 63}, D.W. Denning^{4, 5, 6, 63}, A.H. Groll^{7, 62, 63}, K. Lagrou^{8, 62, 63}, C. Lass-Flörl^{9, 62, 63}, R.E. Lewis^{10, 62}, P. Munoz^{11, 12, 13, 62, 63}, P.E. Verweij^{14, 62, 63}, A. Warris^{15, 62, 63}, F. Ader^{16, 17, 65}, M. Akova^{18, 62, 63}, M.C. Arendrup^{19, 62, 63}, R.A. Barnes^{20, 63}, C. Beigelman-Aubry^{21, 65}, S. Blot^{22, 23, 65}, E. Bouza^{11, 12, 13, 62, 63}, R.J.M. Brüggemann^{24, 62}, D. Buchheidt^{25, 62, 63}, J. Cadranet^{26, 65}, E. Castagnola^{27, 62}, A. Chakrabarti^{28, 63}, M. Cuenca-Estrella^{29, 62, 63}, G. Dimopoulos^{30, 65}, J. Fortun^{31, 62, 63}, J.-P. Gangneux^{32, 62, 63}, J. Garbino^{33, 62, 63}, W.J. Heinz^{1, 62, 63}, R. Herbrecht^{34, 62}, C.P. Heussel^{35, 63}, C.C. Kibbler^{36, 63}, N. Klimko^{37, 63}, B.J. Kullberg^{24, 62, 63}, C. Lange^{38, 39, 40, 65}, T. Lehrnbecher^{41, 63}, J. Löffler^{1, 62, 63}, O. Lortholary^{42, 62, 63}, J. Maertens^{43, 62, 63}, O. Marchetti^{44, 45, 62, 63}, J.F. Meis^{46, 62, 63}, L. Pagano^{47, 63}, P. Ribaud⁴⁸, M. Richardson^{4, 5, 6, 62, 63}, E. Roilides^{49, 50, 62, 63}, M. Ruhnke^{51, 62, 63}, M. Sanguinetti^{52, 62, 63}, D.C. Sheppard^{53, 62, 63}, J. Sinkó^{54, 62}, A. Skiada^{55, 62, 63}, M.J.G.T. Vehreschild^{56, 57, 58, 63}, C. Viscoli^{59, 62, 63}, O.A. Cornely^{56, 58, 60, 61, 62, 63, 64, *}

Table 15

Indications for testing for azole resistance in clinical *Aspergillus* isolates

Population	Intention	Intervention	SoR	QoE	Comment	Ref.
All clinically relevant <i>Aspergillus</i> isolates (in patient groups or regions with known azole resistance)	Identify azole resistance	Reference MIC testing	A	II	In situations where rapid testing is available	[105,111,114,116,300,474–484]
Clinically relevant <i>Aspergillus</i> isolates in patient groups with high prevalence of azole resistance or patients unresponsive to treatment	Identify isolates with intrinsic resistance	Species identification to complex level	A	III	Some species are intrinsically resistant—e.g. <i>A. calidoustus</i> (azole resistant) and <i>A. terreus</i> (AmB resistant)	[103,485]
Clinically relevant <i>A. fumigatus</i> isolates	Identify azole-resistant <i>A. fumigatus</i>	Routine azole agar screening	B	III	Identifies resistant colonies that require MIC-testing	[118,486]
All isolates—resistance surveillance	Determine the local epidemiology of azole resistance	Periodical reference MIC testing of <i>A. fumigatus</i> complex	A	II	Test at least 100 isolates	[105,111,114,300,477–480,482–484]
Azole-resistant isolates	Determine nature and trends in Cyp51A mutation distribution	Cyp51A-gene mutation analysis	A	II	Test resistant isolates from surveillance survey	[107]

Abbreviations: AmB, Amphotericin B; MIC, minimum inhibitory concentration; QoE, Quality of evidence; SoR, Strength of recommendation.

ESCMID[†] and ECMM[‡] joint clinical guidelines for the diagnosis and management of mucormycosis 2013

O. A. Cornely^{1,†,‡,§}, S. Arikan-Akdagli^{2,†,§}, E. Dannaoui^{3,§}, A. H. Groll^{4,†,‡,§}, K. Lagrou^{5,†,§}, A. Chakrabarti^{6,§}, F. Lanternier^{7,§}, L. Pagano⁹, A. Skiada¹⁰, M. Akova², M. C. Arendrup¹¹, T. Boekhout^{12,13,14}, A. Chowdhary¹⁵, M. Cuenca-Estrella^{16,†,‡}, T. Freiburger^{17,18,†}, J. Guinea^{19,†,‡}, J. Guarro^{20,†}, S. de Hoog^{12,†}, W. Hope^{21,†}, E. Johnson^{22,†}, S. Kathuria¹⁵, M. Lackner^{23,†}, C. Lass-Flörl^{23,†,‡}, O. Lortholary^{7,†,‡}, J. F. Meis^{24,25,†,‡}, J. Meletiadis^{26,†}, P. Muñoz^{19,†}, M. Richardson^{27,28,†,‡}, E. Roilides^{29,†,‡}, A. M. Tortorano^{30,†}, A. J. Ullmann^{31,†,‡}, A. van Diepeningen¹², P. Verweij^{25,32,†,‡} and G. Petrikos^{33,†,‡,§}

TABLE 5. Recommendations on susceptibility testing in mucormycosis

Population	Intention	Method/Finding	SoR	QoE	Comment	References
Any	To guide treatment	EUCAST/CLSI reference microdilution methods	C	IIu	Clinical relevance uncertain. No data available to correlate MIC and outcome	79,80,83
Any	To guide treatment	Correlation of MIC with <i>in vivo</i> outcome	C	IIu	For <i>Apophysomyces elegans</i> , limited retrospective data suggest correlation	83
Any	To guide treatment	Correlation of MIC/MFC with <i>in vivo</i> outcome	B	III	Animal, posaconazole better in <i>Rhizopus microsporus</i> and <i>Rhizopus oryzae</i> strains MIC 0.25 µg/mL than in those with MICs 2 µg/mL	82,84,85
Any	To establish epidemiological knowledge	Susceptibility testing	A	IIu	<i>n</i> = 37 <i>n</i> = 36 <i>n</i> = 217 <i>n</i> = 45 <i>n</i> = 77 <i>n</i> = 18, <i>Apophysomyces elegans</i> <i>n</i> = 21 <i>n</i> = 66 Review	86 88 87 21 92 83 91 90 195
Any	To establish epidemiological knowledge	MIC determined by reference method	A	III	e.g. Etest [®] not validated for Mucorales	79,80

MFC, minimum fungicidal concentration; QoE, quality of evidence; SoR, strength of recommendation.

ESCMID and ECMM joint guidelines on diagnosis and management of hyalohyphomycosis: *Fusarium* spp., *Scedosporium* spp. and others

A. M. Tortorano^{1,*†}, M. Richardson^{2,3,*†,‡}, E. Roilides^{4,*†,‡}, A. van Diepeningen^{5,*}, M. Caira^{6,*}, P. Munoz^{7,*†}, E. Johnson^{8,*†}, J. Meletiadis^{9,*†}, Z.-D. Pana^{4,*}, M. Lackner^{10,*†}, P. Verweij^{11,12,*†,‡}, T. Freiburger^{13,*†}, O. A. Cornely^{14,†,‡}, S. Arikan-Akdogan^{15,†}, E. Dannaoui^{16,†}, A. H. Groll^{17,†,‡}, K. Lagrou^{18,†}, A. Chakrabarti¹⁹, F. Lanternier^{20,21}, L. Pagano^{22,†}, A. Skiada^{23,†}, M. Akova^{15,†}, M. C. Arendrup^{24,†,‡}, T. Boekhout^{5,25,26,†}, A. Chowdhary^{27,†}, M. Cuenca-Estrella^{28,†,‡}, J. Guinea^{7,†,‡}, J. Guarro^{29,†}, S. de Hoog^{5,†}, W. Hope^{30,†}, S. Kathuria²⁷, O. Lortholary^{31,32,†,‡}, J. F. Meis^{11,33,†,‡}, A. J. Ullmann^{34,†,‡}, G. Petrikos^{35,*†,‡} and C. Lass-Flörl^{10,*†,‡}

TABLE 4. Summary of recommendations for diagnosis of *Fusarium* infection

<i>Fusarium</i> infection/ Population	Test	SoR	QoE	Comment	References
Any population	Direct microscopy	A	III	Essential investigation	[2]
	Culture (species identification)	A	III	Essential investigation Easily recovered on routine mycological media without cycloheximide	[2,190]
	Histopathology	A	IIu	Essential investigation Features of hyaline septate hyphae (with acute angle branching) are similar to those seen with aspergillosis	[120]
	Immunohistochemistry	C	III	Not yet evaluated	[120]
	β-D-Glucan test/ Galactomannan	B	III	Glucan usually positive in case of invasive fusariosis. <i>Aspergillus</i> galactomannan sometimes positive in patients with fusariosis	[46]
	Pan-fungal PCRs for identification ^a	C	II	In combination with conventional methods High negative predictive values	[36–38]
	Multiplex PCRs ^a	C	III	Not yet validated Cover limited number of species/genera	[30,37,39]
	<i>In situ</i> hybridization	C	III	Not yet evaluated	[191,192]
	Susceptibility testing	C	III	Gives an overview of drug activity and may be helpful in selecting antifungals	[2,181,182,193–195]
	Environmental sampling (and fungal typing)	A	III	In case of an outbreak situation	[33,76]
Haematological patients	Chest computed tomography (CT) scan	A	IIu	None of patients had normal CT Pulmonary nodules in 82% of patients	[26]

QoE, quality of evidence; SoR, strength of recommendation.

^aThird-party appraisal of results and harmonization of PCR-based techniques are necessary before any clear recommendations can be made regarding clinical utility.

ESCMID and ECMM joint guidelines on diagnosis and management of hyalohyphomycosis: *Fusarium* spp., *Scedosporium* spp. and others

A. M. Tortorano^{1,*†}, M. Richardson^{2,3,*†,‡}, E. Roilides^{4,*†,‡}, A. van Diepeningen^{5,*}, M. Caira^{6,*}, P. Munoz^{7,*†}, E. Johnson^{8,*†}, J. Meletiadis^{9,*†}, Z.-D. Pana^{4,*}, M. Lackner^{10,*†}, P. Verweij^{11,12,*†,‡}, T. Freiburger^{13,*†}, O. A. Cornely^{14,†,‡}, S. Arikian-Akdogan^{15,†}, E. Dannaoui^{16,†}, A. H. Groll^{17,†,‡}, K. Lagrou^{18,†}, A. Chakrabarti¹⁹, F. Lanternier^{20,21}, L. Pagano^{22,†}, A. Skiada^{23,†}, M. Akova^{15,†}, M. C. Arendrup^{24,†,‡}, T. Boekhout^{5,25,26,†}, A. Chowdhary^{27,†}, M. Cuenca-Estrella^{28,†,‡}, J. Guinea^{7,†,‡}, J. Guarro^{29,†}, S. de Hoog^{5,†}, W. Hope^{30,†}, S. Kathuria²⁷, O. Lortholary^{31,32,†,‡}, J. F. Meis^{11,33,†,‡}, A. J. Ullmann^{34,†,‡}, G. Petrikos^{35,*†,‡} and C. Lass-Flörl^{10,*†,‡}

TABLE 7. Summary of recommendations for diagnosis of Scedosporium infections

Population	Test	SoR	QoE	Comment	References
Any population	Direct microscopy	A	III	Essential investigation	[85]
	Culture (species identification by morphology and physiological characteristics)	A	III	Essential investigation Selective media supplemented with cycloheximide or benomyl (10 mg/L, Sce ⁻ Sel ⁺) allows growth of <i>Scedosporium</i> over other filamentous fungi from bronchial secretions.	[85,122,123,136,206–210]
	Molecular-based identification methods	C	III	Accurate species assignment is important for guiding clinical management	[121–123,211]
	Histopathology	A	III	Hyaline thin-walled septate hyphae, 2–5 µm wide similar to those seen with aspergillosis and other hyalohyphomycoses	[120,212,213]
	Pan-fungal PCR ^a	C	III	Irregular branching	[36–38]
	Multiplex PCR ^a	C	III	Molecular tests could be used in combination with conventional laboratory tests.	[30,31,37,39]
	In situ hybridization	C	III	Molecular tests could be used in combination with conventional laboratory tests.	[191,192]
	Species identification (MALDI TOF and PCR)	C	III	Low sensitivity	[124,214–219]
	Physiological typing	C	III	Not yet validated	[220]
	In vitro susceptibility testing	C	III	In case of an outbreak situation	[221–223]
				Gives an overview of drug activity and therefore may support choice of antifungals	

MALDI-TOF, matrix-assisted laser desorption ionization time-of-flight mass spectrometry; QoE, quality of evidence; SoR, strength of recommendation.

^aThird-party appraisal of results and harmonization of PCR-based techniques are necessary before any clear recommendations can be made regarding clinical utility.

Klinik direnç sınır değerleri / "ECV" ("ECOFF")

GÜNCEL: Türe özgü
Yönteme özgü
İnkübasyon süresine özgü

Progress in Antifungal Susceptibility Testing of *Candida* spp. by Use of Clinical and Laboratory Standards Institute Broth Microdilution Methods, 2010 to 2012

M. A. Pfaller^{a,b} and D. J. Diekema^b

JMI Laboratories, North Liberty, Iowa, USA,^a University of Iowa Carver College of Medicine, Iowa City, Iowa, USA^b

Journal of Clinical Microbiology p. 2846–2856

September 2012 Volume 50 Number 9

Wild-Type MIC Distributions and Epidemiological Cutoff Values for Amphotericin B and *Aspergillus* spp. for the CLSI Broth Microdilution Method (M38-A2 Document)[∇]

A. Espinel-Ingroff,^{1*} M. Cuenca-Estrella,² A. Fothergill,³ J. Fuller,⁴ M. Ghannoum,⁵
E. Johnson,⁶ T. Pelaez,⁷ M. A. Pfaller,⁸ and J. Turnidge⁹

Wild-Type MIC Distributions and Epidemiological Cutoff Values for the Triazoles and Six Aspergillus spp. for the CLSI Broth Microdilution Method (M38-A2 Document)[∇]

A. Espinel-Ingroff,^{1*} D. J. Diekema,² A. Fothergill,³ E. Johnson,⁴ T. Pelaez,⁵
M. A. Pfaller,² M. G. Rinaldi,³ E. Canton,⁶ and J. Turnidge⁷

JOURNAL OF CLINICAL MICROBIOLOGY, Sept. 2010, p. 3251–3257

Vol. 48, No. 9

Multicenter Study of Isavuconazole MIC Distributions and Epidemiological Cutoff Values for Aspergillus spp. for the CLSI M38-A2 Broth Microdilution Method

A. Espinel-Ingroff,^a A. Chowdhary,^b G. M. Gonzalez,^c C. Lass-Flörl,^d E. Martin-Mazuelos,^e J. Meis,^{f,g} T. Peláez,^h M. A. Pfaller,ⁱ
J. Turnidge^j

August 2013 Volume 57 Number 8

Antimicrobial Agents and Chemotherapy p. 3823–3828

Multicenter Evaluation of MIC Distributions for Epidemiologic Cutoff Value Definition To Detect Amphotericin B, Posaconazole, and Itraconazole Resistance among the Most Clinically Relevant Species of Mucorales

A. Espinel-Ingroff,^a A. Chakrabarti,^b A. Chowdhary,^c S. Cordoba,^d E. Dannaoui,^e P. Dufresne,^f A. Fothergill,^g M. Ghannoum,^h
G. M. Gonzalez,ⁱ J. Guarro,^j S. Kidd,^k C. Lass-Flörl,^l J. F. Meis,^m T. Pelaez,ⁿ A. M. Tortorano,^o J. Turnidge^p

March 2015 Volume 59 Number 3

Antimicrobial Agents and Chemotherapy

International Evaluation of MIC Distributions and Epidemiological Cutoff Value (ECV) Definitions for *Fusarium* Species Identified by Molecular Methods for the CLSI Broth Microdilution Method

A. Espinel-Ingroff,^a A. L. Colombo,^b S. Cordoba,^c P. J. Dufresne,^d J. Fuller,^e M. Ghannoum,^f G. M. Gonzalez,^g J. Guarro,^h S. E. Kidd,ⁱ  J. F. Meis,^j T. M. S. C. Melhem,^k T. Pelaez,^l M. A. Pfaller,^m M. W. Szesz,ⁿ J. P. Takahaschi,^o  A. M. Tortorano,^p N. P. Wiederhold,^q J. Turnidge^r

February 2016 Volume 60 Number 2

1079–1084.

Antimicrobial Agents and Chemotherapy

CLSI M59

Epidemiological Cutoff Values for Antifungal Susceptibility Testing, 2nd ed.

This document includes epidemiological cutoff values and quality control tables developed according to criteria provided in the Clinical and Laboratory Standards Institute guideline M57.

AMB, Azoller, Kandin- Candida

EUCAST

Candida spp.

EUCAST Antifungal Clinical Breakpoint Table v. 9.0 valid from 2018-02-12

MIC method (EUCAST standardised broth microdilution method)

Medium: RPMI1640-2% glucose, MOPS buffer

Inoculum: Final 0.5×10^5 – 2.5×10^5 cfu/mL

Incubation: 18-24h

Reading: Spectrophotometric, complete (>90%) inhibition for amphotericin B but 50% growth inhibition for other compounds

Quality control: *C. parapsilosis* ATCC 22019 or *C. krusei* ATCC 6258

Antifungal agent	MIC breakpoint (mg/L)															
	<i>C. albicans</i>		<i>C. dubliniensis</i>		<i>C. glabrata</i>		<i>C. krusei</i>		<i>C. parapsilosis</i>		<i>C. tropicalis</i>		<i>C. guilliermondii</i>		Non-species related breakpoints ¹	
	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >
Amphotericin B	1	1	IE	IE	1	1	1	1	1	1	1	1	IE	IE	IE	IE
Anidulafungin	0.032	0.032	IE	IE	0.064	0.064	0.064	0.064	0.002	4	0.064	0.064	IE ²	IE ²	IE	IE
Caspofungin	Note ³	Note ³	IE	IE	Note ³	Note ³	Note ³	Note ³	Note ³	Note ³	Note ³	Note ³	IE ²	IE ²	IE	IE
Fluconazole	2	4	IE	IE	0.002	32	-	-	2	4	2	4	IE ²	IE ²	2	4
Isavuconazole	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE
Itraconazole	0.064	0.064	0.064	0.064	IE ²	IE ²	IE ²	IE ²	0.125	0.125	0.125	0.125	IE ²	IE ²	IE	IE
Micafungin	0.016	0.016	IE	IE	0.032	0.032	IE ⁴	IE ⁴	0.002	2	IE ⁴	IE ⁴	IE ⁴	IE ⁴	IE	IE
Posaconazole	0.064	0.064	0.064	0.064	IE ²	IE ²	IE ²	IE ²	0.064	0.064	0.064	0.064	IE ²	IE ²	IE	IE
Voriconazole ⁶	0.064 ⁵	0.25 ⁵	0.064	0.25	IE	IE	IE	IE	0.125 ⁵	0.25 ⁵	0.125 ⁵	0.25 ⁵	IE ²	IE ²	IE	IE

AMB & Azoles -Asp

EUCAST

Aspergillus spp.

EUCAST Antifungal Clinical Breakpoint Table v. 9.0 valid from 2018-02-12

MIC method (EUCAST standardised broth microdilution method)

Medium: RPMI1640-2% glucose, MOPS as buffer

Inoculum: Final 1×10^5 – 2.5×10^5 cfu/mL

Incubation: 48h

Reading: Visual, complete inhibition for amphotericin B and azoles (MIC), aberrant growth endpoint for echinocandins (MEC).

Quality control: *A. fumigatus* ATCC 204305, *A. flavus* ATCC 204304, *A. fumigatus* F 6919, *A. flavus* CM 1813, *C. parapsilosis* ATCC 22019 (read after 18-24 h) or *C. krusei* ATCC 6258 (read after 18-24 h).

Antifungal agent	MIC breakpoint (mg/L)											
	<i>A. flavus</i>		<i>A. fumigatus</i>		<i>A. nidulans</i>		<i>A. niger</i>		<i>A. terreus</i>		Non-species related breakpoints ¹	
	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >	S ≤	R >
Amphotericin B	IE ²	IE ²	1	2	Note ³	Note ³	1	2	-	-	IE	IE
Anidulafungin	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE
Caspofungin	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE
Fluconazole	-	-	-	-	-	-	-	-	-	-	-	-
Isavuconazole	IE ²	IE ²	1	1	0.25	0.25	IE ²	IE ²	1	1	IE	IE
Itraconazole ⁴	1	2	1	2	1	2	IE ^{2,5}	IE ^{2,5}	1	2	IE ⁵	IE ⁵
Micafungin	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE	IE
Posaconazole ⁴	IE ²	IE ²	0.125 ⁶	0.25 ⁶	IE ²	IE ²	IE ²	IE ²	0.125 ⁶	0.25 ⁶	IE	IE
Voriconazole ⁴	IE ²	IE ²	1	2	IE	IE	IE ²	IE ²	IE ²	IE ²	IE	IE

Sunum Planı

Temel bilgiler



YÖNTEM ve
DEĞERLENDİRME



DİRENÇ VARLIĞI /
ORANI



ÖNERİLER

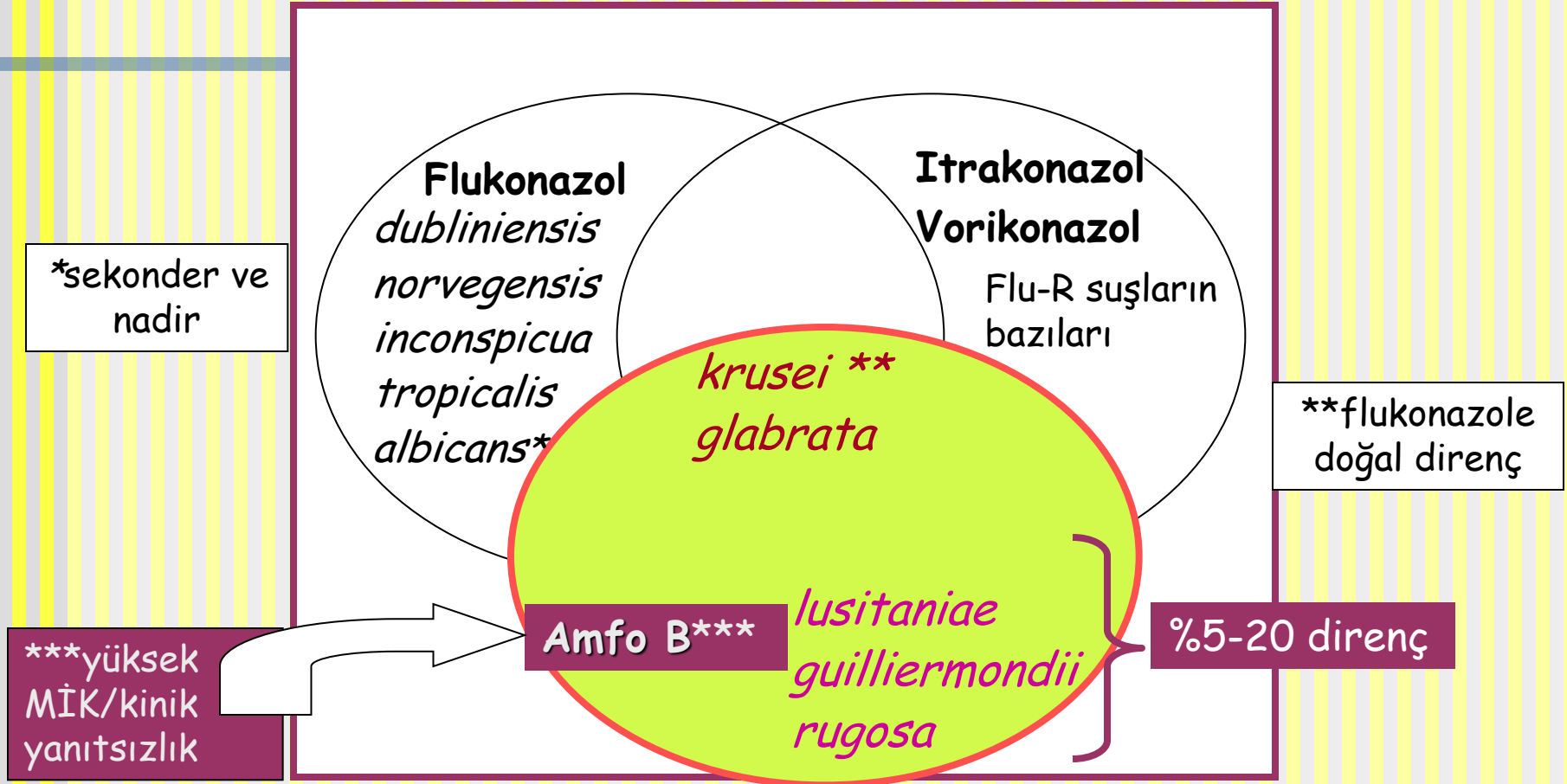
A microscopic image showing a dense population of Candida cells. The cells are small, oval-shaped, and appear to be budding or forming chains. They are stained a deep purple or magenta color, contrasting with the lighter, yellowish-orange background of the tissue or medium. The overall texture is granular and somewhat mottled.

Direnç

VE

Candida

AMB, Azol - Candida



CD101, a long-acting echinocandin, and comparator antifungal agents tested against a global collection of invasive fungal isolates in the SENTRY 2015 Antifungal Surveillance Program

Michael A. Pfaller ^{a,b}, Shawn A. Messer ^a, Paul R. Rhomberg ^a, Mariana Castanheira ^{a,*}

International Journal of Antimicrobial Agents 50 (2017) 352–358

589 *Candida* suşu
C. glabrata, **Flukonazol** direnci: %6.6
C. parapsilosis, **Flukonazol** direnci: % 3.6
C. glabrata, **Kandin** direnci: %0.8

Table 3

FKS alterations detected in *Candida* spp. isolates displaying non-wild-type echinocandin MIC values.

State, Country	Organism	MIC according to CLSI method (µg/mL):				1,3-β-D-glucan synthase mutations:			
		CD101	Anidulafungin	Caspofungin	Micafungin	<i>fks1</i> HS1	<i>fks1</i> HS2	<i>fks2</i> HS1	<i>fks2</i> HS2
NY, USA	<i>Candida albicans</i>	0.25	0.25	1	1	S645P	WT	NT	NT
TX, USA	<i>Candida glabrata</i>	0.25	0.25	0.06	0.03	WT	WT	WT	WT
CA, USA	<i>Candida glabrata</i>	1	1	1	0.25	F625S	NT	WT	WT
VA, USA	<i>Candida glabrata</i>	0.12	0.12	0.06	0.12	WT	WT	WT	WT

HS=hot spot; NT=not tested; WT=wild type.

C. albicans n=304
C. glabrata n=121
C. parapsilosis n=83
C. tropicalis n=55

C. glabrata: Ekinokandin direnci

Oranlar; *Coğrafi Farklılıklar*

A.B.D.

Son 10 yılda %2 den > %13'e (A.B.D.)

Alexander B et al. CID 2013; 56: 1724

%3.1-%3.6 (CDC) A.B.D.-4 şehir

58: 4690

Avrupa

< %1

MSH2 DNA «mismatch
repair» (MMR) gen
delesyonları sonucu
ortaya çıkan "mutator
fenotip" → Flu,
Ekino, AMB direnci

16; 59:312

Ekinokandin

Ekinokandin direnci suşlarının %36'
sında

Pham et al. AAC 2014; 58: 4690

Değişen sınır değerler ile değişen duyarlılık kategorileri (CLSI MD ve DD ile)

Klinik *Candida glabrata* İzolatlarında Flukonazol ve Vorikonazol Duyarlılığının Saptanmasında Mikrodilüsyon ve Disk Difüzyon Yöntemlerinin Karşılaştırılması ve Yeni CLSI Direnç Sınır Değerleri ile Duyarlılık Sonuçlarındaki Değişimin Belirlenmesi

Gülşen HAZIROLAN¹, Zeynep SARIBAŞ², Sevtap ARIKAN AKDAĞLI²

Tablo IV. Önceki ve yeni CLSI direnç sınır değerlerine göre flukonazol mikrodilüsyon ve disk difüzyon test sonuçları (%) (n= 70)

MİKRODİLÜSYON (24 saat)					DİSK DİFÜZYON (24 saat)				
		Önceki CLSI					Önceki CLSI		
		H	DBD	D			H	DBD	D
Yeni CLSI	DBD	94.3	5.7	-	Yeni CLSI	DBD	81.4	5.7	-
	D	-	-	-		D	-	-	12.9
MİKRODİLÜSYON (48 saat)					DİSK DİFÜZYON (48 saat)				
		Önceki CLSI					Önceki CLSI		
		H	DBD	D			H	DBD	D
Yeni CLSI	DBD	80	15.7	-	Yeni CLSI	DBD	68.6	15.7	-
	D	-	-	4.3		D	-	-	15.7
H: Duyarlı; DBD: Doza bağlı duyarlı; D: Dirençli.									

Table. The number of *Candida* isolates with available AST results interpreted according to CBPs^{3,4} or ECVs⁴.

In presence of established CBPs, the isolates were categorized as susceptible (S), dose-dependent susceptible (S-DD) or resistant (R). For the species with no established CBPs and available ECVs for the drugs under study, the categories were determined as wild-type (WT) or non-wild-type (non-WT). The number of isolates determined by using ECVs are shaded in blue.

No. of <i>Candida</i> isolates tested	No. of isolates in each interpretative category (Fluconazole)			No. of <i>Candida</i> isolates tested	No. of isolates in each interpretative category (Voriconazole)			No. of <i>Candida</i> isolates tested	No. of isolates in each interpretative category (Echinocandin)		
	S / WT	S-DD	R / Non-WT		S / WT	S-DD	R / Non-WT		S / WT	I	R / Non-WT
<i>C.albicans</i> (130)	130	-	-	<i>C.albicans</i> (129)	129	-	-	<i>C.albicans</i> (109)	108	-	1
<i>C.parapsilosis</i> (46)	32	1	13	<i>C.parapsilosis</i> (45)	42	3	-	<i>C.parapsilosis</i> (39)	39	-	-
<i>C.glabrata</i> (33)	-	32	1	<i>C.glabrata</i> (33)	32	-	1	<i>C.glabrata</i> (29)	29	-	-
<i>C.tropicalis</i> (23)	23	-	-	<i>C.tropicalis</i> (24)	24	-	-	<i>C.tropicalis</i> (21)	21	-	-
<i>C.kefyr</i> (7)	7	-	-	<i>C.kefyr</i> (7)	7	-	-	<i>C.kefyr</i> (6)	5	-	1
<i>C.krusei</i> (4)	-	-	4	<i>C.krusei</i> (2)	2	-	-	<i>C.krusei</i> (2)	2	-	-
<i>C.lusitanae</i> (4)	4	-	-	<i>C.lusitanae</i> (4)	4	-	-	<i>C.lusitanae</i> (4)	4	-	-
<i>C.dubliniensis</i> (3)	3	-	-	<i>C.dubliniensis</i> (3)	3	-	-	<i>C.dubliniensis</i> (2)	2	-	-
<i>C.guilliermondii</i> (2)	2	-	-	<i>C.guilliermondii</i> (2)	2	-	-	<i>C.guilliermondii</i> (1)	1	-	-
<i>C.pelliculosa</i> (1)	-	-	1	<i>C.pelliculosa</i> (1)	1	-	-	<i>C.pelliculosa</i> (1)	1	-	-
<i>C.rugosa</i> * (1)	-	-	-	<i>C.rugosa</i> * (1)	-	-	-	<i>C.rugosa</i> * (1)	-	-	-

Yeni sınır değerler ile duyarlılık kategorileri

***Candida* Türlerinin Triazol Antifungal Duyarlılık
Profilleri: Antifungal Direncin Belirlenmesinde Yeni
CLSI Türe Özgü Klinik Direnç Sınır Değerleri ve
Epidemiyolojik Eşik Değerlerinin Uygulanması***

.....

Buna göre, 5 *Candida* izolatı (3 *C.albicans*, 1 *C.parapsilosis* ve 1 *C.glabrata*) flukonazole dirençli; birer *C.lusitaniae*, *C.dubliniensis* ve *C.kefyr* izolatı ise vahşi olmayan tip olarak saptanmıştır. Ayrıca 4 *Candida* izolatı (2 *C.albicans*, ve 2 *C.tropicalis*) vorikonazole dirençli ve 9 *Candida* izolatı (5 *C.glabrata*, 3 *C.lusitaniae*, ve 1 *C.kefyr*) vahşi olmayan tip olarak değerlendirilmiştir.

12 merkez

n= 1992

1996-2017

Hacettepe Üniv. - Mikoloji Laboratuvarı
CLSI referans mikrodilüsyon

Kandidemi - Çok Merkezli
Antifungal Direnç

Hacettepe Üniv, Dokuz Eylül Üniv, Ege Üniv, Erciyes Üniv., Gazi Üniv., İstanbul Üniv. İstanbul Tıp Fak.,
Marmara Üniv., Ondokuz Mayıs Üniv., Osmangazi Üniv., Sağlık Bilimleri Üniv. Gülhane Tıp Fak., Selçuk
Üniv, Uludağ Üniv.

	n
C.albicans	852
C.parapsilosis	575
C.glabrata	216
C.tropicalis	203
C.krusei	52
C.kefyr	33
C.lusitaniae	23
C.guilliermondii	16
C.inconspicua/norvegensis	7
C.dublinsiensis	6
C.pelliculosa	3
C.rugosa	2
C.utilis	2
C.lipolytica	1
C.sake	1

1992

Arıkan Akdağlı S
Yayınlanmamış Veri

12 merkez

n= 1992

1996-2017

Hacettepe Üniv. - Mikoloji Laboratuvarı

CLSI referans mikrodilüsyon

Kandidemi - Çok Merkezli

Antifungal Direnç

Resistance %, n

CLSI M27-S4

Pfaller
2012

Fluconazole

%

n

	S	SDD	R	WT	non-WT	n	S	SDD	R	WT	non-WT
C.albicans	99.8	0.2	0.0	96.6	3.4	852	850	2	0	823	29
C.parapsilosis	89.0	3.3	7.7	89.0	11.0	575	512	19	44	512	63
C.glabrata	0.0	99.1	0.9	99.1	0.9	216	0	214	2	214	2
C.tropicalis	100.0	0.0	0.0	100.0	0.0	203	203	0	0	203	0
C.krusei	0.0	0.0	100.0	94.2	5.8	52	0	0	52	49	3
C.kefyr	-	-	-	100.0	0.0	33	-	-	-	33	0
C.lusitaniae	-	-	-	95.7	4.3	23	-	-	-	22	1
C.guilliermondii	-	-	-	100.0	0.0	16	-	-	-	16	0
C.inconspicua/norvegensis	-	-	-	-	-	7	-	-	-	-	-
C.dublinsiensis	-	-	-	100.0	0.0	6	-	-	-	6	0
C.pelliculosa	-	-	-	100.0	0.0	3	-	-	-	3	0
C.rugosa	-	-	-	-	-	2	-	-	-	-	-
C.utilis	-	-	-	-	-	2	-	-	-	-	-
C.lipolytica	-	-	-	-	-	1	-	-	-	-	-
C.sake	-	-	-	-	-	1	-	-	-	-	-

1992

Arıkan Akdağlı S. Yayınlanmamış Veri

12 merkez

n= 1992

1996-2017

Hacettepe Üniv. - Mikoloji Laboratuvarı
CLSI referans mikrodilüsyon

Kandidemi - Çok Merkezli
Antifungal Direnç

Resistance %, n

Voriconazole	%					n					
	S	SDD	R	WT	non-WT	n	S	SDD	R	WT	non-WT
C.albicans	100.0	0.0	0.0	99.9	0.1	852	852	0	0	851	1
C.parapsilosis	97.9	2.1	0.0	97.9	2.1	575	563	12	0	563	12
C.glabrata	-	-	-	99.5	0.5	216	-	-	-	215	1
C.tropicalis	100.0	0.0	0.0	100.0	0.0	203	203	0	0	203	0
C.krusei	100.0	0.0	0.0	100.0	0.0	52	52	0	0	52	0
C.kefyr	-	-	-	100.0	0.0	33	-	-	-	33	0
C.lusitaniae	-	-	-	100.0	0.0	23	-	-	-	23	0
C.guilliermondii	-	-	-	100.0	0.0	16	-	-	-	16	0
C.inconspicua/norvegensis	-	-	-	-	-	7	-	-	-	-	-
C.dublinsiensis	-	-	-	100.0	0.0	6	-	-	-	6	0
C.pelliculosa	-	-	-	100.0	0.0	3	-	-	-	3	0
C.rugosa	-	-	-	-	-	2	-	-	-	-	-
C.utilis	-	-	-	-	-	2	-	-	-	-	-
C.lipolytica	-	-	-	-	-	1	-	-	-	-	-
C.sake	-	-	-	-	-	1	-	-	-	-	-

Resistance %, n

Itraconazole	%						n				
	S	SDD	R	WT	non-WT	n	S	SDD	R	WT	non-WT
C.albicans	99.4	0.6	0.0	99.4	0.6	852	847	5	0	847	5
C.parapsilosis	-	-	-	99.8	0.2	575	-	-	-	574	1
C.glabrata	-	-	-	100.0	0.0	216	-	-	-	216	0
C.tropicalis	-	-	-	100.0	0.0	203	-	-	-	203	0
C.krusei	-	-	-	100.0	0.0	52	-	-	-	52	0
C.kefyr	-	-	-	-	-	33	-	-	-	-	-
C.lusitaniae	-	-	-	100.0	0.0	23	-	-	-	23	0
C.guilliermondii	-	-	-	100.0	0.0	16	-	-	-	16	0
C.inconspicua/norvegensis	-	-	-	-	-	7	-	-	-	-	-
C.dublinsiensis	-	-	-	100.0	0.0	6	-	-	-	6	0
C.pelliculosa	-	-	-	-	-	3	-	-	-	-	-
C.rugosa	-	-	-	-	-	2	-	-	-	-	-
C.utilis	-	-	-	-	-	2	-	-	-	-	-
C.lipolytica	-	-	-	-	-	1	-	-	-	-	-
C.sake	-	-	-	-	-	1	-	-	-	-	-
						1992					

12 merkez
n= 1992
1996-2017
Hacettepe Üniv. - Mikoloji Laboratuvarı
CLSI referans mikrodilüsyon

Kandidemi - Çok Merkezli
Antifungal Direnç

Resistance %, n

Posaconazole	%						n					
	S	SDD	R	WT	non-WT	n	S	SDD	R	WT	non-WT	
C.albicans	-	-	-	100.0	0.0	852	-	-	-	852	0	
C.parapsilosis	-	-	-	96.5	3.5	575	-	-	-	555	20	
C.glabrata	-	-	-	100.0	0.0	216	-	-	-	216	0	
C.tropicalis	-	-	-	100.0	0.0	203	-	-	-	203	0	
C.krusei	-	-	-	98.1	1.9	52	-	-	-	51	1	
C.kefyr	-	-	-	100.0	0.0	33	-	-	-	33	0	
C.lusitaniae	-	-	-	100.0	0.0	23	-	-	-	23	0	
C.guilliermondii	-	-	-	100.0	0.0	16	-	-	-	16	0	
C.inconspicua/norvegensis	-	-	-	-	-	7	-	-	-	-	-	
C.dublinsiensis	-	-	-	100.0	0.0	6	-	-	-	6	0	
C.pelliculosa	-	-	-	100.0	0.0	3	-	-	-	3	0	
C.rugosa	-	-	-	-	-	2	-	-	-	-	-	
C.utilis	-	-	-	-	-	2	-	-	-	-	-	
C.lipolytica	-	-	-	-	-	1	-	-	-	-	-	
C.sake	-	-	-	-	-	1	-	-	-	-	-	
						1992						

12 merkez
n= 1992
1996-2017
Hacettepe Üniv. - Mikoloji Laboratuvarı
CLSI referans mikrodilüsyon

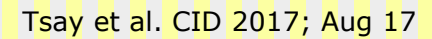
Kandidemi - Çok Merkezli
Antifungal Direnç

Resistance %, n

Micafungin	%						n				
	S	SDD	R	WT	non-WT	n	S	I	R	WT	non-WT
C.albicans	100.0	0.0	0.0	99.6	0.4	852	852	0	0	849	3
C.parapsilosis	99.8	0.2	0.0	100.0	0.0	575	574	1	0	575	0
C.glabrata	100.0	0.0	0.0	99.1	0.9	216	216	0	0	214	2
C.tropicalis	100.0	0.0	0.0	100.0	0.0	203	203	0	0	203	0
C.krusei	100.0	0.0	0.0	76.9	23.1	52	52	0	0	40	12
C.kefyr	-	-	-	100.0	0.0	33	-	-	-	33	0
C.lusitaniae	-	-	-	100.0	0.0	23	-	-	-	23	0
C.guilliermondii	100.0	0.0	0.0	100.0	0.0	16	16	0	0	16	0
C.inconspicua/norvegensis	-	-	-	-	-	7	-	-	-	-	-
C.dubliniensis	-	-	-	100.0	0.0	6	-	-	-	6	0
C.pelliculosa	-	-	-	-	-	3	-	-	-	-	-
C.rugosa	-	-	-	-	-	2	-	-	-	-	-
C.utilis	-	-	-	-	-	2	-	-	-	-	-
C.lipolytica	-	-	-	-	-	1	-	-	-	-	-
C.sake	-	-	-	-	-	1	-	-	-	-	-
						1992					

Resistance %, n

Amphotericin B	%						n				
	S	SDD	R	WT	non-WT	n	S	SDD	R	WT	non-WT
C.albicans	-	-	-	100.0	0.0	852	-	-	-	852	0
C.parapsilosis	-	-	-	100.0	0.0	575	-	-	-	575	0
C.glabrata	-	-	-	100.0	0.0	216	-	-	-	216	0
C.tropicalis	-	-	-	100.0	0.0	203	-	-	-	203	0
C.krusei	-	-	-	100.0	0.0	52	-	-	-	52	0
C.kefyr	-	-	-	-	-	33	-	-	-	-	-
C.lusitaniae	-	-	-	100.0	0.0	23	-	-	-	23	0
C.guilliermondii	-	-	-	100.0	0.0	16	-	-	-	16	0
C.inconspicua/norvegensis	-	-	-	-	-	7	-	-	-	-	-
C.dublinsiensis	-	-	-	100.0	0.0	6	-	-	-	6	0
C.pelliculosa	-	-	-	-	-	3	-	-	-	-	-
C.rugosa	-	-	-	-	-	2	-	-	-	-	-
C.utilis	-	-	-	-	-	2	-	-	-	-	-
C.lipolytica	-	-	-	-	-	1	-	-	-	-	-
C.sake	-	-	-	-	-	1	-	-	-	-	-
						1992					



www.cdc.gov/fungal/diseases/candidiasis/c-auris-drug-resistant.html

Simultaneous Emergence of Multidrug-Resistant *Candida auris* on 3 Continents Confirmed by Whole-Genome Sequencing and Epidemiological Analyses

Shawn R. Lockhart,¹ Kizee A. Etienne,¹ Snigdha Vallabhaneni,¹ Joveria Farooqi,⁴ Anuradha Chowdhary,⁶ Nelesh P. Govender,⁷ Arnaldo Lopes Colombo,⁸ Belinda Calvo,⁹ Christina A. Cuomo,² Christopher A. Desjardins,² Elizabeth L. Berkow,¹ Mariana Castanheira,³ Rindizani E. Magobo,⁷ Kauser Jabeen,⁴ Rana J. Asghar,⁵ Jacques F. Meis,^{10,11} Brendan Jackson,¹ Tom Chiller,¹ and Anastasia P. Litvintseva¹

CID 2017:64 (15 January)

Methods. To understand the global emergence and epidemiology of *C. auris*, we obtained isolates from 54 patients with *C. auris* infection from Pakistan, India, South Africa, and Venezuela during 2012–2015 and the type specimen from Japan. Patient information was available for 41 of the isolates. We conducted antifungal susceptibility testing and whole-genome sequencing (WGS).

Results. Available clinical information revealed that 41% of patients had diabetes mellitus, 51% had undergone recent surgery, 73% had a central venous catheter, and 41% were receiving systemic antifungal therapy when *C. auris* was isolated. The median time from admission to infection was 19 days (interquartile range, 9–36 days), 61% of patients had bloodstream infection, and 59% died. Using stringent break points, 93% of isolates were resistant to fluconazole, 35% to amphotericin B, and 7% to echinocandins; 41% were resistant to 2 antifungal classes and 4% were resistant to 3 classes. WGS demonstrated that isolates were grouped into unique clades by geographic region. Clades were separated by thousands of single-nucleotide polymorphisms, but within each clade isolates were clonal. Different mutations in *ERG11* were associated with azole resistance in each geographic clade.

Conclusions. *C. auris* is an emerging healthcare-associated pathogen associated with high mortality. Treatment options are limited, due to antifungal resistance. WGS analysis suggests nearly simultaneous, and recent, independent emergence of different clonal populations on 3 continents. Risk factors and transmission mechanisms need to be elucidated to guide control measures.

Why is *Candida auris* a problem?



It causes serious infections. *C. auris* can cause bloodstream infections and even death, particularly in hospital and nursing home patients with serious medical problems. More than 1 in 3 patients with invasive *C. auris* infection (for example, an infection that affects the blood, heart, or brain) die.



It's often resistant to medicines. Antifungal medicines commonly used to treat *Candida* infections often don't work for *Candida auris*. Some *C. auris* infections have been resistant to all three types of antifungal medicines.



It's becoming more common. Although *C. auris* was just discovered in 2009, it has spread quickly and caused infections in more than a dozen countries.



It's difficult to identify. *C. auris* can be misidentified as other types of fungi unless specialized laboratory technology is used. This misidentification might lead to a patient getting the wrong treatment.



It can spread in hospitals and nursing homes. *C. auris* has caused outbreaks in healthcare facilities and can spread through contact with affected patients and contaminated surfaces or equipment. Good hand hygiene and cleaning in healthcare facilities is important because *C. auris* can live on surfaces for several weeks.

Scientists are still learning about *Candida auris*

CDC and public health partners are working hard to better understand *C. auris* and answer the following questions so that we can continue to help protect people from this serious infection:

- Why is *C. auris* resistant to antifungal medicines?
- Why did *C. auris* start causing infections in recent years?
- Where did *C. auris* originally come from, and why has it appeared in many regions of the world at the same time?

What is CDC doing?

CDC is collaborating closely with partners to better respond, contain spread, and prevent future infections by:

- Advising healthcare workers and infection control staff on ways to stop the spread of *C. auris* and continually updating this guidance as we learn more about the infection.
- Working with state and local health agencies, healthcare facilities, and clinical microbiology laboratories to ensure that laboratories are using proper methods to detect *C. auris*.
- Testing *C. auris* strains to monitor for resistance to antifungal medicines.
- Examining the DNA of *C. auris* strains using whole genome sequencing to better understand how this germ is spreading in the United States and around the world.
- Working with public health partners in the United States and internationally to learn more about how *C. auris* spreads in healthcare facilities and to eliminate it from those facilities.



"News from ESCMID-EFISG": *Candida auris* (January 2017)

- "... Due to the difficulty of the correct identification of *C. auris*, if proper identification methods are not available, we recommend the referral of suspected invasive isolates to a reference mycology laboratory.
- For more information you can visit the following links:
- · Risk Assessment of the European Center for Disease Control (ECDC)
- · Guidance for the laboratory investigation, management and infection prevention and control from cases of *Candida auris* elaborated by Public Health England (PHE)
- · CDC *Candida auris* [website](#) with links to [Interim Recommendations](#), as well as links to papers on a global [WGS analysis](#) and investigation of the first seven US [cases](#)

This Newsletter is issued on behalf of EFISG by the ESCMID Executive Office. It contains announcements of EFISG-related matters and other information of interest to professionals in the infection field.

Candida auris

- Yıl 2009, Japonya (Avrupa: İngiltere, 2015)
- Çok ilaca dirençli
- Yüksek geçiş oranı, salgınlar
- İnvazif, (yara, kulak) enf.
- Tanımlama: MALDI-TOF / rDNA D1-D2 veya ITS sekans analizi
- Biyokimyasal stripler/VITEK-2:
C. haemulonii, C. famata, C. sake, S. cerevisiae, R. glutinis

Simultaneous Emergence of Multidrug-Resistant *Candida auris* on 3 Continents Confirmed by Whole-Genome Sequencing and Epidemiological Analyses

Clinical Infectious Diseases® 2017;64(2):134–40

Lockhart et al.

Table 2. Antifungal Susceptibility Data for 54 *Candida auris* Isolates

Antifungal	MIC Range, µg/mL	MIC ₅₀ , µg/mL	MIC ₉₀ , µg/mL
Fluconazole	4–256	128	256
Voriconazole	0.03–16	2	8
Itraconazole	0.125–2	0.5	1
Posaconazole	0.06–1	0.5	1
Caspofungin	0.03–16	0.25	1
Anidulafungin	0.125–16	0.5	1
Micafungin	0.06–4	0.25	2
Flucytosine	0.125–128	0.125	0.5
Amphotericin B	0.38–4	1	2

Abbreviations: MIC, minimum inhibitory concentration; MIC₅₀, MIC for 50% of isolates; MIC₉₀, MIC for 90% of isolates.

Antifungal Susceptibility Testing

Antifungal susceptibility testing was performed on 54 isolates. The MIC range and the MICs for 50% and 90% of isolates are shown in Table 2, and the MIC distribution is shown in Supplemental Table 2. Using stringent break points, 50 isolates (93%) were resistant to fluconazole, 29 (54%) to voriconazole (≥ 2 µg/mL), 19 (35%) to amphotericin B (7 from Pakistan and 12 from India), 4 (7%) to echinocandins (2 from India and 2 from South Africa), and 3 (6%) (from India) were resistant to flucytosine. Two isolates, both from India, were resistant to fluconazole, voriconazole, echinocandins, and amphotericin B. In all, 22 (41%) isolates were resistant to ≥ 2 classes of antifungals.

4 filogenik grup (Coğrafi):

Doğu Asya

Güney Asya

Afrika

Güney Amerika

Comparison of EUCAST and CLSI Reference Microdilution MICs of Eight Antifungal Compounds for *Candida auris* and Associated Tentative Epidemiological Cutoff Values

M. C. Arendrup,^{a,b,c} Anupam Prakash,^d Joseph Meletiadis,^{e,f} Cheshta Sharma,^d
 Anuradha Chowdhary^d

99% endpoints), and via the derivatization method (dECOFFs). The CLSI and EUCAST MIC distributions were wide, with several peaks for all compounds except amphotericin B, suggesting possible acquired resistance. Modal MIC, geometric MIC, MIC₅₀, and MIC₉₀ values were ≤ 1 2-fold dilutions apart, and no significant differences were found. The quantitative agreement was best for amphotericin B (80%/97% within $\pm 1/\pm 2$ dilutions) and lowest for isavuconazole and anidulafungin (58%/76% to 75% within $\pm 1/\pm 2$ dilutions). We found that 90.2%/100% of the isolates were amphotericin B susceptible based on CLSI/EUCAST methods, respectively (i.e., with MICs of ≤ 1 mg/liter), and 100%/97.6% were fluconazole nonsusceptible by CLSI/EUCAST (MICs > 2). The ECOFFs (in milligrams per liter) were similar across the three different methods for itraconazole (ranges for CLSI/EUCAST, 0.25 to 0.5/0.5 to 1), posaconazole (0.125/0.125 to 0.25), amphotericin B (0.25 to 0.5/1 to 2), micafungin (0.25 to 0.5), and anidulafungin (0.25 to 0.5/0.25 to 1). In contrast, the estimated ECOFFs were dependent on the method applied for voriconazole (1 to 32) and isavuconazole (0.125 to 4). CLSI and EUCAST MICs were remarkably similar and confirmed uniform fluconazole resistance and variable acquired resistance to the other agents.

TABLE 1 MIC distributions of antifungal drugs for *C. auris* isolates (*n* = 123) tested by using the CLSI and EUCAST methods

Drug and AFST method	MIC (mg/liter) ^a															MIC range (no. of dilutions ^b)	GM	MIC ₅₀	MIC ₉₀
	0.002	0.004	0.008	0.016	0.032	0.064	0.125	0.25	0.5	1	2	4	8	16	32				
FLU																			
CLSI											-	10	7	15	(91)	4 to ≥64 (5)	43.38	≥64	≥64
EUCAST	NT ^c	NT	NT	NT					1		2		2	10	(108)	0.5 to ≥64 (8)	53.74	≥64	≥64
ITC																			
CLSI					25	12	57	20	7	1	1					0.032 to 2 (7)	0.11	0.125	0.25
EUCAST	NT	NT	(4)	9	4	14	36	35	20	1						≤0.008 to 1 (8)	0.13	0.125	0.5
VRC																			
CLSI					1	11	14	10	27	19	24	13	1	3		0.032 to 16 (10)	0.66	0.5	4
EUCAST	NT	NT	(1)		2	1	17	12	35	37	13	5				≤0.008 to 4 (10)	0.54	0.5	2
ISA																			
CLSI				26	7	23	20	30	12		3	2				0.015 to 4 (9)	0.095	0.125	0.5
EUCAST	NT	NT	(22)	1	19	9	20	20	21	6	5					≤0.008 to 2 (9)	0.090	0.125	0.5
PSC																			
CLSI				73	4	26	9	7		1	1	1	1			0.015 to 8 (9)	0.035	0.016	0.125
EUCAST	NT	NT	(22)	19	33	33	12	3	1							≤0.008 to 0.5 (7)	0.033	0.032	0.125
AMB																			
CLSI							2	16	58	35	4	6	2			0.125 to 8 (7)	0.66	0.5	2
EUCAST	NT	NT						1	15	107						0.25 to 1 (3)	0.91	1	1
AFG																			
CLSI				1		8	61	24	20	2		-	7			0.015 to 8 (10)	0.22	0.125	0.5
EUCAST	1			2	11	34	30	12	12	11	2	8				0.002 to 2 (12)	0.17	0.125	1
MFG																			
CLSI				4	4	47	49	9	2	1			7			0.015 to 8 (10)	0.12	0.125	0.25
EUCAST	1			1	5	29	69	9				8				0.002 to 4 (12)	0.13	0.125	0.25

^aModal MICs are indicated with underlined numbers and gray shading, and values in parentheses represent the number of isolates with an MIC equal or less than the MIC indicated due to truncation. Additional peaks are illustrated by underlining.

^bThe number of dilutions each MIC distribution spanned is given in parentheses.

^cNT, not tested.

TABLE 3 CLSI and EUCAST tentative statistical, derivatization, and visual ECOFFs for *Candida auris*, using three different endpoints for the statistical methods

		Statistical ECOFF at indicated endpoint ^a						dECOFF via derivatization method	ECOFF via visual eyeball method ^b
		95%		97.5%		99%			
Drug and AFST method	Modal MIC (mg/liter)	ECOFF Finder	MicDat1.23 software	ECOFF Finder	MicDat1.23 software	ECOFF Finder	MicDat1.23 software		
FLC									
CLSI	64	NA	64	NA	64	NA	64	128	ND
EUCAST	64	NA	64	NA	64	NA	64	128	ND
ITC									
CLSI	0.125	0.5	0.5	0.5	0.5	1	1	0.25	0.5
EUCAST	0.125	1	1	1	1	2	2	1	0.5
VRC									
CLSI	0.5	8	8	16	16	32	16	1	ND
EUCAST	1	4	4	4	4	8	8	2	ND
ISA									
CLSI	0.25	1	1	2	1	2	2	0.5	ND
EUCAST	0.5	0.125	2	0.25	4	0.25	4	1	1
POS									
CLSI	0.016	0.125	0.125	0.125	0.125	0.25	0.25	0.125	ND
EUCAST	0.032/0.64	0.125	0.125	0.25	0.25	0.25	0.25	0.125	0.25
AMB									
CLSI	0.5	1	2	2	2	2	2	2	2
EUCAST	1	NA	1	NA	1	NA	1	2	1
AFG									
CLSI	0.125	0.25	0.5	0.25	0.5	0.25	1	0.25	0.5
EUCAST	0.06	0.25	1	0.25	1	0.5	2	0.25	1
MFG									
CLSI	0.125	0.25	0.25	0.25	0.25	0.25	0.5	0.25	0.5
EUCAST	0.125	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.5

^aThe statistical ECOFF determination method we used is described in reference 21, and the derivatization ECOFF determination method is described in reference 22. The ECOFF Finder program (21) will soon be freely available at the EUCAST website (www.eucast.org). NA, not available; the ECOFF Finder program could not provide an ECOFF.

^bND, not determined; an ECOFF could not be determined by the visual method when distributions were truncated or bi- or trimodal with no clear main wild-type population.



Direnç

VE

Aspergillus

Epidemiology and molecular mechanisms of antifungal resistance in *Candida* and *Aspergillus*

Sarah Santos Gonçalves,¹ Ana Carolina Remondi Souza,¹ Anuradha Chowdhary,² Jacques F. Meis^{3,4} and Arnaldo Lopes Colombo¹

Mycoses, 2016, **59**, 198–219

Table 3 Primary resistance among *Aspergillus* species.

Species	Invasive disease	Resistance profile	Section	References
<i>A. lentulus</i>	Relatively common	Primary resistance to AMB, azoles and varied resistance to CFG	<i>Fumigati</i>	[48, 301–304]
<i>A. udagawae</i>	Rare	Primary resistance to AMB and VRC	<i>Fumigati</i>	[66, 301, 302, 305, 306]
<i>N. pseudofisherii</i>	Rare	Varied susceptibility to AMB and lower susceptibility to azoles	<i>Fumigati</i>	[48, 302, 307]
<i>A. fumigatiaffinis</i>	Not reported	Primary resistance to AMB and lower resistance to azoles	<i>Fumigati</i>	[49]
<i>A. viridinutans</i>	Rare	Lower susceptibility to AMB and azoles	<i>Fumigati</i>	[48, 308]
<i>A. flavus</i>	Relatively common	Primary resistance to AMB and varied susceptibility to CFG	<i>Flavi</i>	[54, 66, 302, 309–313]
<i>A. allilaceus</i>	Relatively common	Lower susceptibility to AMB and CFG	<i>Flavi</i>	[302, 314–317]
<i>A. terreus</i>	Relatively common	Primary resistance to AMB	<i>Terrei</i>	[66, 276, 280, 318–321]
<i>A. alabamensis</i>	Not reported	Primary resistance to AMB	<i>Terrei</i>	[322]
<i>A. niger</i>	Rare	Varied and lower susceptibility to azoles	<i>Nigri</i>	[60, 310, 323–325]
<i>A. calidoustus</i>	Rare	Primary resistance to AMB and azoles and varied susceptibility to CFG	<i>Ustis</i>	[77, 302, 326, 327]
<i>A. versicolor</i>	Rare	Varied susceptibility to AMB and lower susceptibility to azoles	<i>Nidulantes</i>	[302, 325, 328]
<i>A. sydowii</i>	Rare	Varied susceptibility to AMB and lower susceptibility to azoles	<i>Nidulantes</i>	[60, 302, 328]

AMB, amphotericin B; CFG, caspofungin; IA, invasive aspergillosis; VRC, voriconazole.

1997: İlk itrakonazol-R *Aspergillus fumigatus*
:Yüksek ($\geq 16 \mu\text{g/ml}$) MİK, klinik
yanıtsızlık, destekleyen hayvan modeli
sonuçları

Emergence of Azole Resistance in *Aspergillus fumigatus* and Spread of a Single Resistance Mechanism

-I

Eveline Snelders^{1,2}, Henrich A. L. van der Lee^{1,2}, Judith Kuijpers^{1,2}, Anthonius J. M. M. Rijs^{1,2}, János Varga^{3,4}, Robert A. Samson³, Emilia Mellado⁵, A. Rogier T. Donders⁶, Willem J. G. Melchers^{1,2}, Paul E. Verweij^{1,2*}

ITRA DİRENCİ

1994-2007 izolatları
n=1912

Hollanda, tek merkez
Tarama testinde 8 µg/ml itra
varlığında üreme,
MİK: ≥ 4 µg/ml

GENOTİPLENDİRME ve DİRENC EPİDEMİYOLOJİSİ ÇALIŞMALARI

n=464
Çok merkez; Hollanda
ve ayrıca 6
Avrupa ülkesi

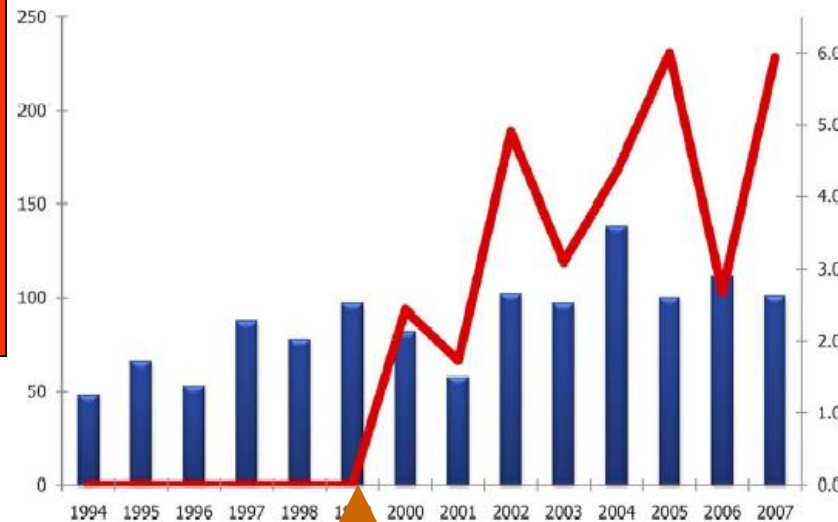


Figure 1. Epidemiology of ITZ Resistance in the *A. fumigatus* Isolates

Blue bars represent the number of patients with a positive *A. fumigatus* culture (left y-axis) and the red line represents the percentage of those patients with an ITZ+ isolate (right y-axis). The x-axis is the year.

doi:10.1371

Itra R oranı:
%1.7-6

1999'dan sonraki yıllarda izole edilen suşlarda itra direnci ortaya çıkıyor ve artarak devam ediyor.

Emergence of Azole Resistance in *Aspergillus fumigatus* and Spread of a Single Resistance Mechanism

-II

Eveline Snelders^{1,2}, Henrich A. L. van der Lee^{1,2}, Judith Kuijpers^{1,2}, Anthonius J. M. M. Rijs^{1,2}, János Varga^{3,4}, Robert A. Samson³, Emilia Mellado⁵, A. Rogier T. Donders⁶, Willem J. G. Melchers^{1,2}, Paul E. Verweij^{1,2*}

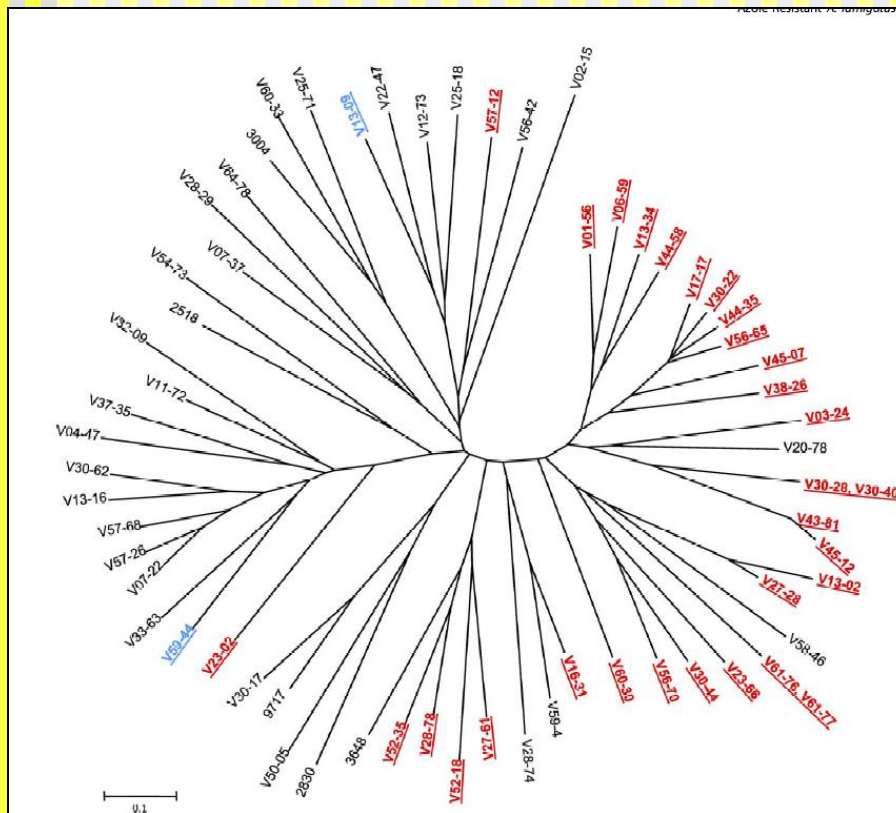


Figure 3. Genotypic Relatedness of 32 ITZ+ *A. fumigatus* Isolates and 32 ITZ− Controls

The numbers are the identification numbers of the individual isolates. The ITZ+ *A. fumigatus* with the L98H substitution and a tandem repeat in the promoter region of the *cyp51A* gene are underlined and printed in red. The ITZ+ isolates with other or unknown resistance mechanisms are underlined and printed in blue. The ITZ− isolates are printed in black.

doi:10.1371/journal.pmed.0050219.g003

Itra-R suşların genotipik ilişkileri:

Kırmızı: Cyp51A geninde L98H değişimi ile birlikte genin promoter bölgesinde 'tandem repeat' olan Itra-R suşlar

Suşların çoğunluğu. Bu suşlar, genotipik olarak birbirleriyle ilişkili. Bir kısmı, daha önce hiç azol almamış hastalardan...

J.W.M. van der Linden, M.C. Arendrup,
A. Warris, K. Lagrou, H. Pelloux, P.M. Hauser,
E. Chrysanthou, E. Mellado, S.E. Kidd,
A.M. Tortorano, E. Dannaoui, P. Gaustad,
J.W. Baddley, A. Uekötter, C. Lass-Flörl,
N. Klimko, C.B. Moore, D.W. Denning,
A.C. Pasqualotto, C. Kibbler, S. Arian-Akdagli,
D. Andes, J. Meletiadis, L. Naumiuk,
M. Nucci, W.J.G. Melchers, P.E. Verweij

Prospective Multicenter International Surveillance of Azole Resistance in *Aspergillus fumigatus*

To investigate azole resistance in clinical *Aspergillus* isolates, we conducted prospective multicenter international surveillance. A total of 3,788 *Aspergillus* isolates were screened in 22 centers from 19 countries. Azole-resistant *A. fumigatus* was more frequently found (3.2% prevalence) than previously acknowledged, causing resistant invasive and noninvasive aspergillosis and severely compromising clinical use of azoles.

A. fumigatus - itrakonazol direnç oranı

ÜLKE	SUŞUN KAYNAĞI	R %	REFERANS
Danimarka	Klinik	4.5	Mortensen JCM 2011
Fransa	Klinik	0.85-4.6	Alanio JAC 2011; Burgel AAC 2012;
Hindistan	Klinik	1.9	Chowdhary JAC 2012
Hollanda	Klinik	5.3-38	Van der Linden Emerg Infect Dis 2011,; van Ingen JAC 2015
İngiltere	Klinik	5	Howard Emerg Infect Dis 2009
Japonya	Klinik	7	Tashiro AAC 2012
Danimarka	Çevre	5.8-14.2	Mortensen AAC 2010
Hollanda	Çevre	12.2	Snelders Appl &Env Microbiol 2009
İran	Çevre	3.3	Badali Mycoses 2013

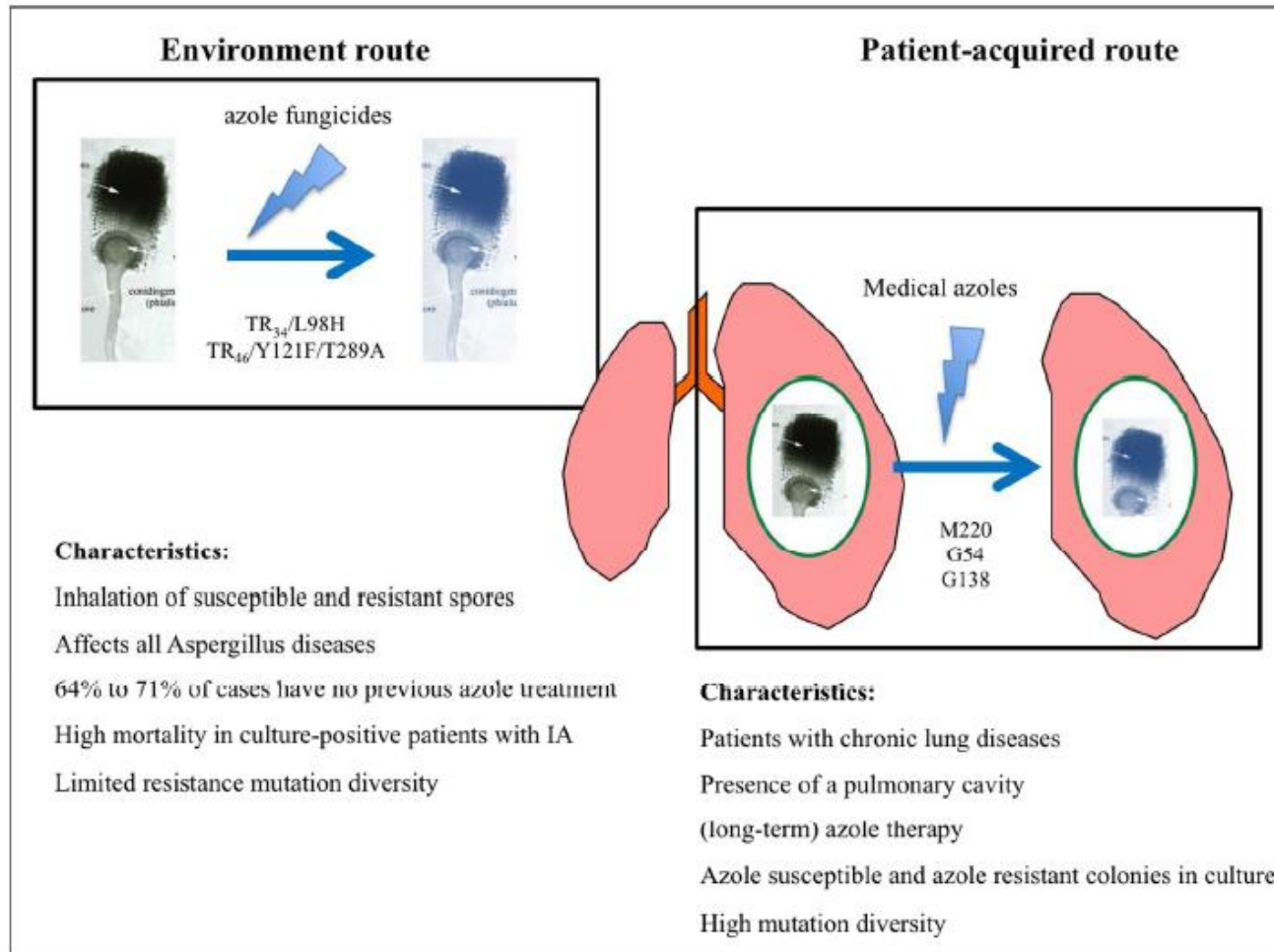
First determination of azole resistance in *Aspergillus fumigatus* strains carrying the TR34/L98H mutations in Turkey

Gülşah Ece Özmerdiven^a, Seçil Ak^b, Beyza Ener^{a,*}, Harun Ağca^a, Burcu Dalyan Cilo^a, Berrin Tunca^b, Halis Akalın^c

J Infect Chemother 21 (2015) 581–586

exhibited good activity against all isolates whereas the azoles were resistant. Sequence analysis of the promoter region and *CYP51A* gene indicated the presence of TR34/L98H in 86.8% (n = 66) of isolates. This initial analysis of the resistance mechanism of *A. fumigatus* from Turkey revealed a common TR34/L98H mutation in the *cyp51A* gene.

Olgular azol-R *Aspergillus* suşlarını ne yolla alıyor olabilir?



Characteristics of azole resistant *Aspergillus* infections by the environmental- and patient route.

Table 1. Common resistance mechanisms reported in the *cyp51A* gene of clinical and environmental *Aspergillus fumigatus*

Promotor tandem repeat (TR) insertion and Point Mutation	Country	Medical azoles affected	Reference
TR34/L98H	Netherlands, UK, Germany, Belgium, France, Spain, Italy, Austria, Denmark, Poland, Ireland, China, USA, India, Japan, Colombia, Taiwan, Turkey, Kuwait, Iran, Pakistan, Australia, Tanzania	Itraconazole, voriconazole, posaconazole, isavuconazole	24,55,56,57,59,60, 62,63,64,65,66,67, 68,69,70,71,72,73,74,76
TR46/Y121F/T289A	Netherlands, UK, Germany, Belgium, France, Spain, Denmark, Ireland, China, USA, India, Japan, Colombia, Tanzania	Itraconazole (variable), voriconazole, posaconazole (variable), isavuconazole	24,55,56,58,61, 62,75,76,77
TR53	Netherlands, Colombia	Itraconazole, voriconazole, posaconazole, isavuconazole	24,77
G54	Netherlands, UK, Germany, France, Spain, India, Romania, China, USA, Tanzania	Itraconazole, posaconazole	24
M220	Netherlands, UK, Germany, Belgium, France, Spain, China, USA	Itraconazole, posaconazole	24
Y121F	France	Voriconazole	92

Hangi triazole direnç ?

	ITR	VOR	POS	ISV
TR*34/L98H	R	R	R	R
TR46/Y121F/T289A	V	R	R/V	
TR53	R	R	↓ S	
G54	R	S	R	
M220	R	↓ S	V	
G448	↓ S	R	↓ S	

*CYP51A geninin promoter kısmında 34 , 46 ya da bp'lik bir diziden iki kopya ("tandem repeat") bulunması ve CYP51A ekspresyonunda artış

Meis et al. Philos Trans R Soc Lond B Biol Sci. Clinical implications of globally emerging azole resistance in *A. fumigatus*

A-Alanin
F-Fenilalanin
G-Glisin
H-Histidin
L-Lösin
M-Metionin
T-Treonin
Y-Tirozin

**Retrospektif
H.Ü. n=215
1997-2015**

**Agar tarama $\Rightarrow$ EUCAST mikrodilüsyon
cyp51A sekans analizi**

Retrospektif
H.Ü. , n=215 , 1997-2015
Agar tarama ⇒ EUCAST mikrodilüsyon
cyp51A sekans analizi

Klinik A. fumigatus izolatları
Azol direnci

İzolat no.										
4C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	
6C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	
7C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	
10C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	
16C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	Q141H
22C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	
23C	Y46F	G89G	V172M	T248N	E255D	L358L	K427E	C454C	Y431D	

Emergence of Echinocandin Resistance due to a Point Mutation in the *fks1* Gene of *Aspergillus fumigatus* in a Patient with Chronic Pulmonary Aspergillosis

Cristina Jiménez-Ortigosa^{1*}, Caroline Moore², David W. Denning³ and David S. Perlin¹

We have identified the first case of a *fks1* hot spot 1 point mutation causing echinocandin resistance in a clinical *Aspergillus fumigatus* isolate recovered from a chronic pulmonary aspergillosis patient with an aspergilloma who first failed azole and polyene therapy, and then subsequently failed micafungin treatment.

Table 2. MIC/MEC distributions of the antifungal drugs tested in the study for the *Aspergillus* spp. clinical isolates. Alterations in the Cyp51A and Fks1 together with the sequence type (ST) are also shown. Three wild type (WT) strains have also been included for comparison purposes.

#	Lab no.	Date received	ID	MIC/MEC (mg/L)								Cyp51A*	Fks1*	ST
				ISA	ITR	POS	VOR	CSF	MCF	AMB	TERB			
1	22432	2/9/2010	<i>Aspergillus fumigatus</i>	0.25	0.25	0.12	0.25	0.12	0.03	2	1	Y46F, V172M, T248N, E255D, K427E	ND ^b	9
2	23525	5/4/2010	<i>Aspergillus fumigatus</i>	0.25	0.25	0.12	0.12	0.12	0.03	2	0.5	Y46F, V172M, T248N, E255D, K427E	ND	9
3	24053A	6/15/2010	<i>Aspergillus fumigatus</i>	0.25	0.25	0.25	0.25	0.12	0.03	2	1	Y46F, V172M, T248N, E255D, K427E	S53G	12
4	24053B	6/15/2010	<i>Aspergillus fumigatus</i>	0.5	0.25	0.25	1	2	2	2	1	Y46F, V172M, T248N, E255D, K427E	S53G and F675S	9
5	24555	7/28/2010	<i>Aspergillus fumigatus</i>	0.25	0.25	0.25	0.25	0.12	0.03	2	0.5	Y46F, V172M, T248N, E255D, K427E	ND	9
6	29576A	7/27/2011	<i>Aspergillus flavus</i>	0.25	0.5	0.25	0.25	0.12	0.03	2	0.06	WT ^a	ND	ND
7	29576B	7/27/2011	<i>Aspergillus fumigatus</i>	0.25	0.25	0.25	0.25	0.12	0.03	2	0.5	Y46F, V172M, T248N, E255D, K427E	ND	9
8	30906	10/25/2011	<i>Aspergillus fumigatus</i>	0.25	0.5	0.25	0.12	0.12	0.03	2	1	Y46F, V172M, T248N, E255D, K427E	ND	12
9	33460	4/10/2012	<i>Aspergillus fumigatus</i>	0.25	0.5	0.25	0.12	0.12	0.03	2	1	Y46F, V172M, T248N, E255D, K427E	ND	9
10	45755	12/30/2013	<i>Aspergillus fumigatus</i>	2	1	1	0.5	0.12	0.03	2	1	Y46F, V172M, T248N, E255D, K427E	ND	12
11	53619	1/28/2015	<i>Aspergillus fumigatus</i>	0.25	0.5	0.25	0.25	0.12	0.03	2	0.5	Y46F, V172M, T248N, E255D, K427E	ND	9
12	62194	12/24/2015	<i>Aspergillus fumigatus</i>	0.25	0.5	0.25	0.12	0.12	0.03	2	0.5	Y46F, V172M, T248N, E255D, K427E	ND	9
	ATCC13073		<i>Aspergillus fumigatus</i>	0.25	0.25	0.25	0.25	0.12	0.03	2	1	ND	S53G	ND
	R21		<i>Aspergillus fumigatus</i>	0.25	0.25	0.25	0.25	0.12	0.03	2	2	ND	WT	ND
	Af293		<i>Aspergillus fumigatus</i>	0.12	0.12	0.12	0.12	0.06	0.03	1	0.5	ND	WT	ND

*Ref strain used Af293

^bND = not determined

^aRef strain used *A. flavus* NRRL3357

Sunum Planı

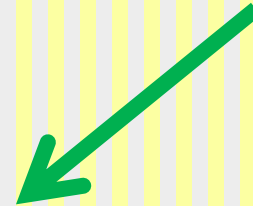
Temel bilgiler



YÖNTEM ve
DEĞERLENDİRME



DİRENÇ VARLIĞI /
ORANI



ÖNERİLER

Clinical Practice Guideline for the Management of Candidiasis: 2016 Update by the Infectious Diseases Society of America

Peter G. Pappas,¹ Carol A. Kauffman,² David R. Andes,³ Cornelius J. Clancy,⁴ Kieren A. Marr,⁵ Luis Ostrosky-Zeichner,⁶ Annette C. Reboli,⁷ Mindy G. Schuster,⁸ Jose A. Vazquez,⁹ Thomas J. Walsh,¹⁰ Theoklis E. Zaoutis,¹¹ and Jack D. Sobel¹²

Clinical Infectious Diseases® 2016;63(4):433–42

Practice Guidelines for the Diagnosis and Management of Aspergillosis: 2016 Update by the Infectious Diseases Society of America

Thomas F. Patterson,^{1,a} George R. Thompson III,² David W. Denning,³ Jay A. Fishman,⁴ Susan Hadley,^{5,b} Raoul Herbrecht,⁶ Dimitrios P. Kontoyiannis,⁷ Kieren A. Marr,⁸ Vicki A. Morrison,⁹ M. Hong Nguyen,¹⁰ Brahm H. Segal,¹¹ William J. Steinbach,¹² David A. Stevens,¹³ Thomas J. Walsh,¹⁴ John R. Wingard,¹⁵ Jo-Anne H. Young,¹⁶ and John E. Bennett^{17,a}

ESCMID-EFISG Aspergillus Guideline 2018

Recommendations for identification of species by molecular methods for prediction of primary AMB resistance

Population	Intention	Intervention	SoR	QoE	Reference	Comment
Clinically relevant <i>Asp</i> isolate	Predict primary AMB resistance according to the species	Species ID by using β tubulin or calmodulin sequencing	B	III	Alastruey-Izquierdo AAC 2013 Blum AAC 2013 Colozza Intl J Antimicrob Agents 2012 Escibano CMI 2012 Baddley JCM 2009 Lass-Flörl Br J Haem 2005 Arikan Med Mycol 2008 Barchiesi JAC2013 (murine) Hadrach Med Mycol 2012 Shivaprakash DMID 2011 Arendrup EUCAST-AFST CMI 2012 Espinel-Ingroff AAC 2011	<i>A. terreus</i> <i>A. flavus</i>

ESCMID-EFISG Aspergillus Guideline 2018

Intrinsic resistance – Antifungal regimen

Intention	Intervention	SoR	QoE	Comment
Change tx. if AMB MIC ≥ 1 mg/L.	Replace AMB with azole, if azole-S	B	II	
Treat <i>Aspergillus</i> disease due to:	Voriconazole	A	II	Avoid polyenes
<i>A. terreus</i>	Isavuconazole	A	II	
	Posaconazole	B	III	Avoid polyenes
	Itraconazole	B	III	Avoid polyenes
<i>A. calidoustus</i>	Lipid AMB	A	II	Avoid azoles
<i>A. tubingensis</i> (<i>A. niger</i> complex)	Avoid azole monotherapy	C	III	No data: higher azole MICs are common but clinical impact unknown
<i>A. lentulus</i> (<i>A. fumigatus</i> complex)				
<i>A. alliaceus</i> (<i>A. flavus</i> complex)	Avoid AmB	C	III	
<i>A. niger</i> complex	Avoid itra & isavu	B	III	MICs are in general 1 step higher compared to <i>A. fumigatus</i> for ISAV, POS and VCZ and 2 steps higher for ITZ but limited clinical data.
<i>A. nidulans</i>	Prefer VCZ	C	IIIa	Elevated AmB MICs, poor clinical responses in CGD

Azol direnci klinik yanıtsızlığa yol açan
bağımsız bir risk faktörü müdür?

Direnç oranı ?

Primer tedavi seçeneğinin değiştirilmesi gerekliliği ?

Impact of azole resistance on *Aspergillus* guidelines

Georgiadou and Kontoyiannis Ann NY Acad Sci **2012**; 1272: 15-22

Table 1. Variability of azole resistance prevalence in *Aspergillus* isolates among medical centers and published studies: a multifactorial issue

Parameters

Patient referral patterns (acute versus chronic aspergillosis)
Selected versus random specimens
Time factors (recent versus older series)
Small collections of isolates (e.g., <100 isolates)
Differences in screening methods (microdilution versus plate method)
Exclusion of cryptic *Aspergillus* species
Rates using different denominators (patient versus isolates)
Different susceptibility methods and inocula
Different breakpoints^a and definitions of resistance
Differences in geographic location of the center

Conclusions

Resistance of *Aspergillus* spp. to azoles might be on the rise; nevertheless, inferences are drawn from incomplete observational data, and it is rather early to consider that azole resistance is becoming a menacing worldwide phenomenon. A better understanding of the epidemiology and the biologic plausibility of resistance would assist in the development of better diagnostic strategies (probably PCR-based assays) for both prevention and treatment. Currently, we need better field studies but not necessarily modification of guidelines.

European Centre for Disease Prevention and Control

Risk assessment on the impact of environmental usage of triazoles on the development and spread of resistance to medical triazoles in *Aspergillus* species Stockholm: ECDC; 2013

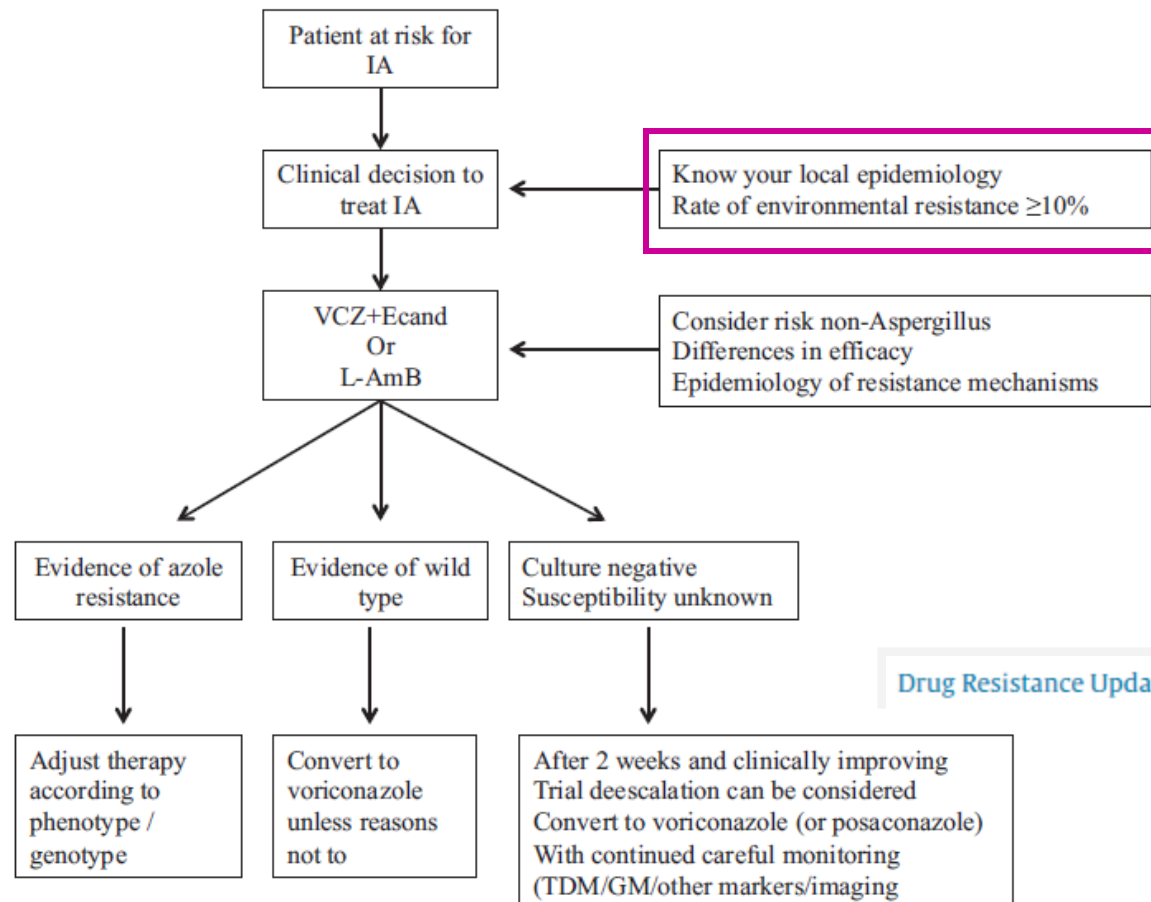
www.ecdc.europa.eu/en/publications/publications/risk-assessment-impact-environmental-usage-of-triazoles-on-aspergillus-spp-resistance-to-medical-triazoles.pdf

Direnç nasıl gelişiyor ?
"Çevre"de direnç: Klinik etki ? Ekonomik etki ?

("This report of the European Centre for Disease Prevention and Control (ECDC) was coordinated by Niels Kleinkauf and Dominique Monnet. Contributing authors Niels Kleinkauf, Paul E. Verweij, Maiken C. Arendrup, Peter J. Donnelly, Manuel Cuenca-Estrella, Bart Fraaije, Willem J.G. Melchers, Niels Adriaenssens, Gert H.J. Kema, Andrew Ullmann, Paul Bowyer, David W. Denning This report was sent for consultation to the European Food Safety Authority - EFSA.")

International expert opinion on the management of infection caused by azole-resistant *Aspergillus fumigatus*

Paul E. Verweij^{a,*}, Michelle Ananda-Rajah^b, David Andes^c, Maiken C. Arendrup^d, Roger J. Brüggemann^e, Anuradha Chowdhary^f, Oliver A. Cornely^g, David W. Denning^h, Andreas H. Grollⁱ, Koichi Izumikawa^j, Bart Jan Kullberg^k, Katrien Lagrou^l, Johan Maertens^m, Jacques F. Meis^{a,n}, Pippa Newton^h, Iain Page^h, Seyedmojtaba Seyedmousavi^a, Donald C. Sheppard^o, Claudio Viscoli^p, Adilia Warris^q, J. Peter Donnelly^r



Drug Resistance Updates 21–22 (2015) 30–40

Fig. 3. Management of patients with clinical suspicion of IPA in regions with environmental resistance of $\geq 10\%$. IA, invasive aspergillosis; L-AmB, liposomal amphotericin B; VCZ, voriconazole; Ecand, echinocandin; TDM, therapeutic drug monitoring; GM, galactomannan.

Azole resistance surveillance in *Aspergillus fumigatus*: beneficial or biased?

Paul E. Verweij^{1,2*}, Pieter P. A. Lestrade¹, Willem J. G. Melchers^{1,2} and Jacques F. Meis^{2,3}

J Antimicrob Chemother 2016; **71**: 2079–2082

Table 1. Characteristics of three surveillance strategies for azole resistance

Surveillance strategy	Aim	Advantage	Disadvantage	Bias
Screening of unselected clinical <i>A. fumigatus</i> isolates	Trends in resistance frequency Trends in distribution of resistance mutations Detection of new resistance mutations	A substantial number of isolates may be screened Relatively straightforward to conduct	Includes isolates that are clinically not relevant or from colonized/untreated patients May not correspond to resistance frequency in specific risk groups	Denominator bias Laboratory practice and procedure bias in multicentre design
Identify patients with IA based on a positive <i>Aspergillus</i> culture	Determine the frequency of resistance in patients with culture-positive IA	Provides a relatively high sample size, when patients are included independent of risk category	Includes only a minority of IA patients Variability in underlying diseases Variability in management/sampling strategy	Case definition, case ascertainment and sampling bias Laboratory practice and procedure bias in multicentre design
Determine the frequency of IA within a cohort of patients	To determine the prevalence of azole resistance in a specific risk group	Gives an accurate indication of resistance prevalence Relatively uniform management and sampling strategy in single-centre design	Requires a substantial effort to evaluate and classify patients May take a long time to include a meaningful number of cases Variability in management/sampling strategy in multicentre design	Case definition, case ascertainment and sampling bias Laboratory practice and procedure bias in multicentre design

Azole resistance surveillance in *Aspergillus fumigatus*: beneficial or biased?

Paul E. Verweij^{1,2*}, Pieter P. A. Lestrade¹, Willem J. G. Melchers^{1,2} and Jacques F. Meis^{2,3}

J Antimicrob Chemother 2016; **71**: 2079–2082

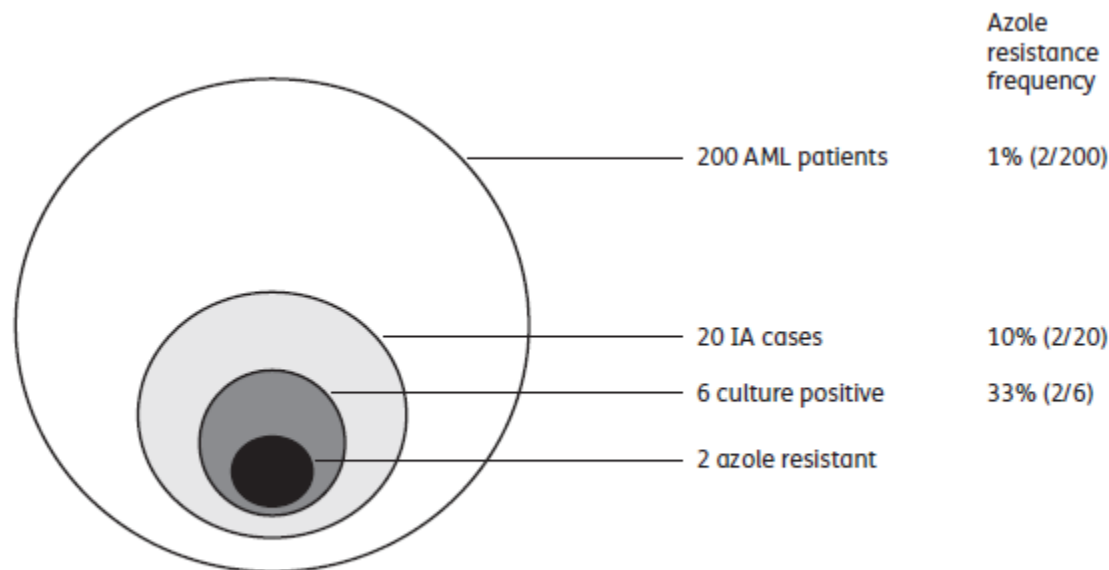


Figure 1. Impact of the denominator on the resistance frequency in azole resistance surveillance. Resistance rates are calculated for a theoretical population of 200 AML patients, with a 10% prevalence of IA and 33% culture positivity.

“.....**Azole resistance frequencies have been based on** screening of unselected *A. fumigatus* isolates, on the number of azole-resistant cases within a cohort of patients with a specific *Aspergillus* disease, or on analysis of patients within a specific risk group.....”

Changes in In Vitro Susceptibility Patterns of *Aspergillus* to Triazoles and Correlation With Aspergillosis Outcome in a Tertiary Care Cancer Center, 1999–2015

Sang Taek Heo,^{1,2,a} Alexander M. Tatara,^{1,3,a,c} Cristina Jiménez-Ortigosa,⁴ Ying Jiang,¹ Russell E. Lewis,^{1,b} Jeffrey Tarrand,⁵ Frank Tverdek,⁶ Nathaniel D. Albert,¹ Paul E. Verweij,⁷ Jacques F. Meis,⁷ Antonios G. Mikos,³ David S. Perlin,⁴ and Dimitrios P. Kontoyiannis¹

Methods. We examined changes over time in triazole minimum inhibitory concentrations (MICs) of 290 sequential *Aspergillus* isolates recovered from respiratory sources during 1999–2002 (before introduction of the *Aspergillus*-potent triazoles voriconazole and posaconazole) and 2003–2015 at MD Anderson Cancer Center. We also tested for polymorphisms in ergosterol biosynthetic genes (*cyp51A*, *erg3C*, *erg1*) in the 37 *Aspergillus fumigatus* isolates isolated from both periods that had non-wild-type (WT) MICs. For the 107 patients with hematologic cancer and/or HSCT with invasive pulmonary aspergillosis, we correlated in vitro susceptibility with 42-day mortality.

Results. Non-WT MICs were found in 37 (13%) isolates and was only low level (MIC <8 mg/L) in all isolates. Higher-triazole MICs were more frequent in the second period and were *Aspergillus*-species specific, and only encountered in *A. fumigatus*. No polymorphisms in *cyp51A*, *erg3C*, *erg1* genes were identified. There was no correlation between in vitro MICs with 42-day mortality in patients with invasive pulmonary aspergillosis, irrespective of antifungal treatment. Asian race (odds ratio [OR], 20.9; 95% confidence interval [CI], 2.5–173.5; $P = .005$) and azole exposure in the prior 3 months (OR, 9.6; 95% CI, 1.9–48.5; $P = .006$) were associated with azole resistance.

Conclusions. Non-WT azole MICs in *Aspergillus* are increasing and this is associated with prior azole exposure in patients with hematologic cancer or HSCT. However, no correlation of MIC with outcome of aspergillosis was found in our patient cohort.

Table 4. Crude Mortality Within 42 Days in Patients With Invasive Pulmonary Aspergillosis According to Various Minimum Inhibitory Concentration Cutoffs of the *Aspergillus* Isolates^a

	Rate of Mortality Within 42 Days, no./No. (%)		
MIC Cutoff, $\mu\text{g/mL}$	MIC $\leq$ Cutoff	MIC > Cutoff	P Value ^c
Itraconazole			
1	30/83 (36)	7/19 (37)	.95
2	33/91 (36)	4/11 (36)	>.99
3	35/95 (37)	2/7 (29)	>.99
4	37/102 ^b (36)	NA	...
Voriconazole			
1	30/86 (35)	7/16 (44)	.50
2	36/98 (37)	1/4 (25)	>.99
3	36/100 (36)	1/2 (50)	>.99
Posaconazole			
0.25	31/85 (36)	6/17 (35)	.93
0.5	34/93 (37)	3/9 (33)	>.99
1	36/97 (37)	1/5 (20)	.65
2	37/102 (36)	NA	...
Isavuconazole			
1	37/102 (36)	NA	...

Abbreviations: MIC, minimum inhibitory concentration; NA, not applicable.

^aClinical and Laboratory Standards Institute criteria.

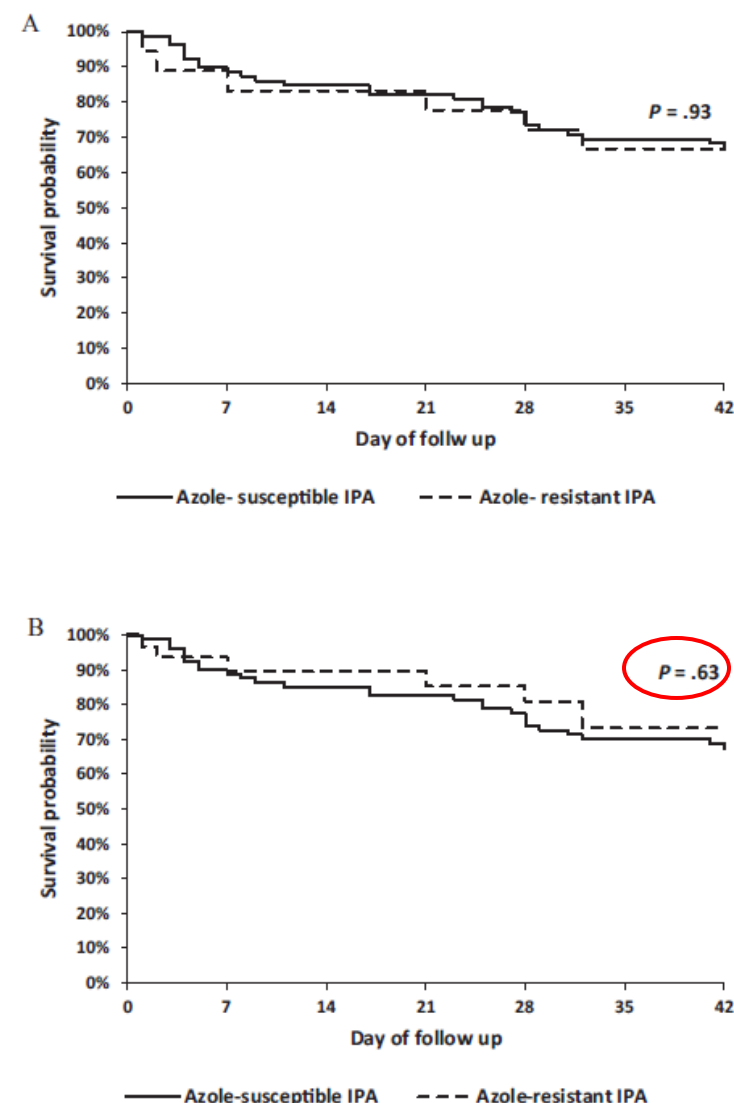


Figure 2. Survival curves for patients who had either azole wild-type or azole non-wild-type invasive pulmonary aspergillosis (IPA). *A*, Kaplan-Meier survival curve. *B*, Adjusted Kaplan-Meier survival curve (adjusted for history of neutropenia, lymphopenia, intensive care unit admission, and time period of IPA diagnosis).

PCR-based detection of *Aspergillus fumigatus* Cyp51A mutations on bronchoalveolar lavage: a multicentre validation of the AsperGenius assay[®] in 201 patients with haematological disease suspected for invasive aspergillosis

G. M. Chong^{1*}, M. T. van der Beek², P. A. von dem Borne³, J. Boelens⁴, E. Steel⁵, G. A. Kampinga⁶, L. F. R. Span⁷, K. Lagrou⁸, J. A. Maertens⁹, G. J. H. Dingemans¹⁰, G. R. Gaajetaan¹⁰, D. W. E. van Tegelen¹⁰, J. J. Cornelissen¹¹, A. G. Vonk¹² and B. J. A. Rijnders¹

Results: Two hundred and one patients each contributed one BAL sample, of which 88 were positive controls and 113 were negative controls. The optimal cycle threshold cut-off value for the *Aspergillus* species PCR was <38. With this cut-off, the PCR was positive in 74/88 positive controls. The sensitivity, specificity, positive predictive value and negative predictive value were 84%, 80%, 76% and 87%, respectively. 32/74 BAL samples were culture negative. Azole treatment failure was observed in 6/8 patients with a RAM compared with 12/45 patients without RAMs ($P=0.01$). Six week mortality was 2.7 times higher in patients with RAMs (50.0% versus 18.6%; $P=0.07$).

Conclusions: The AsperGenius[®] assay had a good diagnostic performance on BAL and differentiated WT from *Aspergillus fumigatus* with RAMs, including in culture-negative BAL samples. Most importantly, detection of RAMs was associated with azole treatment failure.

SONUÇ ve ÖNERİLER

- Tanımlama: Primer direnç
- AFDT: Standart yöntem (endikasyonlar)
 - Güncel direnç sınır değerleri; "ECV"
 - Direnç oranı
 - Multipl faktörler

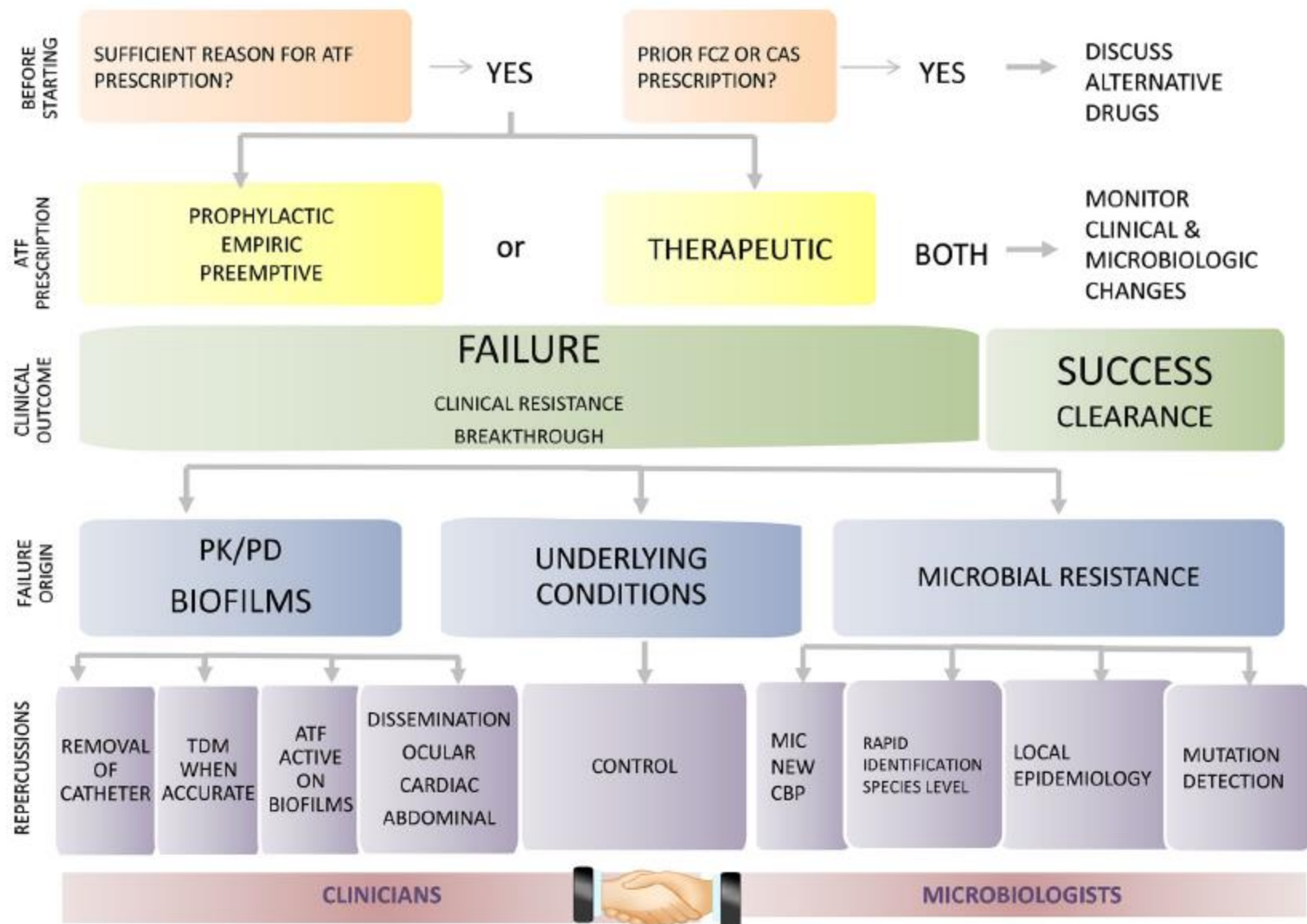


Fig. 3 Bedside strategy for circumventing antifungal drug resistance in 2014. *ATF* antifungal drug, *FCZ* fluconazole, *CAS* caspofungin, *PK/PD* pharmacokinetics and pharmacodynamics, *TDM* therapeutic drug monitoring, *MIC* minimal inhibitory concentration, *CBP* clinical breakpoint