



Evolving fungal landscape in Asia

Professor Yee-Chun Chen

Professor of Medicine

National Taiwan University Hospital and College of Medicine;

Investigator, National Institute of Infectious Diseases and Vaccinology

National Health Research Institutes, Taiwan



The Evolving Fungal Landscape in Asia

Yee-Chun Chen, M.D., PhD.

Department of Medicine, National Taiwan University Hospital and College of Medicine; National Institute of Infectious Diseases and Vaccinology, National Health Research Institutes, Taiwan



Outlines

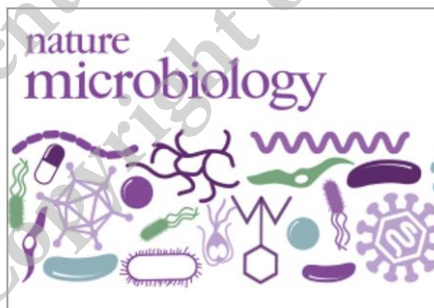
- Stop neglecting fungi
- Emerging: *Candida auris*
- Unexpected: fungal infections after natural disasters
- One health: azole-resistant *Aspergillus fumigatus*
- Conclusion

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editorial

Stop neglecting fungi



Fungal pathogens are virtually ignored by the press, the public and funding bodies, despite posing a significant threat to public health, food biosecurity and biodiversity.

<https://www.nature.com/articles/nmicrobiol2017120>

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Stop neglecting fungi

Indeed, in comparison to the threat from drug-resistant bacterial infections or viral outbreaks, diseases caused by fungi, fungal drug resistance and the development of new antifungal therapeutics gets little coverage. Yet in this case, no news is certainly not good news, and the disparity relative to other infectious disease agents unjustified.



Few realize that over 300 million people suffer from serious fungal-related diseases, or that fungi collectively kill over 1.6 million people annually, which is more than malaria and similar to the tuberculosis death toll.

The Burden of Fungal Disease (LIFE, 2017); <http://go.nature.com/2sMKpuN>
 Burden of HIV-associated histoplasmosis compared with tuberculosis in Latin America: a modelling study.
 Lancet Infect Dis 2018; 18: 1150

The overall burden of fungal diseases is challenging to quantify, because they are likely substantially underdiagnosed.

Kaitlin Benedict K et al., Clin Infect Dis 2018; cy776,

Disease (most common species)	Location	Estimated life-threatening infections/ year at that location*	Mortality rates (% in infected populations)*
Opportunistic invasive mycoses			
Aspergillosis (<i>Aspergillus fumigatus</i>)	Worldwide	>200,000	30–95
Candidiasis (<i>Candida albicans</i>)	Worldwide	>400,000	46–75
Cryptococcosis (<i>Cryptococcus neoformans</i>)	Worldwide	>1,000,000	20–70
Mucormycosis (<i>Rhizopus oryzae</i>)	Worldwide	>10,000	30–90
Pneumocystis (<i>Pneumocystis jirovecii</i>)	Worldwide	>400,000	20–80
Endemic dimorphic mycoses*			
Blastomycosis (<i>Blastomyces dermatitidis</i>)	Midwestern and Atlantic United States	~3,000	<2–68
Coccidioidomycosis (<i>Coccidioides immitis</i>)	Southwestern United States	~25,000	<1–70
Histoplasmosis (<i>Histoplasma capsulatum</i>)	Midwestern United States	~25,000	28–50
Paracoccidioidomycosis (<i>Paracoccidioides brasiliensis</i>)	Brazil	~4,000	5–27
Penicilliosis (<i>Penicillium marneffe</i>)	Southeast Asia	>8,000	2–75

*Most of these figures are estimates based on available data, and the logic behind these estimates can be found in the text and in the Supplementary Materials. †Endemic dimorphic mycoses can occur at many locations throughout the world. However, data for most of those locations are severely limited. For these mycoses, we have estimated the infections per year and the mortality at a specific location, where the most data are available.

Impact of local epidemiology on global health:

Importation through travel, returned immigrants, global trade
International healthcare, solid organ transplantation, etc.

Brown GD, et al. Sci Transl Med 2012;4;

More updated data: Bongomin F, et al. J Infect 2017;3:57

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World Health
Organization

Current WHO Initiatives on fungal infection

- HIV Department have recommendation for screening, treatment, prevention of:
 - *Pneumocystis pneumonia*
 - *Cryptococcus neoformans*
 - *Candida* (thrush)
- Neglected tropical disease
 - Mycetoma, 2016
 - Chromoblastomycosis, 2017
- Antimicrobial resistance
 - Surveillance of bloodstream infection due to *Candida* spp, 2018

<https://www.who.int/glass/events/AMR-in-invasive-candida-infections-meeting/en/>

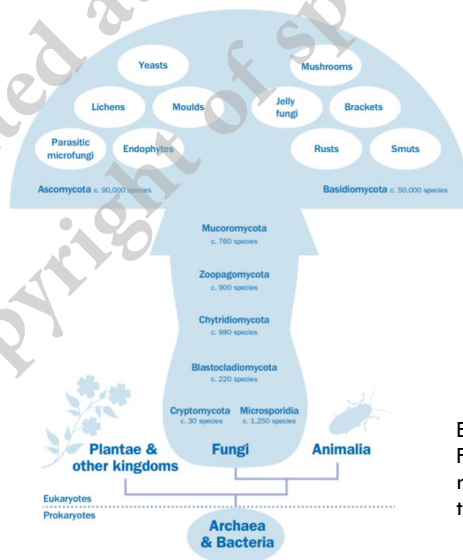
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Unmet Medical Needs

- Increased incidences of invasive fungal diseases in developed countries due to higher survival of susceptible populations
- Remained high mortality/morbidity
- Existing treatment options are limited
 - few antifungal families/targets of action
 - efficacies vary depending on the infecting species
 - pharmacokinetic and –dynamic considerations
- Emergence of antifungal resistance

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Kingdom Fungi in the Tree of Life



Evolutionary studies have shown that Fungi and Animalia are more closely related to each other than to any of the other kingdoms of life.

https://stateoftheworldfungi.org/2018/reports/SOTWFungi_2018_Full_Report.pdf

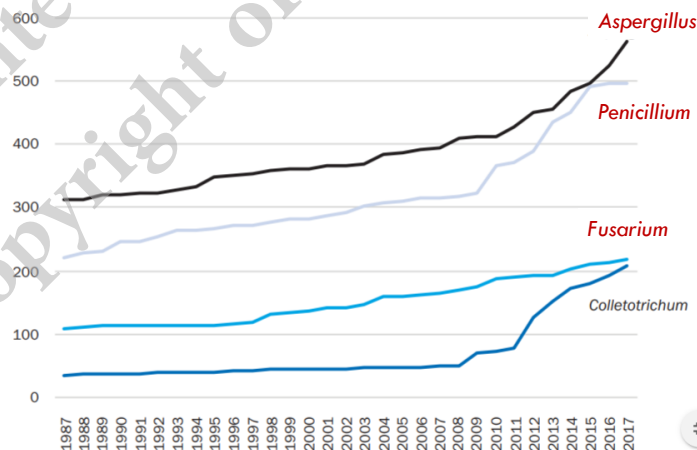
Global Warming Will Bring New Fungal Diseases for Mammals

- The estimated number of fungal species is 1.5 million, occupying a broad variety of habitats.
- About 300 fungal species are reported to be pathogenic to humans. These fungi have the ability to survive and grow at the high body temperatures of endothermic animals.
- Global warming may increase the prevalence of fungal disease in humans as fungi adapt to survival in warmer temperatures.

Garcia-Solache, M. A., and A. Casadevall. 2010. mBio 1(1):e00061

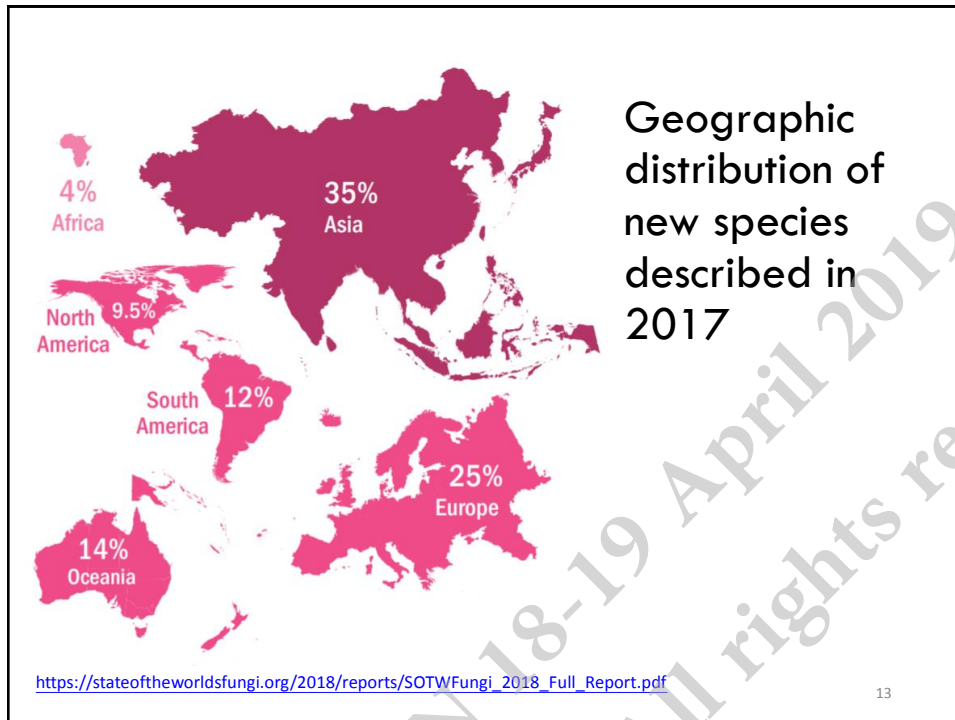
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Numbers of accepted species for four economically important fungal genera



https://stateoftheworldfungi.org/2018/reports/SOTWFungi_2018_Full_Report.pdf

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A deadly frog-killing fungus probably originated in East Asia



RISKY TRAVELER A genetic study now traces an amphibian-killing fungus to East Asia, from where international trade in such showy pets as this oriental fire-bellied toad could have helped launch the pathogen worldwide.

<https://www.sciencenews.org/article/deadly-frog-killing-bd-fungus-probably-originated-east-asia>

S.J. O'Hanlon et al. [Recent Asian origin of chytrid fungi causing global amphibian declines](#). *Science* 2018;360:621.

K. Lips. [The hidden biodiversity of amphibian pathogens](#). *Science* 2018;360:604.

Emerging: *Candida auris*

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The New York Times

April 6, 2019

DEADLY GERMS, LOST CURES **A Mysterious Infection, Spanning the Globe in a Climate of Secrecy**

> 587 *C. auris* infections since 2013.

Recently *C. auris* reached New York (N=309), New Jersey (N=104) and Illinois (N=144), leading the CDC, USA, to add it to a list of germs deemed “urgent threats.”

<https://www.nytimes.com/2019/04/06/health/drug-resistant-candida-auris.html>

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2015-2016

Intensive care unit closed after new deadly superbug emerges in the UK

2009/1996

CANDIDA AURIS
A Dangerous 'Superbug' Fungus Outbreak

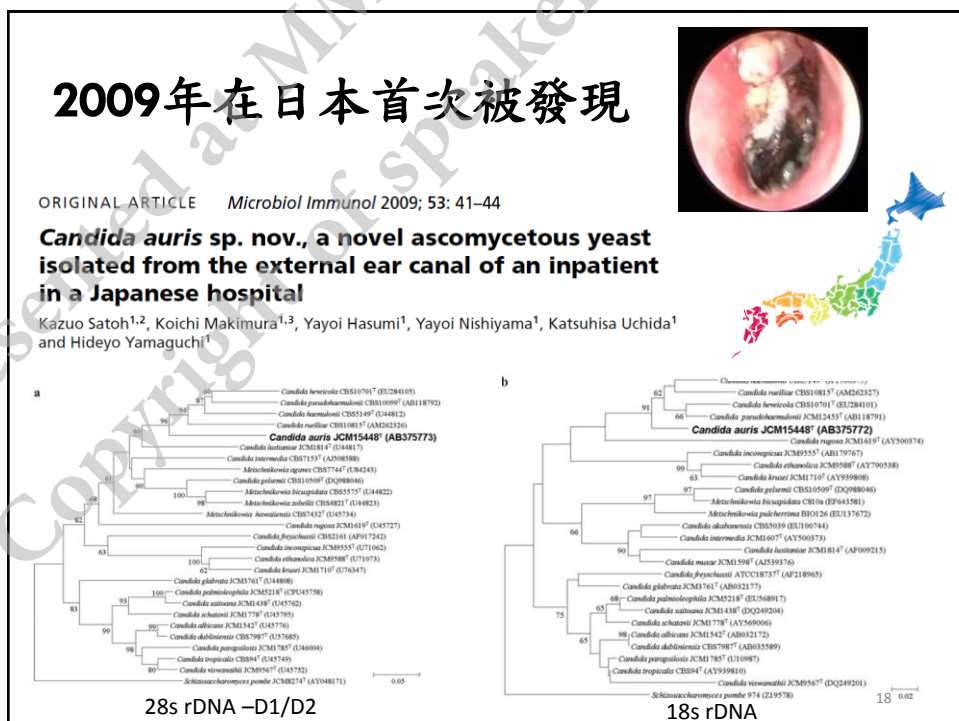
Dr. Axe

<https://draxe.com/candida-auris/>

傳播媒體新議題
April 12, 2019

New York Times: Superbug C. auris identified in 617 people, CDC says

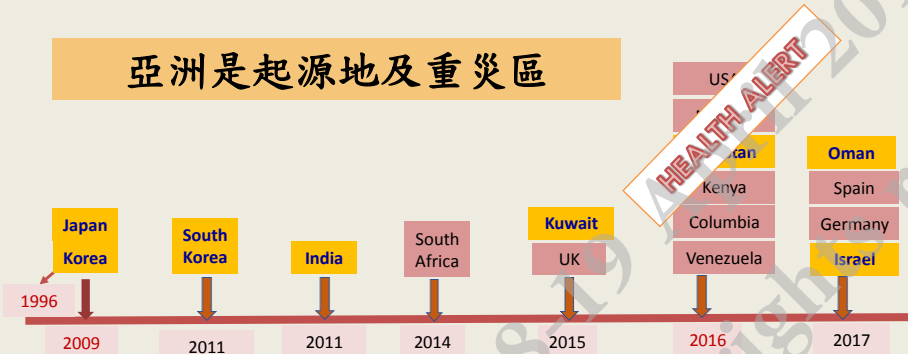
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新興病菌成為全球危機

最早已知的菌株來自於1996年的南韓，已於歐、亞、美、非二十餘國迅速傳播並造成感染及重症死亡

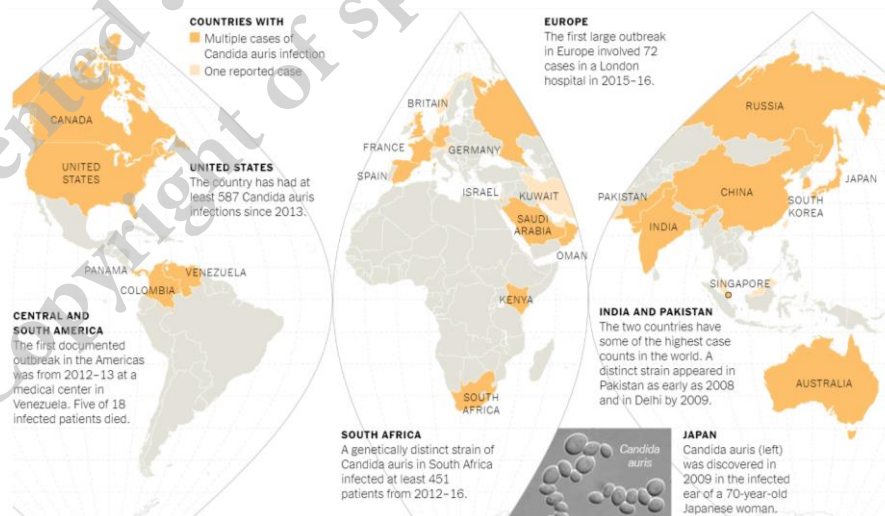
亞洲是起源地及重災區



Chakrabarti A, et al. Intensive Care Med 2015; 41: 285
Chowdhary A, et al. J Hosp Infect 2016; 94: 209

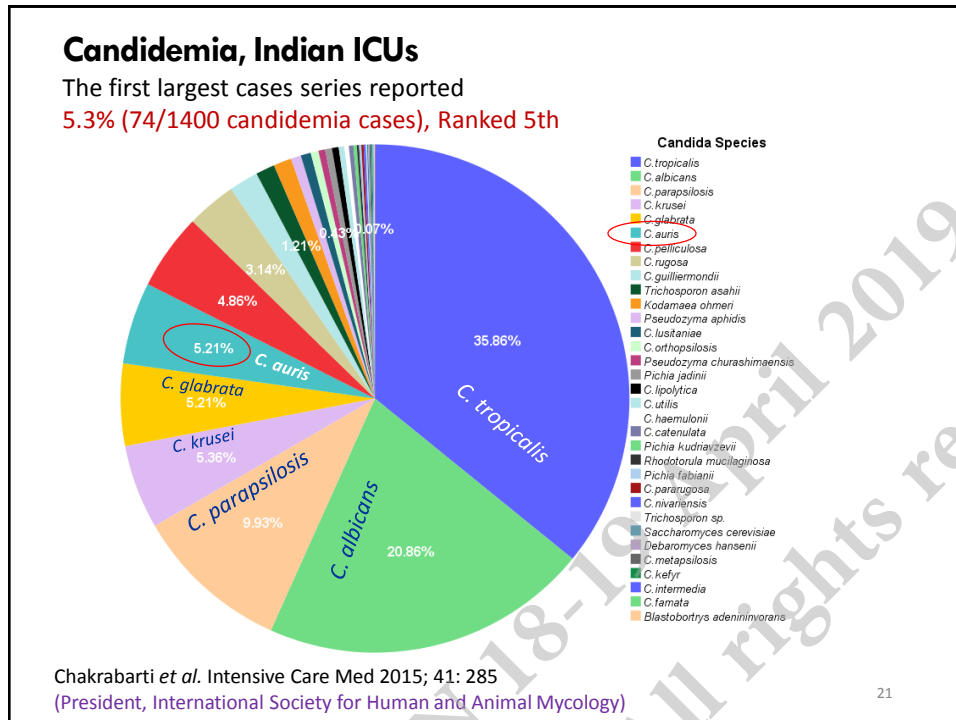
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Countries from which *C. auris* cases have been reported



<https://www.nytimes.com/2019/04/06/health/drug-resistant-candida-auris.html>

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Unique features from largest series in Indian ICUs

- Significant risk factors in Indian ICUs

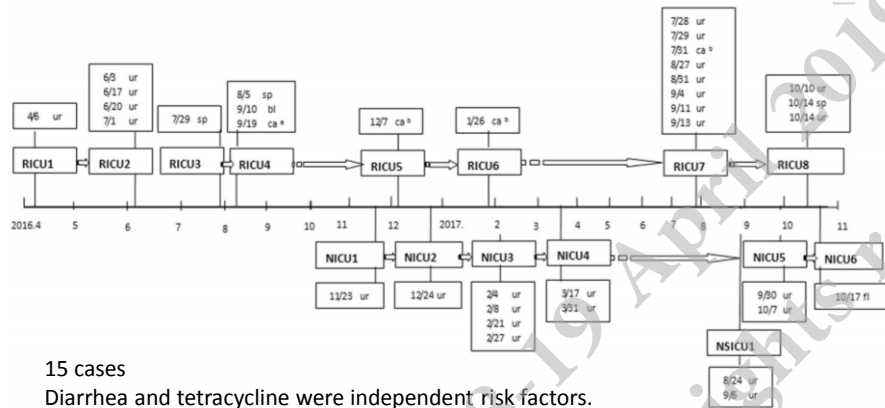
1. prior antifungal exposure ($P < 0.001$)
2. underlying respiratory illness ($P < 0.002$)
3. vascular surgery ($P < 0.048$)
4. multiple interventions ($P < 0.007$)
5. public-sector hospital ($P < 0.006$)

**Patients with sepsis,
undergoing
invasive
management for
longer periods &
exposed to
antifungal agents**

Rudramurthy S, et al. J Antimicrob Chemother 2017; 72: 1794

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First cases of *C. auris* infection or colonization, Shenyang, China



15 cases

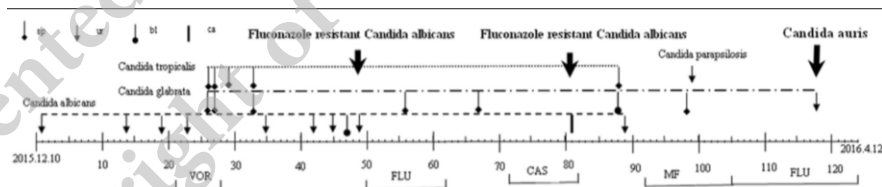
Diarrhea and tetracycline were independent risk factors.

Tian S, et al. Emerg Infect Dis 2018;7:128

RICU, Respiratory ICU; NICU, neuroICU; NSICU, neurosurgery ICU
Sp, sputum; ur, urine; bl, blood; ca, urinary catheter

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First case, Shenyang, China



Tian S, et al. Emerg Infect Dis 2018;7:128

Vor, voriconazole; FLU, fluconazole; CAS, caspofungin; MF, micafungin;
Sp, sputum; ur, urine; bl, blood; ca, urinary catheter

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Available online at www.sciencedirect.com

ScienceDirect

journal homepage: www.jfma-online.com

Review Article

Are we ready for the global emergence of multidrug-resistant *Candida auris* in Taiwan?



Po-Liang Lu^{a,b,c}, Wei-Lun Liu^{d,e}, Hsiu-Jung Lo^{f,g},
Fu-Der Wang^{h,i}, Wen-Chien Ko^{j,k}, Po-Ren Hsueh^{l,m},
Mao-Wang Hoⁿ, Chun-Eng Liu^o, Yen-Hsu Chen^{a,b,c},
Yee-Chun Chen^{f,m,*}, Yin-Ching Chuang^p, Shan-Chwen Chang^m

^a Division of Infectious Diseases, Department of Internal Medicine, Kaohsiung Medical University Hospital, Kaohsiung, Taiwan

^b School of Medicine, College of Medicine, Kaohsiung Medical University, Kaohsiung, Taiwan

^c Department of Biological Science and Technology, College of Biological Science and Technology,

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Prevalence of *C. auris* among 5064 clinical isolates based on multicenter surveillance in Taiwan

Investigator(s)	Source of isolates	Specimens types	Year	Results
HJ Lo	TSARY National surveillance ^b	Randomly collected <i>Candida</i> clinical isolates (1999) or <i>Candida</i> isolates from sterile sites and non-sterile sites (2002, 2006, 2010, and 2014)	1999 2002 2006 2010 2014	0/660 0/945 0/1015 0/1130 0/1168
WL Liu, FD Wang, MW Ho, YH Chen, CE Liu	CMMC, VGH-TPE, CMUH, KMH, CCH	Blood isolates, hospital wide, rare <i>Candida</i> species ^c	January 2011–June 2014	0/52 ^d
YC Chen, PR Hsueh	NTUH	Blood isolates, hospital wide, rare <i>Candida</i> species ^c	2011–2016	0/57 ^d
WL Liu	CMMC, Liouying campus	Blood isolates, hospital wide, rare <i>Candida</i> species ^c	2007–2014	0/21 ^{d 37}
MC Li WC Ko	NCKUH	Blood isolates, hospital wide, rare <i>Candida</i> species ^c	2011–2016	0/37

Abbreviation: TSARY, Taiwan Surveillance of Antimicrobial Resistance of Yeasts; CMMC: Chi Mei Medical Center; VGH-TPE: Taipei Veterans General Hospital; CMUH: China Medical University Hospital; KMH: Kaohsiung Medical University Hospital; CCH: Changhua Christian Hospital; NTUH, National Taiwan University Hospital; NCKUH, National Cheng Kung University Hospital.

^a Data from personal communication with the principal investigators at each hospital or research site. These data are generated based on DNA sequencing of the internal transcribed spacer regions of the nuclear rRNA gene operon and the D1/D2 domain of the large ribosomal subunit of 26S rDNA.

^b Multicenter in different geographic location of Taiwan.³⁶

^c *Candida* species other than *C. albicans*, *C. tropicalis*, *C. parapsilosis*, *C. glabrata*, and *C. krusei*.

^d One isolate per patient.

Lu P-L, et al., Are we ready for the global emergence of multidrug-resistant *Candida auris* in Taiwan?, Journal of the Formosan Medical Association (2017), <https://doi.org/10.1016/j.jfma.2017.10.005>

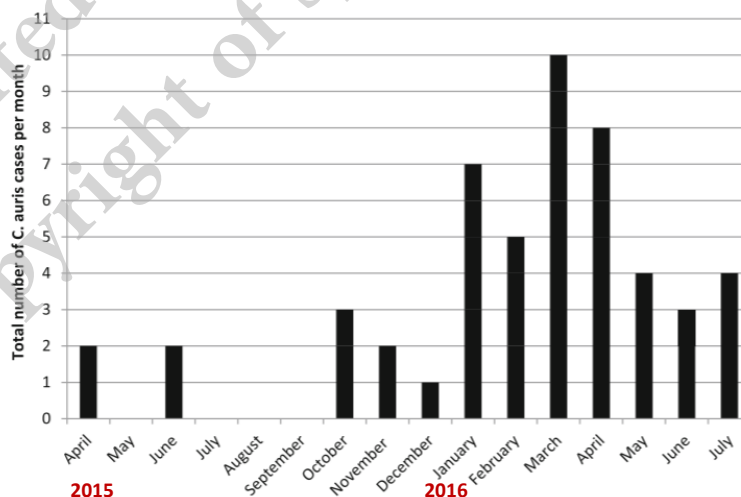
The first clinical isolate of *C. auris* in Taiwan

- A 55-year-old man with a medical history of diabetes mellitus and pemphigus vulgaris underwent treatment with azathioprine and prednisolone. Between November 9, 2017 and December 30, 2017, he was hospitalized at CMMC due to pemphigus vulgaris-related skin and soft tissue infection caused by MRSA, and received anti-MRSA treatment.
- On April 11, 2018, several ruptured vesicles with erythematous changes and purulent discharge over the face were noted. Bacterial cultures were obtained.
- *Candida* along with MRSA were recovered from BAP. This isolate did not grow on Mycosel agar containing cycloheximide (400 mg/L) in the culture media that is routinely used for primary isolation of fungi in the laboratory.

Tan HJ, et al. International Journal of Antimicrobial Agents (accepted)

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First hospital outbreak *C. auris* in a European hospital



Antimicrob Resist Infect Control 2016;5:35.

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By June 2016, the hospital had seen 72 cases of *C. auris*, and decided to shut down its intensive care unit for 11 days to address the contamination



Royal Brompton Hospital near London, UK - a National Health specialist center for cardio-thoracic surgery with 296 beds that draws wealthy patients from the Middle East and around Europe.

<https://www.nytimes.com/2019/04/06/health/drug-resistant-candida-auris.html>

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THE NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

A *Candida auris* Outbreak and Its Control in an Intensive Care Setting

- Oxford University Hospitals, UK; Feb 2015 and Aug 2017
- 70 patients with *C. auris* colonization/infection (7 patients, 10%)
- 94% admitted to the **neuroICU** before diagnosis
- Predictors of *C. auris* colonization or infection (multivariate analysis)
 - The use of **reusable skin-surface axillary temperature probes** (odds ratio, 6.80, $P < 0.001$)
 - Systemic fluconazole exposure (odds ratio, 10.34, $P = 0.01$)
- No attributable mortality

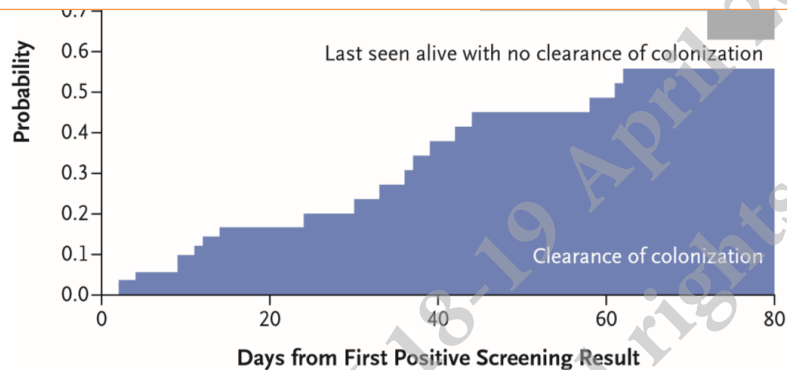
N Engl J Med 2018;379:1322

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Duration of *C. auris* carriage

The median duration of carriage among patients remaining alive was

- 61 days when two consecutive negative screening results were used to define clearance of colonization
- 82 days when three consecutive negative results were used

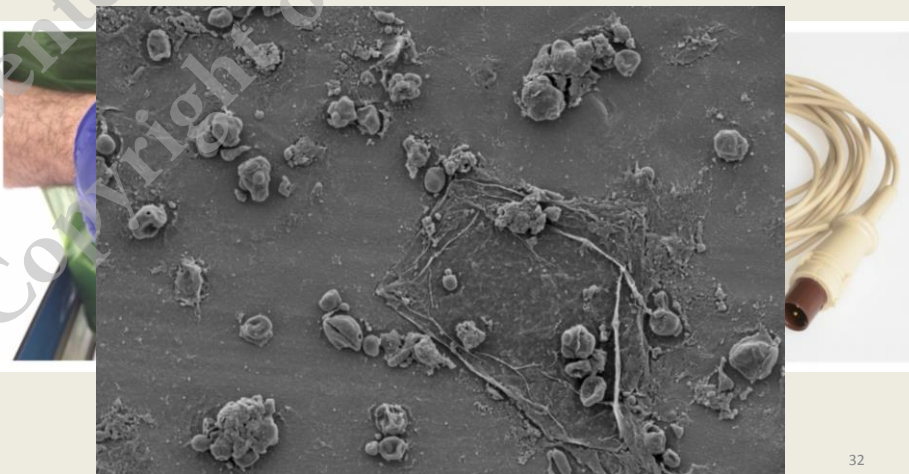


N Engl J Med 2018;379:1322

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Skin surface temperature probes (axillary), neurolCU, Oxford

Used routinely in ventilated patients for continuous temperature monitoring



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Number of *C. auris* clinical cases

- 0
- 1
- 2-10
- 11-10
- 51-100
- 101 or more

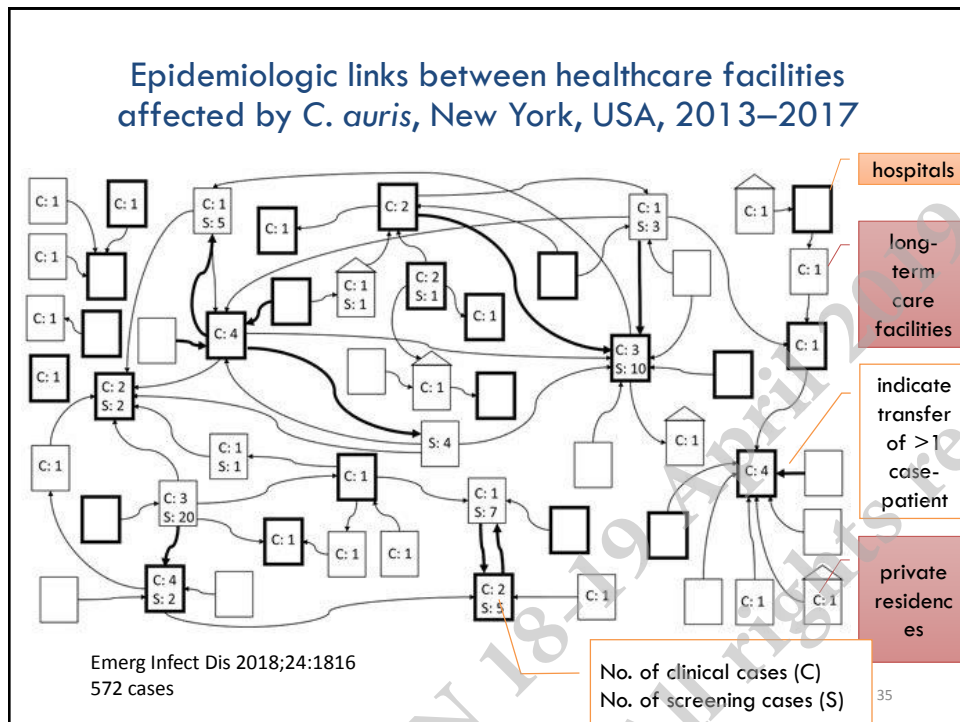
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Single *C. auris* case reported

Transmission or multiple cases of *C. auris* reported

U.S. *C. auris* cases linked to healthcare stays in these countries

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Environmental contamination with *C. auris* in healthcare facilities, New York, USA

Category, object or surface	No. samples negative by culture & PCR/No. samples evaluated (%)
Near-patient surfaces and objects in rooms	145/178 (82)
Other surfaces and objects in rooms	163/187 (87)
Equipment in room	30/35 (86)
Equipment outside of room	243/260 (94)

Emerg Infect Dis 2018;24:1816

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Everything was positive

Multidrug resistance

Environmental
Contamination

Not limited in the
acute-care hospitals

Persistent carriage

N Engl J Med 2018;379:1322; Emerg Infect Dis 2018;24:1816

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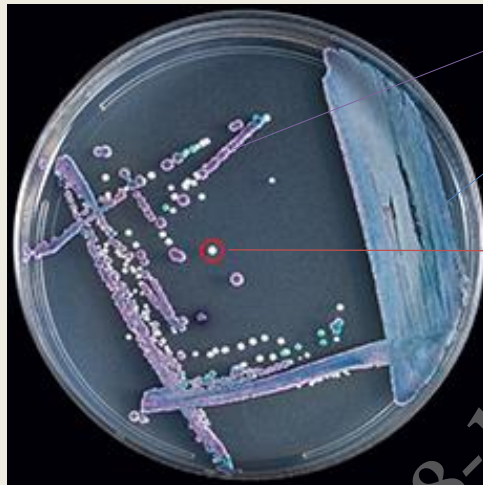
Common misidentifications based on the identification method used

Identification Method	Organism <i>C. auris</i> can be misidentified as
Vitek 2 YST	<i>Candida haemulonii</i> <i>Candida duobushaemulonii</i>
API 20C	<i>Rhodotorula glutinis</i> (characteristic red color not present) <i>Candida sake</i>
BD Phoenix yeast identification system	<i>Candida haemulonii</i> <i>Candida catenulata</i>
MicroScan	<i>Candida famata</i> <i>Candida guilliermondii</i> [*] <i>Candida lusitanae</i> [*] <i>Candida parapsilosis</i> [*]
RapID Yeast Plus	<i>Candida parapsilosis</i> [*]

^{*}*C. guilliermondii*, *C. lusitanae*, and *C. parapsilosis* generally make pseudohyphae on cornmeal agar. If hyphae or pseudohyphae are not present on cornmeal agar, this should raise suspicion for *C. auris* as *C. auris* typically does not make hyphae or pseudohyphae. However, some *C. auris* isolates have formed hyphae or pseudohyphae. Therefore, it would be prudent to consider any *C. guilliermondii*, *C. lusitanae*, and *C. parapsilosis* isolates identified on MicroScan or any *C. parapsilosis* isolates identified on RapID Yeast Plus as possible *C. auris* isolates and forward them for further identification. <https://www.cdc.gov/fungal/candida-auris/recommendations.html>

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Mixed culture on CHROMagar Candida



Candida glabrata
(purple)

Candida tropicalis
(navy blue)

Candida auris
(white)

<https://www.cdc.gov/fungal/candida-auris/recommendations.html>

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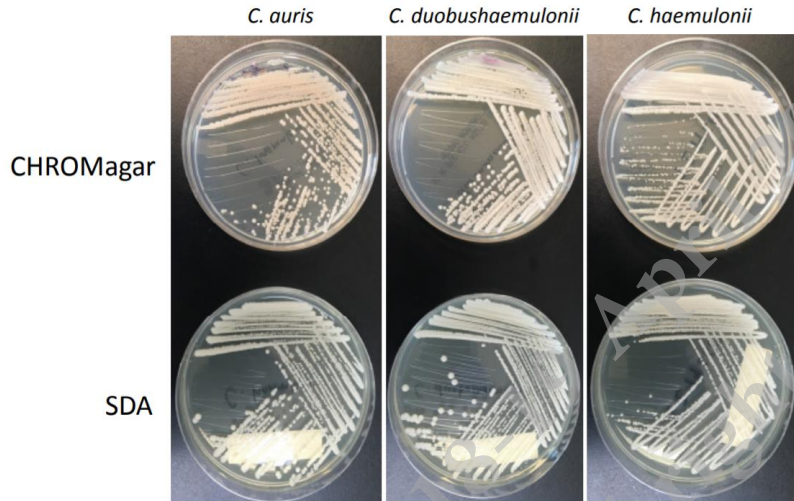
Candida auris on CHROMagar Candida, displaying multiple color morphs



<https://www.cdc.gov/fungal/candida-auris/recommendations.html>

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Like most other non-albicans *Candida* species, *C. auris* isolates are germ tube test negative and colonies appear pale purple or pink on CHROMagar *Candida* medium.



http://www.chromagar.com/fichiers/1526603271AMMI_C_auris_Identification.pdf?PHPSESSID=0541bde3c8c5cdb9fd5caa84968aeabe

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The commercially available biochemical test kits and MALDI-TOF instruments used in clinical laboratories for the identification of yeast isolates may lack *C. auris* in their database, these assays may misidentify *C. auris*.

Each box represents one testing site:



MALDI-TOD MS +
update library

http://www.chromagar.com/fichiers/1526603271AMMI_C_auris_Identification.pdf?PHPSESSID=0541bde3c8c5cdb9fd5caa84968aeabe

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Algorithm to identify *Candida auris* based on phenotypic laboratory method and initial species identification

PURPOSE

Candida auris is a multidrug-resistant yeast that has been found in multiple countries, including the United States. *C. auris* can cause invasive infections, be passed from person to person, and persist in the environment. Its severity, communicability, and drug resistance makes correctly identifying *C. auris* crucial to treating patients and preventing infections. However, this is challenging because traditional phenotypic methods frequently misidentify *C. auris*. This algorithm details the steps needed to determine the correct *Candida* spp. based on the tests and equipment available in your lab.

TABLE OF CONTENTS – ALGORITHMS BY METHOD

1. Bruker Biotyper MALDI-TOF
2. bioMérieux VITEK MS MALDI-TOF
3. VITEK 2 YST
4. API 20C
5. BD Phoenix
6. MicroScan
7. RapID Yeast Plus
8. Summary of this algorithm in table form

Please note that these algorithms are based on our current knowledge about misidentification of *C. auris* and may change as we learn new information.

<https://www.cdc.gov/fungal/diseases/candidiasis/pdf/Testing-algorithm-by-Method-temp.pdf>

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C. auris - an emerging fungus that presents a serious global health threat



<https://www.cdc.gov/fungal/diseases/candidiasis/candida-auris-qanda.html>

Unique features

1. Is often multidrug-resistant
2. Is difficult to identify
3. Has caused outbreaks in healthcare settings (delayed diagnosis, prolonged carriage, environment contamination)



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Unexpected

Invasive fungal infections after natural disasters

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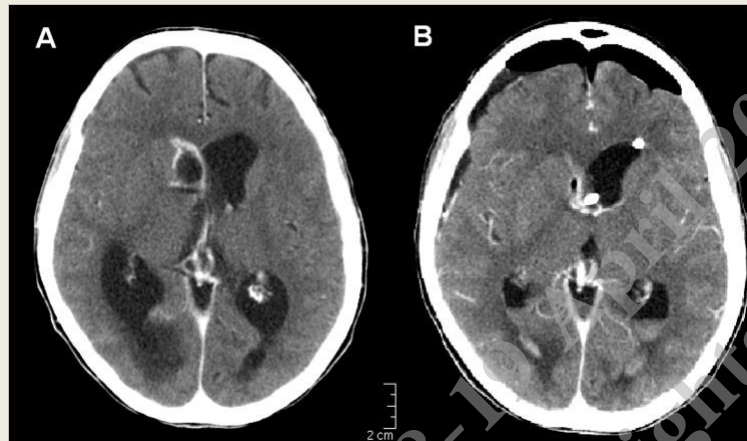
Necrotizing fasciitis caused by *Apophysomyces elegans* complicating soft-tissue and pelvic injuries in an Indian Ocean Tsunami survivor, Thailand



42 days liposomal amphotericin B

Snell & Yavakoli Plastic Reconstruct Surg 2007; 119: 448-449

Brain abscess caused by *Scedosporium apiospermum* in a tsunami survivor



Garzoni et al. Emerg Infect Dis 2005;11:1591

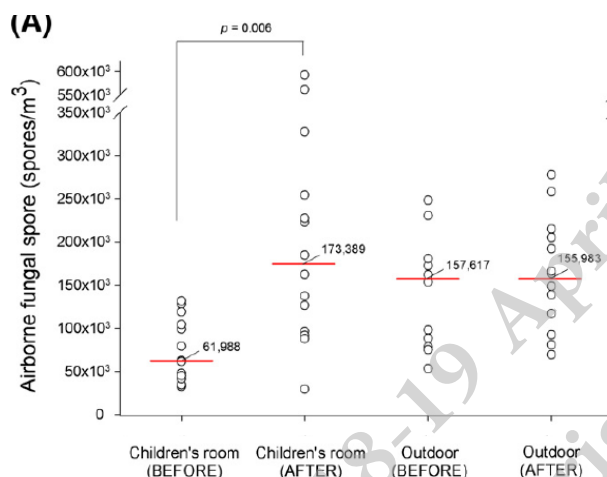
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Aspergillus meningitis after the administration of spinal anesthesia for cesarean section after tsunami, Sri Lanka

Characteristic or Test	Patient No.				
	1	2	3	4	5
Clinical features					
Age — yr	26	21	27	29	38
Date of exposure	June 21	June 22	June 20	June 22	July 17
Date of onset of symptoms	July 3	July 2	July 5	July 3	July 25
Neck stiffness and Kernig's sign	Absent	+++	++	+	+
Lateral rectus palsy	Absent	Absent	Bilateral	Bilateral	Absent
Stroke	Thalamic infarction on July 9	Thalamic infarction on July 31	Absent	Absent	Ventricular hemorrhage on Aug. 10
Complications	Partial seizures, diabetes insipidus	Partial seizures	Deep-vein thrombosis	Papilledema, hydrocephalus	Coagulopathy, polyuria
Laboratory results					
Random blood glucose — mg/dl	115	90	133	120	109
Cerebrospinal fluid					
Protein — mg/dl	68	49	134	33	28
Glucose — mg/dl	56	25	21	45	61
Neutrophils — no. per mm ³	300	400	20	572	0
Lymphocytes — no. per mm ³	2	175	700	858	225
Gram's stain	Negative	Negative		Negative	Negative
Cytologic findings	Negative	Negative	Positive for fungal spores	Negative	Negative
Fungal culture of cerebrospinal fluid	Negative	Negative	Negative	<i>Aspergillus fumigatus</i>	Negative
Postmortem fungal culture of brain specimen	<i>A. fumigatus</i>	<i>A. fumigatus</i>			<i>A. fumigatus</i>

Gunaratne PS, et al. N Engl J Med. 2007;356:754

Changes in profiles of airborne fungi in flooded homes after typhoon, Taiwan



Hsu et al. Sci Total Environ 2011;409:1677

Percentage profiles of individual fungal species before and after typhoon

Fungal species	Indoor, mean \pm SD (%)	
	Before typhoon (n = 11)	After typhoon (n = 11)
<i>Aspergillus niger</i>	0.22 \pm 0.38	1.59 \pm 2.65
<i>Aspergillus flavus</i>	0.04 \pm 0.14	0.00 \pm 0.00
<i>Aspergillus glaucus</i>	0.00 \pm 0.00	0.10 \pm 0.33
<i>Aspergillus versicolor</i>	0.00 \pm 0.00	0.83 \pm 1.11
<i>Aspergillus terreus</i>	0.00 \pm 0.00	9.79 \pm 15.30
<i>Aspergillus candidus</i>	0.62 \pm 0.85	1.73 \pm 3.40
Other <i>Aspergillus</i> spp.	0.00 \pm 0.00	11.34 \pm 15.84
<i>Penicillium</i>	1.62 \pm 2.71	1.80 \pm 3.62
<i>Cladosporium</i>	34.76 \pm 22.91	2.72 \pm 5.47
<i>Alternaria</i>	1.16 \pm 1.18	0.28 \pm 0.92
<i>Paecilomyces</i>	1.22 \pm 1.50	4.20 \pm 4.62
<i>Curvularia</i>	0.13 \pm 0.42	0.00 \pm 0.00
<i>Fusarium</i>	0.57 \pm 1.52	0.78 \pm 1.80
<i>Drechslera</i>	0.66 \pm 1.59	0.48 \pm 0.88
Non-sporing	43.52 \pm 22.65	46.79 \pm 22.90
Yeast	14.67 \pm 12.98	11.01 \pm 10.43
<i>Trichoderma</i>	0.59 \pm 0.80	0.00 \pm 0.00
<i>Zygosporium</i>	0.00 \pm 0.00	0.00 \pm 0.00
Unknown	0.23 \pm 0.66	6.57 \pm 10.20

Hsu et al. Sci Total Environ 2011;409:1677

Invasive fungal infections after natural disasters

- Route of transmission (pattern of infection)
 - Inhalation (respiratory)
 - Trauma (soft tissue infection, CNS)
 - Near-drowning (respiratory)
 - Indoor exposures (respiratory)
 - Healthcare—associated

Benedict & Park. Emerg Infect Dis 2014;20:349

One Health

Agricultural fungicides and invasive fungal infections in humans

- To minimize agricultural losses from fungal diseases, fungicides are routinely applied to economically valuable crops.
- Although controversial, there is concern that extensive use of agricultural triazoles can induce the resistance of *A. fumigatus* to medically important triazoles such as itraconazole, voriconazole, and posaconazole.
- In the Netherlands, the prevalence of itraconazole resistance of *A. fumigatus* isolated from humans was 6.0%, and 94% of these resistant isolates contain a 34-base pair tandem repeat and a point mutation in *cyp51A* (TR₃₄/L98H mutation). The TR₃₄/L98H mutant also confers reduced susceptibility or resistance to voriconazole and posaconazole. *A. fumigatus* TR₃₄/L98H mutants have also been isolated from humans in several other European countries and Asia.

Gauthier GM, Keller NP. Fungal Genet Biol. 2013 Dec;61:146

Azole Resistance in *Aspergillus fumigatus*

Global 3-6%, Mostly TR₃₄/L98H, which is link to use of azole fungicides

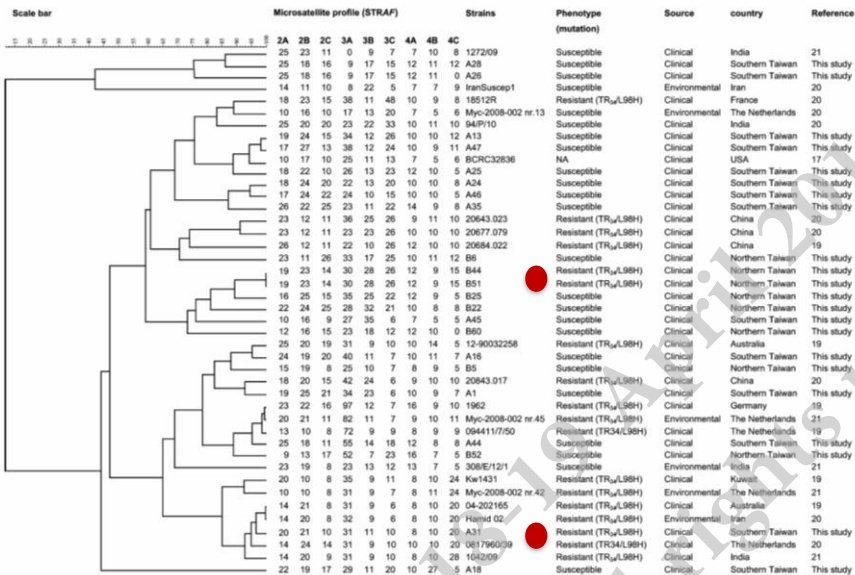
Shaded areas show countries that have reported the TR₃₄/L98H and TR₄₆/Y121F/T289A resistance mechanism in clinical or environmental *A. fumigatus* isolates



Verweij et al. Clin Infect Dis. 2016;62:362; Meis et al. Philos Trans R Soc Lond B Biol Sci. 2016;371(1709).

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Taiwan and Global *A. fumigatus* isolates



The prevalence rates of azole resistance were 7.9% and 6.5% based on isolates and patients respectively.

Wu CJ et al. Mycoses, 2015;58: 544.

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Azole Resistance in *Aspergillus fumigatus*: Can We Retain the Clinical Use of Mold-Active Antifungal Azoles?

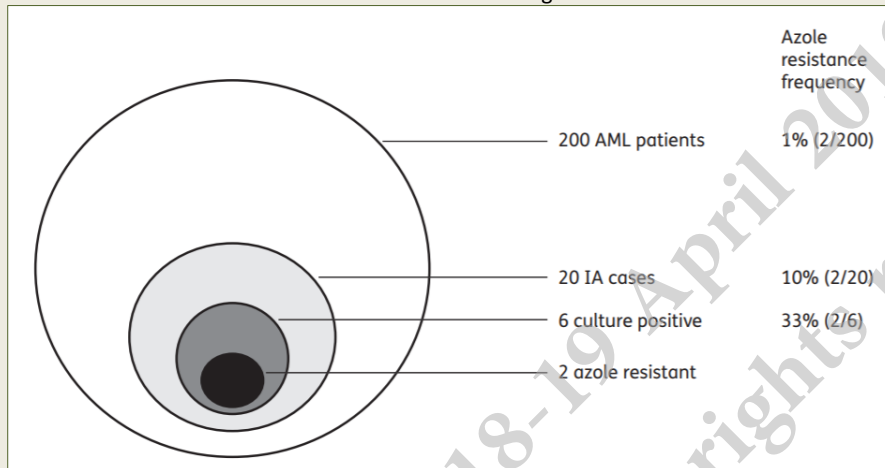
- An expert panel recommended reconsidering the use of azole monotherapy, i.e., voriconazole, in regions with azole resistance rates exceeding 10%.
- Alternative empirical therapy included either liposomal amphotericin B or voriconazole and echinocandin combination therapy, but the efficacy of these alternative treatment options was the subject of much debate.

Verweij PE, Ananda-Rajah M, Andes D et al. International expert opinion on the management of infection caused by azole-resistant *Aspergillus fumigatus*. Drug Resist Updat 2015; 21

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Impact of the denominator on the resistance frequency in azole resistance surveillance

Characteristics of three surveillance strategies for azole resistance



Verweij et al. JAC 2016;71:2079

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Conclusion

Invisible and unexpected

Think fungus

Call for action

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Invisible and Unexpected

Invisible

unaware

What the mind does
not know, the eye
does not see

Misidentification

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Asia on High Alert for Fungus

FUNGAL DISEASE AWARENESS WEEK



SEPTEMBER 23-27, 2019

www.cdc.gov/fungal



<https://www.cdc.gov/fungal/awareness-week.html>

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Call for Action

Infection prevention and control

Increase in vigilance

Diagnostic stewardship

Strengthen capability and capacity for medical mycology

Need for a One Health strategy

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Thanks for your attention.

YC Chen at
NTUCM