

Insights into Antifungal Resistance & Newer (Other) Moulds

Nathan P. Wiederhold, PharmD
Fungus Testing Laboratory (FTL)



UT Health
San Antonio

Department of Pathology
& Laboratory Medicine
South Texas Reference Laboratories

Disclosures

Funding to FTL

- Astellas
- bioMerieux
- Cepheid
- Cidara
- F2G
- Pfizer
- Viamet

Speakers Bureau

- Gilead

Member, CLSI Antifungal
Susceptibility Subcommittee

Road Map

- Antifungal activity/resistance in newer/ cryptic species (& other moulds)
 - ❖ *Aspergillus fumigatus* & azole resistance
 - ❖ Cryptic *Aspergillus* spp.
 - ❖ Mucorales
 - ❖ *Fusarium* spp.
 - ❖ *Scedosporium* spp.
 - ❖ Other moulds (intrinsic microbiologic resistance)
- Challenges with mould susceptibility testing & interpretation of results

Azole Resistant *A. fumigatus*

Continent/Country	Percent Resistance	Isolate Source
<p><i>J Antimicrob Chemother</i> 2017; 72: 2443–2446 doi:10.1093/jac/dkx168 Advance Access publication 1 June 2017</p> <p>Journal of Antimicrobial Chemotherapy</p> <p>Isolation of azole-resistant <i>Aspergillus fumigatus</i> from the environment in the south-eastern USA</p> <p>Steven F. Hurst¹, Elizabeth L. Berkow¹, Katherine L. Stevenson², Anastasia P. Litvintseva¹ and Shawn R. Lockhart^{1*}</p>		
United States	0.6 – 11.8%	Clinical & Environmental
Colombia (TR ₅₃)	3.3%	Environmental
Africa (TR ₃₄ /L98H, TR ₄₆ /Y121F/T289A)		
Tanzania	13.9%	Environmental
Asia & Australia (G54, M222, G448S, TR ₃₄ /L98H, TR ₄₆ /Y121F/T289A)		
Australia	2.6%	Clinical
Taiwan, China, India, Iran, Japan, Kuwait, Pakistan	1.9 – 11.1%	Clinical & Environmental

Rivero-Menendez *J Fungi* 2016;21: doi:10.3390/jof2030021. Wu et al. *Mycoses* 2015;58:544-549.

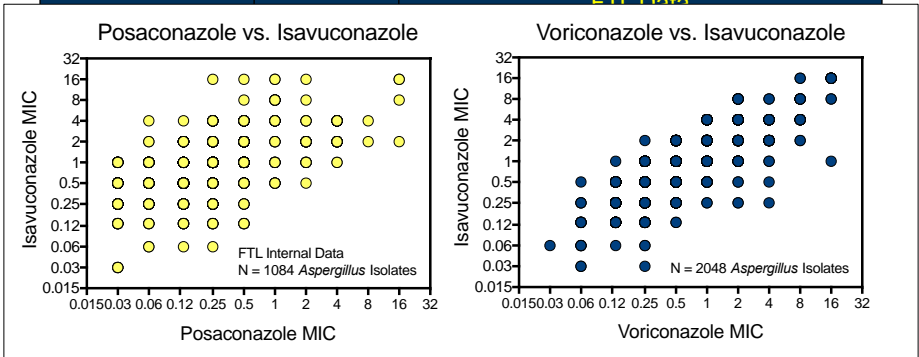
Cryptic *Aspergillus* Species

TRANSNET (US 2001-2006)	218 <i>Aspergillus</i> isolates 11% cryptic species	<i>A. lentulus</i> (1.8%) <i>A. udagawae</i> (1.4%) <i>A. tubingensis</i> (2.8%) <i>A. calidoustus</i> (2.8%)
FILPOP (Spain Oct 2010 & May 2011)	323 <i>Aspergillus</i> isolates 14.5% cryptic species	<i>A. lentulus</i> (1.1%) <i>A. alliaceus</i> (1.1%) <i>A. tubingensis</i> (7.9%) <i>A. calidoustus</i> (1.4%)

Section	Cryptic Species	Antifungal Susceptibility
<i>Fumigati</i>	<i>A. lentulus</i> <i>A. thermomutatus</i> <i>A. udagawae</i> <i>A. felis</i>	Reduced azole susceptibility
<i>Flavi</i>	<i>A. alliaceus</i>	Reduced susceptibility to amphotericin B and echinocandins
<i>Nidulantes</i>	<i>A. spinulosporus</i> <i>A. quadrilineatus</i>	Variable amphotericin B susceptibility azole susceptible
<i>Nigri</i>	<i>A. tubingensis</i>	Variable azole susceptibility
<i>Usti</i>	<i>A. calidoustus</i>	Reduced amphotericin B and azole susceptibility
<i>Versicolores</i>	<i>A. sydowii</i> <i>A. versicolor</i>	Variable amphotericin B susceptibility & reduced azole susceptibility

Aspergillus Species U.S. Isolates (2015-2018)

Section	Percentage	Prevalent spp. in Sections (% Overall) FTL Data
---------	------------	--



<i>Circumdati</i>	1.0%	<i>A. subramanianii</i> (0.3%)
<i>Aspergillus & Cremei</i>	0.6%	---

FTL Internal Data (1889 *Aspergillus* human isolates, excluding those from ears and nails)

Mucorales

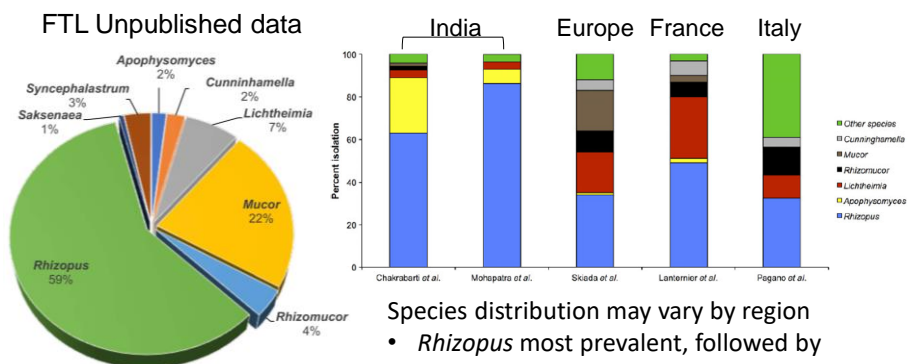
Common causes of non-*Aspergillus* invasive mould infections

Genera	Representative Species (not comprehensive)
<i>Rhizopus</i> (most prevalent)	<i>R. arrhizus</i> / <i>deleamar</i> , <i>R. microsporus</i> , <i>R. schipperae</i>
<i>Mucor</i>	<i>M. circinelloides</i> (f. <i>circinelloides</i> & f. <i>janssenii</i>), <i>M. irregularis</i>
<i>Rhizomucor</i>	<i>R. pusillus</i>
<i>Lichtheimia</i> (formerly <i>Absidia</i>)	<i>L. corymbifera</i> , <i>L. ornata</i> , <i>L. ramosa</i>
<i>Cunninghamella</i>	<i>C. bertholletiae</i> , <i>C. echinulata</i>
<i>Apophysomyces</i>	<i>A. elegans</i> , <i>A. trapeziformis</i> , <i>A. variabilis</i> , <i>A. mexicanus</i>
<i>Saksenaana</i>	<i>S. erythrospora</i> , <i>S. oblongispora</i> , <i>S. vasiformis</i>
<i>Syncephalastrum</i>	<i>S. racemosum</i>

- Risk factors
 - Hematologic malignancies (\pm HSCT)
 - Prolonged neutropenia or corticosteroid use
 - Poorly controlled DM
 - Trauma (traumatic inoculation)

Order Mucorales

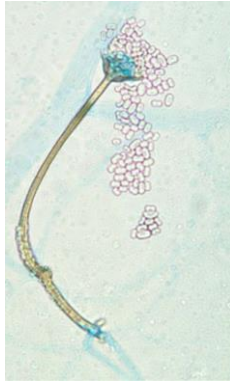
Species Distribution by Geographic Area



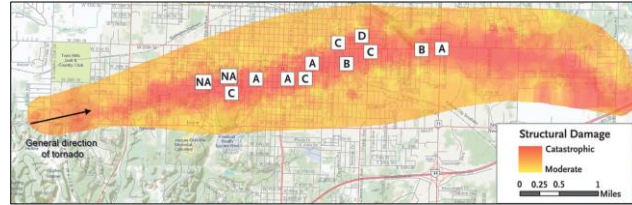
- 655 isolates (2.5 years)
- *Rhizopus* spp. most prevalent
 - 43.1% *R. arrhizus*
 - 22.3% *R. deleamar*

Alvarez et al. *J Clin Microbiol* 2009 47:1650-1660.
 Gomes et al. *Clin Microbiol Rev* 2011;24:411-445.
 Chakrabati & Singh *Mycoses* 2014;57(Suppl. 3):85-90.

Necrotizing Cutaneous Mucormycosis after a Tornado in Joplin, Missouri, in 2011



*Apophysomyces
trapeziformis*



13 patients with soft-tissue infections; 5 died; all in zone with most severe damage

Neblett et al. *N Engl J Med* 2012;367:2214-2225.

Mucorales Antifungal Susceptibility *What about Isavuconazole?*

Study	Arendrup et al. AAC 2015 (CLSI methodology)						Pfaller et al. AAC 2018	
	Amphotericin		Posaconazole		Isavuconazole		Isavuconazole	
Species	MIC Range	MIC50	MIC Range	MIC50	MIC Range	MIC50	MIC Range	MIC50
<i>R. oryzae</i> (arthizus)	0.03-0.25	0.06	0.12-0.5	0.5	0.12-2	1	0.25-4	1
<i>R. microsporus</i>	0.03-0.25	0.12	0.06-0.5	0.25	0.12-1	0.5	0.25->16	1
<i>M. circinelloides</i>	0.03-0.12	0.06	0.125->16	2	1-8	8	2->16	8
<i>Lichtheimia</i> spp.	0.03-0.12	0.03	0.12-0.5	0.25	0.5-2	1	1->16	4
	Chowdhary et al. <i>Mycoses</i> 2014						FTL Unpub. Data	
<i>Apophysomyces</i>	0.06-0.12	---	0.25-1	---	0.25-8	---	1-4	---
<i>Cunninghamella</i>	---	---	---	---	---	---	0.5->16	8
<i>Saksenaia</i>	---	---	---	---	---	---	0.25-2	---
<i>Syncephalastrum</i>	0.03-0.12	0.06	0.06-1	0.5	0.125-8	1	0.25->16	>16

In clinical setting, is reduced isavuconazole susceptibility offset by higher exposure?

Chowdhary et al. *Mycoses* 2014;57(Suppl. 3):97-107. Arendrup et al. *Antimicrob Agents Chemother* 2015;59:7735-7742.
Pfaller et al. *Antimicrob Agents Chemother* 2018 (Epub ahead of print).

Scedosporium & Scedosporiosis

- Invasive infections in immunocompromised patients (may mimic aspergillosis)
- Common colonizers in CF patients

Species	Taxonomic / Nomenclature Changes	FTL % Isolates	Lackner AAC 2012			
			AMB	VOR	ISA	PSC
<i>Scedosporium apiospermum</i>	Previously considered to be anamorph of <i>Pseudallescheria boydii</i>	57.2%	0.5->16	0.25-8	1->16	0.25->16
<i>Scedosporium boydii</i>	Previously <i>Pseudallescheria boydii</i>	22.5%	0.5->16	0.12-2	0.5->16	0.12->16
<i>Scedosporium aurantiacum</i>	Reduced amphotericin & azole susceptibility	1.4%	≥16	0.5-1	4-16	1->16
<i>Scedosporium dehoogii</i>	Rare Reduced voriconazole susceptibility	1.8%	2->16	0.5->16	2->16	0.5->16
<i>Scedosporium minutispora</i>	Previously <i>Pseudallescheria minutispora</i>	0.7%	1-4	0.25-2	2-16	0.5->16
<i>P. ellipsoidea</i>	Still <i>Pseudallescheria</i> species	1.8%	4->16	0.5-4	2->16	0.5->16
<i>Lomentospora prolificans</i>	Previously <i>Scedosporium prolificans</i> - Highly resistant	12.3%	8->16	4->16	8->16	>16

Lackner et al. *Antimicrob Agents Chemother* 2012;56:2635-2642. FTL internal data (N = 285; 9/12 through 8/18)

Fusarium Species & Fusariosis

Significant cause of morbidity and mortality in immunocompromised hosts

- Invasive & disseminated infections (high mortality in neutropenic hosts)
- Keratitis & onychomycosis in immunocompetent hosts
 - Fungal keratitis in contact lenses
 - Peritonitis in peritoneal dialysis patients

Human infections can be caused by 8 species complexes

- *Fusarium solani* spp. complex (FSSC)
- *Fusarium oxysporum* spp. complex (FOSC)
- *Fusarium fujikuroi* spp. complex (FFSC)
- *Fusarium chlamydosporum* spp. complex (FCSC)
- *Fusarium dimerum* spp. complex (FDSC)
- *Fusarium incarnatum-equiseti* spp. complex (FIESC)
- *Fusarium sambucinum* spp. complex (FSAMSC)
- *Fusarium tricinctum* spp. complex (FTSC)

Campo et al. *J Infect* 2010;60:331-337. Kerr et al. *Ann Intern Med* 1983;99:334-336.
O'Donnell et al. *J Clin Microbiol* 2007;45:2235-2248.

Fusarium solani Species Complex

Species	Haplotype	Proposed Name
<i>F. petrophilum</i>	haplotype 1	<i>Neocosmospora petrophila</i>
<i>F. keratoplasticum</i>	haplotype 2	<i>Neocosmospora keratoplastica</i>
<i>F. falciforme</i>	haplotype 3+4	<i>Neocosmospora falciforme</i>
<i>F. solani</i>	haplotype 5	<i>Neocosmospora solani</i>
<i>F. metavorans</i>	haplotype 6	<i>Neocosmospora metavorans</i>
FSSC 7	haplotype 7	<i>Neocosmospora gamsii</i>
<i>F. lichenicola</i>	haplotype 16	<i>Neocosmospora lichenicola</i>
FSSC 20	haplotype 20	<i>Neocosmospora suttoniana</i>
FSSC 43	haplotype 43	<i>Neocosmospora catenata</i>

Genus *Neocosmospora* (FSSC) contains significant plant pathogen species, important agents of veterinary infections, and clinically relevant species that cause infections in immunocompromised hosts.

Lombard et al. *Stud Mycol* 2015;80:189-245. Sandoval-Denis & Crous. *Persoonia* 2018;41:109-129.

Fusarium & Antifungal Susceptibility

Species	Espinel-Ingroff AAC 2015 MIC Ranges (Mode)				Pfaller et al. AAC 2018
	Amphotericin	Itraconazole	Posaconazole	Voriconazole	Isavuconazole
<i>F. solani</i> SC	≤ 0.25 - 16 (2)	0.5 - >16 (16)	1 - >16 (8)	0.5 - >16 (8)	>4
<i>F. oxysporum</i> SC	≤ 0.25 - 16 (2)	1 - >16 (16)	0.5 - >16 (2)	0.5 - >16 (4)	---

Limited in vitro activity of available antifungals against *Fusarium*
Clinical relevance in vitro susceptibility testing?

<i>F. fujikuroi</i> SC	0.5 - 2 (1)	≥16 (16)	0.5 - 4 (2)	2 - 16 (4)	---
<i>Fusarium</i> spp.	---	---	---	---	>4 (8->16; >16)*

Espinel-Ingroff et al. *Antimicrob Agents Chemother* 2015;1079-1084.

Pfaller et al. *Antimicrob Agents Chemother* 2018 (Epub ahead of print). *FTL internal data

Things to Remember...

- Low MIC *does not* predict successful outcomes
- High MIC *does not* predict failure
 - *In vitro* resistance may predict a population less likely to respond
- Other factors are important predictors of outcomes
 - *Host immune status & other comorbidities*
 - *Site of infection*
 - *Time to diagnosis/start of treatment*
 - *Drug concentrations at site of infection*
 - *Drug-drug & drug-disease interactions*

Both antibacterial AND antifungal susceptibility testing share these limitations

Antifungal Susceptibility Moulds Challenges

- No automation for mould susceptibility testing
 - No FDA-cleared panels
 - Some labs use YeastOne Sensititre plates (*Aspergillus*)
 - Utility against other moulds?
- Few clinical laboratories in U.S. perform susceptibility testing against moulds
 - Sendout lab (delays in results)
- Clinical breakpoints not established against most moulds
 - Limited breakpoints against *Candida* species
 - EUCAST breakpoints against some *Aspergillus* spp.

Epidemiologic Cut-Off Values


ECV or ECoff

- **ECVs** separate wild-type isolates from non-wild type isolates
 - Encompass ~95 - 97.5% of isolates in wild type MIC distribution
 - Sensitive measure of emergence of resistance
- **Clinical breakpoints** indicate isolates likely to respond to treatment with a given antimicrobial agent
 - Established using PK/PD, ECV & MIC distributions, clinical data

Genus	Antifungals	CLSI, EUCAST, or Other
<i>Candida</i> (not all species)	Amphotericin, Azoles, Echinocandins (AFG & MFG)	CLSI M59, EUCAST
<i>C. neoformans</i> & <i>C. gatti</i> (certain genotypes)	Amphotericin, Azoles, Flucytosine	CLSI M59, EUCAST
<i>Aspergillus</i> (not all species)	Amphotericin, Azoles, Caspofungin	CLSI M59, EUCAST
<i>Fusarium</i> (by SC)	Amphotericin, Azoles (not Isavuconazole)	Espinel-Ingroff AAC 2015

Limited number ECVs established against moulds


Turnidge and Johnson. *Clin Microbiol Rev* 2007;20:391-408. Pfaller et al. *Drug Resist Updat* 2011;14:164-76.



**Antimicrobial Agents
and Chemotherapy**

MECHANISMS OF RESISTANCE

2018:62:e02599-17.



2 of 3 *S. apiospermum*-specific amino acid changes (Y136F & G464S) corresponded to mutations in *A. fumigatus* CYP51A (Y121F & G448S) linked to azole resistance

Identification of 14- α -Lanosterol Demethylase (CYP51) in *Scedosporium* Species

Anne Bernhardt,^a Wieland Meyer,^{b,c} Volker Rickerts,^a Toni Aebischer,^a Kathrin Tintelnot^a

MICROBIAL DRUG RESISTANCE
Volume 24, Number 6, 2018
© Mary Ann Liebert, Inc.
DOI: 10.1089/mdr.2017.0311

Increased expression of CYP51A observed following exposure to voriconazole, posaconazole, & tebuconazole, but not amphotericin B.

Overinduction of the *CYP51A* Gene After Exposure to Azole Antifungals Provides a First Clue to the Resistance Mechanism in the *Fusarium solani* Species Complex

Mellado et al
Chamilos &
Slaven et al.

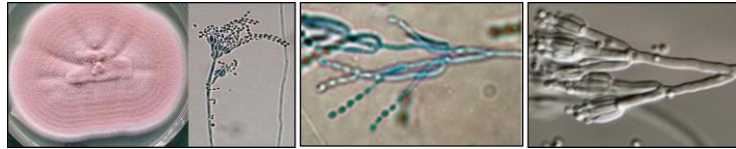
Maurine D'Agostino,¹ Thomas Lemmet,¹ Claire Dufay,¹ Amandine Luc,²
Jean Pol Frippiat,¹ Marie Machouart,^{1,3} and Anne Debourgogne^{1,3}

92-2097.
. PLoS

Pathog 2008;4:e1000200; Camps et al. *PLoS One* 2012;7:e50034; Ammar et al. *PLoS One* 2013;8:e79042; Crowley et al. *J Infect Dis* 2017;216:S436-444.

Other Moulds & Resistance

Species	Reduced Susceptibility/Resistance
<i>Purpureocillium lilacinum</i> (formerly <i>Paecilomyces lilacinus</i>)	Amphotericin B (MIC >16) Low voriconazole MICs
<i>Paecilomyces variotii</i>	Voriconazole & Isavuconazole (MIC 4->16) Low Amphotericin B MICs
Various <i>Penicillium</i> spp. <ul style="list-style-type: none"> • <i>P. citrinum</i> • <i>P. steckii</i> • <i>P. sumatraense</i> (plus others)	Voriconazole (MIC >16) & Sometimes Itraconazole/Posaconazole (variable; MICs can range 4 - >16)



Rasamsonia

Intrinsic Voriconazole Resistance

- Thermotolerant and thermophilic *Penicillium*-like fungi
 - *Rasamsonia argillacea* species complex
 - *R. argillacea*
 - *R. eburnean*
 - *R. piperina*
 - *R. aegroticola*
- Previously known as *Geosmithia*
Often misidentified by morphology as
Penicillium or *Paecilomyces*
- Emerging pathogens reported to cause invasive disease in CGD & patients with hematologic malignancies
 - Chronic colonization in CF patients
 - No clear correlation between colonization & lung deterioration

Houbraken et al. *J Clin Microbiol* 2013;51:22-30.

Fatal Disseminated *R. aegroticola* Infection & Antifungal Susceptibility

- 21 year old CF patient who developed disseminated invasive fungal infection post-lung transplantation
- Cultures pre-transplant: MRSA, *A. fumigatus*, *P. variotii*
- Voriconazole part of antifungal prophylaxis/treatment
 - Days 2 – 100; trough levels 2-5 µg/mL
- *R. aegroticola* cultured from BAL collected pre-mortem & tissue collected at autopsy
 - DNA sequence analysis (previous sputum cultures mis-identified by morphology as *P. variotii*)



Species	Isavuconazole	Itraconazole	Voriconazole	Posaconazole	Amphotericin	Micafungin
<i>R. argillacea</i>	>32	1->16	>32	1->16	0.5-32	0.125-0.25
<i>R. piperina</i>	>32	0.5-2	>32	0.5-1	1	0.03-0.25
<i>R. aegroticola</i>	>32	2->16	>32	1-4	1-8	0.125-0.25
<i>R. aegroticola</i> (pt. isolate)	NT	2	>16	1	2	≤ 0.015

Hong et al. *J Cyst Fibros* 2017;16:e3-7. Steinmann et al. *Antimicrob Agents Chemother* 2016;60:6890-6891.

Antifungals in Development

Aspergillus (Cryptic spp.) *Scedosporium/L. prolificans* Mucorales *Fusarium* Other Moulds

Olorofim

Rezafungin

APX001
(formerly E1210)

APX001(?)

SCY-078

APX001

Olorofim

VT-1598

Olorofim

VT-1598
(*Rhizopus*)

VL-2397

Summary

- Increased recognition of antifungal resistance in moulds
 - *Aspergillus fumigatus* & other *Aspergillus* species
 - *Fusarium*, *Scedosporium* & Mucorales (and others)
- Increased recognition of cryptic species
 - Reduced susceptibility to clinically available agents
- Understanding of clinical significance lags behind microbiologic data
 - Other factors of importance
 - Host response (or lack thereof)
 - Concentrations/exposures achieved at sites of infection