



Taiwan Surveillance of Antimicrobial Resistance of Yeasts

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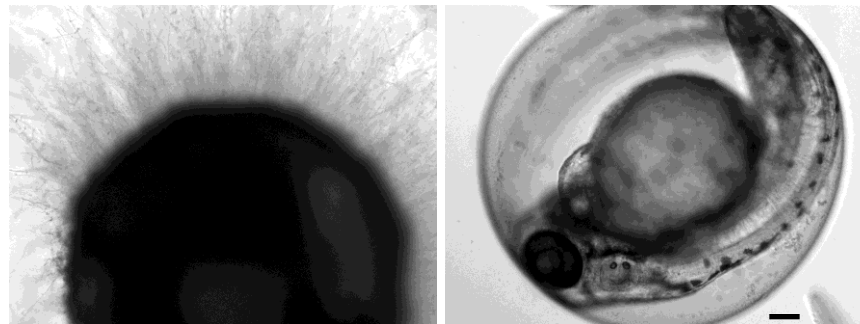
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TSARY Hospitals

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中山醫學大學附設醫院

台中仁愛綜合醫院

台中榮民總醫院

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衛生署花蓮醫院

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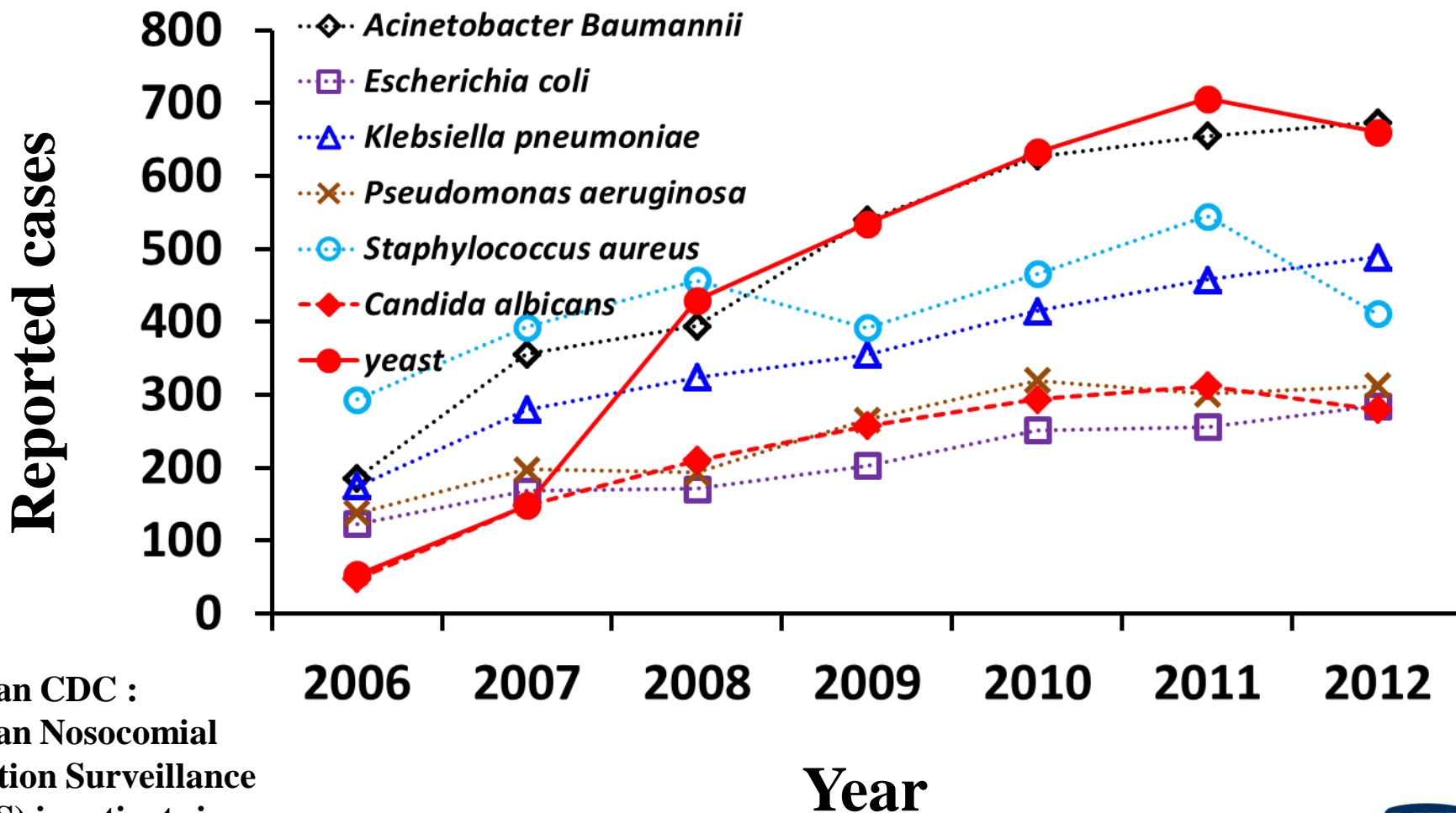
羅東博愛醫院

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Yeast Infections Increased Significantly



Taiwan CDC :
Taiwan Nosocomial
Infection Surveillance
(TNIS) inpatients in
intensive care unit (ICU)





Fungal Infections: Emerging Threats to Human Health

- **1,000,000,000** skin infections
- **100,000,000** Mucosal infections
- **10,000,000** Allergies and SAFS
(Severe asthma with fungal sensitization)
- **1,000,000 death**
 - = tuberculosis
 - > malaria
 - > breast cancer





Candida auris: 新興抗藥念珠致病菌種 I

感染的情況

日本在**2009**年由一位住院病患外耳道分離出*C. auris*來。

2011年韓國報導fluconazole和/或amphotericin B治療*C. auris*菌血症的三位病患中，其中**兩位病人**因治療失敗死亡。

印度、南非、科威特、英國、委內瑞拉、巴西、美國、哥倫比亞、巴基斯坦，西班牙、德國、以色列、挪威及阿曼也偵測到*C. auris*引起菌血症或侵入性感染。Sept. 14, 2017 止，已有**67篇**相關論文

鑑定：

*Candida auris*在傳統形態上或Vitek 2和API系列會被誤判為*Candida haemulonii*、*Candida famata*、*Candida sake*、*Saccharomyces cerevisiae*或*Rhodotourla glutinis*。目前需以**ribosomal DNA** 序列或matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (**MALDI-TOF**)才能正確鑑定出此菌種。

*Candida auris*好發於**住院幾星期後(10 至50天左右)**的病患，*Candida auris*菌血症和**30%到60%**的死亡有關。分子流行病學研究結果顯示每個地區有他自己獨特盛行的菌株品系(**clones**)。





Candida auris: 新興抗藥念珠致病菌種 II

抗藥性：

幾乎全部*C. auris*對**fluconazole**有抗藥性，三分之一的菌株對voriconazole 及 amphotericin B的感受性低(最低抑菌濃度 ≥ 2 mg/L)，有些菌株對 echinocandin感受性也低。目前建議是以**echinocandin**為治療*C. auris*感染的第一線藥。

感染的防治：

因為*C. auris*抗藥性的特質，人體一旦有*C. auris*共生(colonization)後，就不容易將他移除。雖然*C. auris*在呼吸道、尿道與腸道有被偵測到。但是，目前還不清楚一般感染是由自己本身原有共生的菌株(endogenous)或是外來的菌株(exogenous)所造成，還是兩種方式都有可能。

所幸，在五期(1999、2002、2006、2010及2014年) TSARY所收集的約五千株菌株中，並沒有偵測到*C. auris*。

感控雜誌 中華民國 105 年 12 月第二十六卷六期 271-272
Sears D, Schwartz BS (2017). *Candida auris*: an emerging multidrug-resistant pathogen. Int J Infect Dis.





Taiwan Mycology Reference Center

台灣黴菌實驗中心

<http://mycology.nhri.org.tw>

Taiwan Mycology Reference Center (TMRC) provides information about fungal infections.



The screenshot shows the website interface for the Taiwan Mycology Reference Center. The browser window displays the URL <http://mycology.nhri.org.tw/>. The main banner features three images: a microscopic view of a dark, fuzzy fungal mass on the left, a petri dish with various colored fungal cultures in the center, and a microscopic view of a white, branching fungal structure on the right. The text "台灣黴菌諮詢中心" and "Taiwan Mycology Reference Center" is overlaid on the banner. Below the banner is a navigation menu with links: 首頁 / 最新消息 / 諮詢委員 / 服務項目 / 使用者資訊 / 教育訓練 / 研究主題 / 會員資訊 / 工作團隊 / 實驗步驟 / 聯絡我們. The footer contains the text "財團法人國家衛生研究院" (National Health Research Institutes) and "衛生福利部疾病管制署" (Centers for Disease Control and Prevention), along with the NHRI logo. The system tray at the bottom shows the date and time as 2015/10/13, 下午 09:26.



Current Challenges for Managing Fungal Infections

- 1. Risk populations increased.**
- 2. Limited choice of antifungal drugs.**
- 3. Emerging drug resistance problems.**
- 4. Emerging species causing diseases in humans.**
- 5. How to improve diagnosis of fungal infections?**
- 6. How to develop new effective antifungal drugs?**
- 7. How to transmit drug resistant pathogenic fungi?**

How can a basic researcher help?





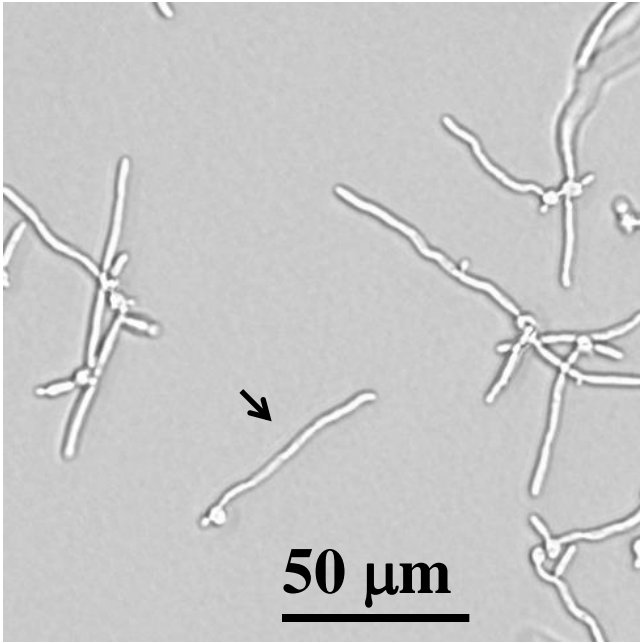
Taiwan Surveillance of Antimicrobial Resistance of Yeasts (TSARY)

	Year				
	1999	2002	2006	2010	2014
Collection period	Apr 15 to Jun 15	Jul 1 to Sep 30			
No. of hospitals	22	24	22	24	24
Sources	All body sites	Sterile sites	Non-Sterile sites		
<i>C. albicans</i>	up to 10	all	up to 10		
Non- <i>C. albicans</i>	up to 40	all	up to 40		
Total	670	945	1015	1130	1165

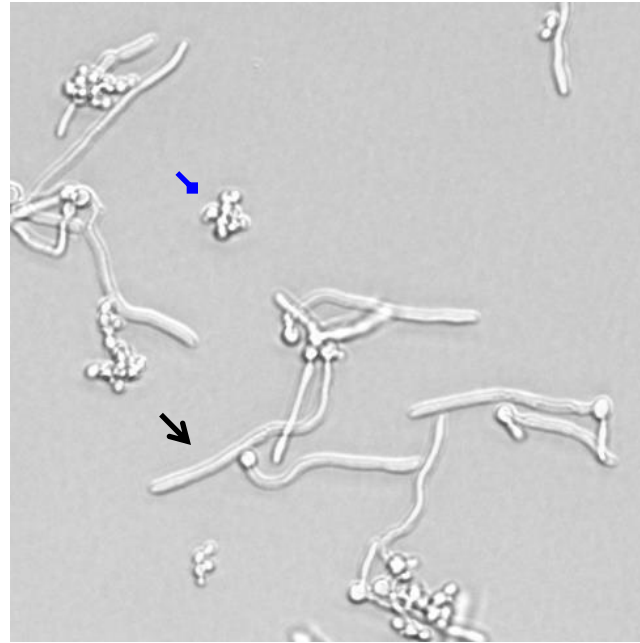
TSARY 2002-2014 have same set of hospitals and were conducted along with TSAR.



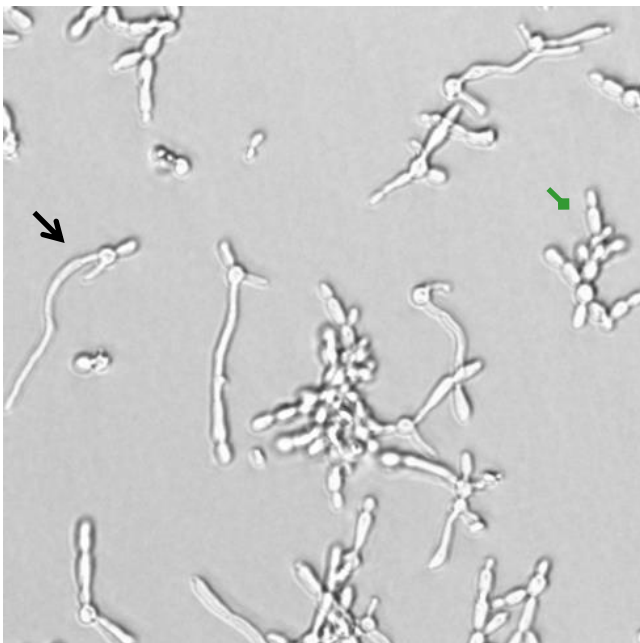
Germ Tube Assay



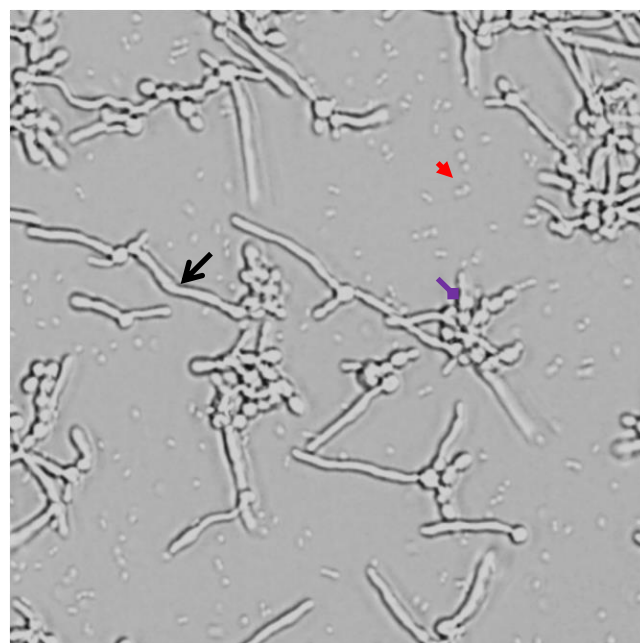
*C.
albicans*



*C.
albicans*
*C.
glabrata*



*C.
albicans*
*C.
tropicalis*



*C.
albicans*
*C.
parapsilosis*
Bacterium

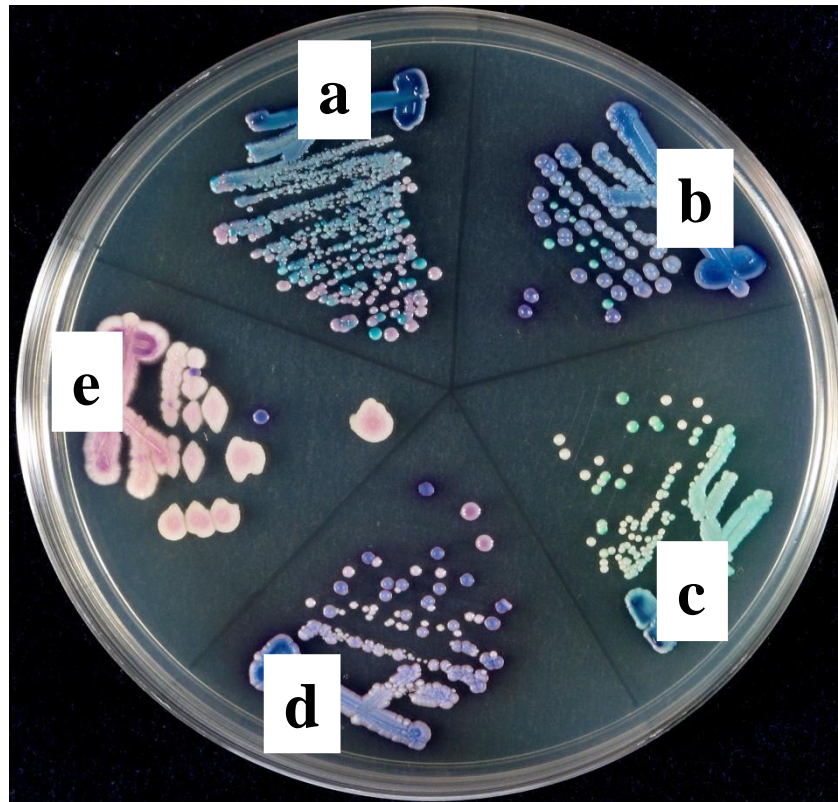


The Power of CHROMagar Candida

C. albicans
C. glabrata

C. krusei
C. tropicalis

C. glabrata
C. tropicalis

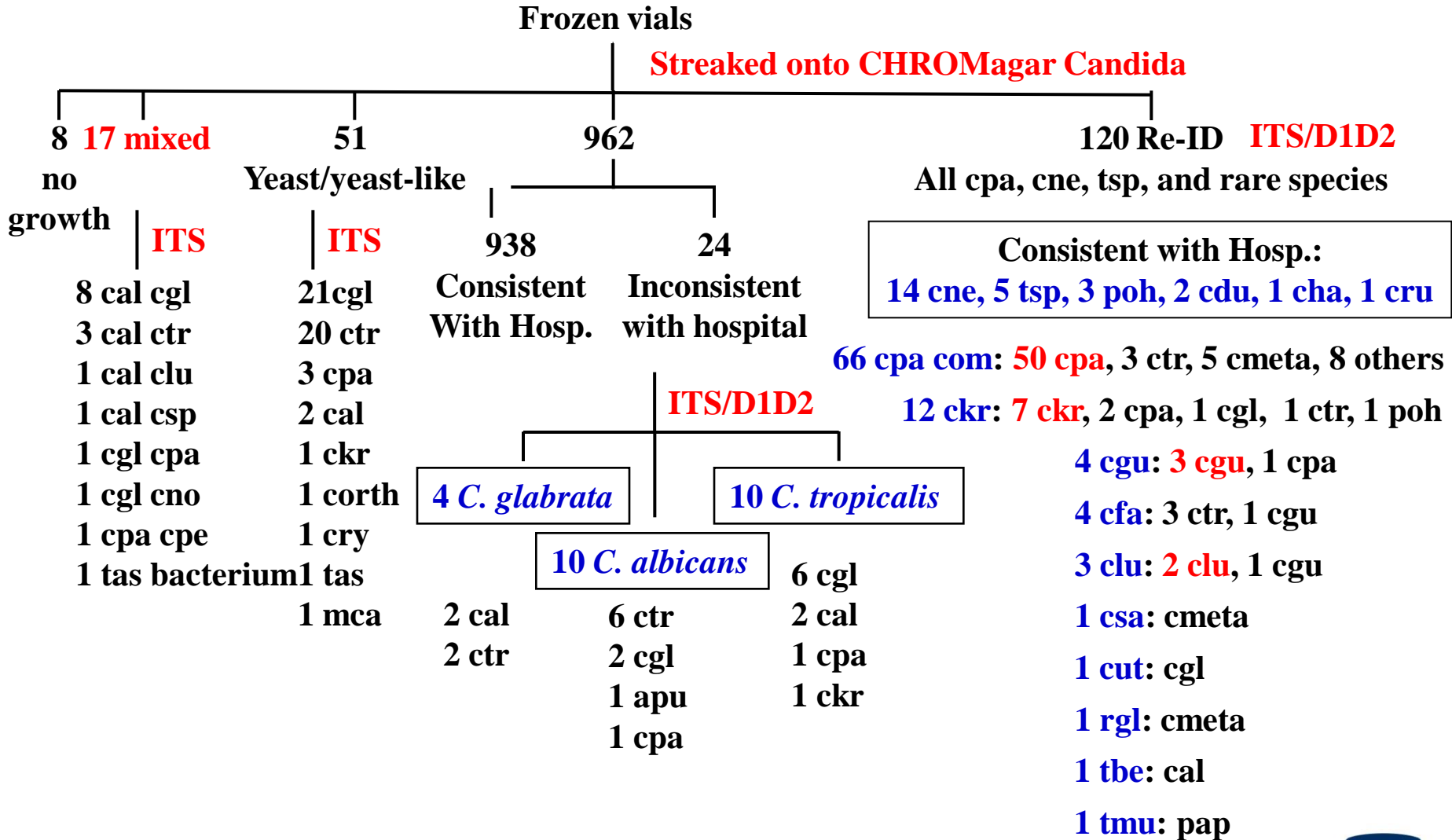


C. albicans
C. tropicalis

C. albicans
C. parapsilosis



TSARY 2014



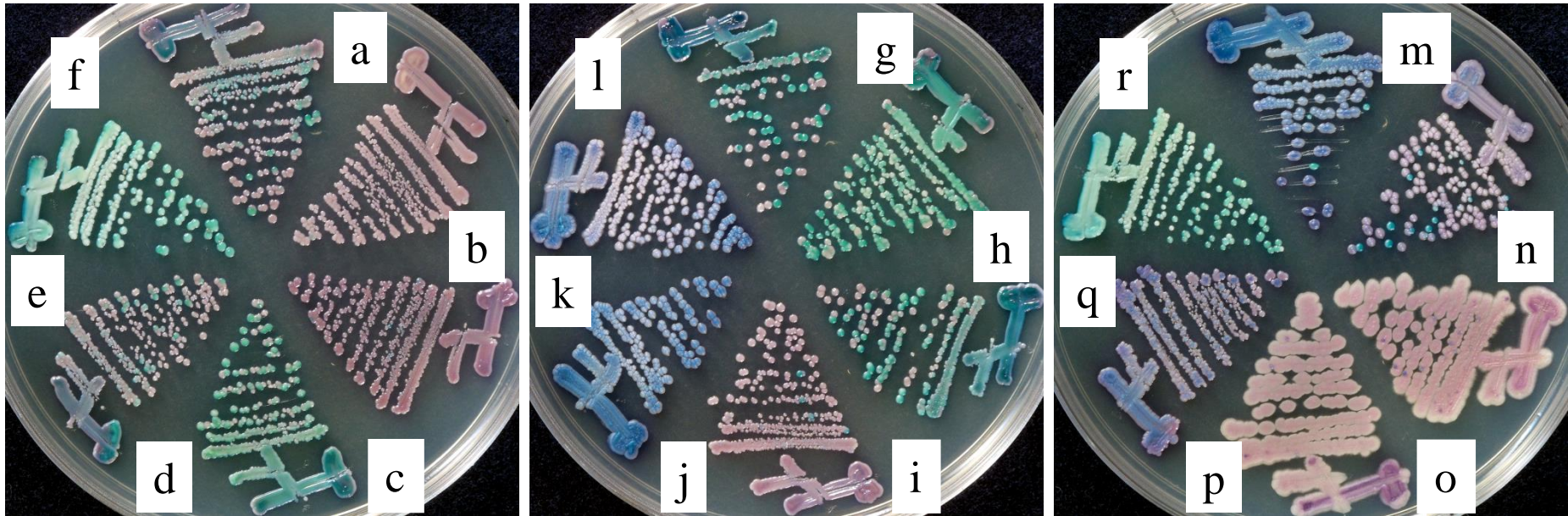
The Distribution of Species in Specimens Having Mixed Yeast Cultures (TSARY 2014)

	cal			ctr			cgl			cpa				
	cgl	cgl	ctr	can	clu	cno	cgl	cgl	cgl	cpa	can	ckr	clu	cpe
	ctr			cpa			cno							
Ascites (1/3)*	0	2	1	0	0	0	0	0	0	0	0	0	0	0
Blood (1/5)	1	2	1	0	0	0	0	0	0	0	0	0	0	1
Sputum (7/12)	0	3	5	0	1	1	0	1	0	0	0	1	0	0
Urine (5/13)	0	5	1	1	0	0	1	0	3	0	1	0	1	0
Wounds (2)	0	0	1	0	0	0	0	0	0	1	0	0	0	0
Tip (1/2)	0	0	2	0	0	0	0	0	0	0	0	0	0	0
Pus (1)	0	0	0	0	0	0	0	0	0	1	0	0	0	0
Stool (1)	0	0	0	0	0	0	0	0	1	0	0	0	0	0
Mouth (1/1)	0	1	0	0	0	0	0	0	0	0	0	0	0	0
TA(1)	0	0	0	0	0	0	0	0	1	0	0	0	0	0
Total (16/41)	1	13	11	1	1	1	1	1	5	2	1	1	1	1

*Number of mix specimen identified by NHRI/total specimens, cal: *C. albicans*, can: *Candida* sp. , cgl: *C. glabrata*, ckr: *C. krusei*, clu: *C. lusitaniae*, cno: *C. norvegensis*, cpa: *C. parapsilosis*, ctr: *C. tropicalis*, cpe: *C. pelliculosa*, TA: Tracheal aspiration.



The Power of CHROMagar Candida



CHROMagar Candida medium is good for specimens from

Sterile sites

Patients having fever after antibacterial treatment



Species Distribution Causing Candidemia

Species	World (1992-2001)	SENTRY (2008–2009)	TSARY (2002-2006)
	Blood N = 6080 CMI 2004	Blood N = 1354 AAC 2011	Invasive infections N= 339 MM 2010
<i>C. albicans</i>	3401 (55.9)	655 (48.4)	182 (53.7)
<i>C. tropicalis</i>	585 (9.6)*	143 (10.6)*	75 (22.1)**
<i>C. glabrata</i>	984 (16.2)	247 (18.2)*	42 (12.4)*
<i>C. parapsilosis</i>	796 (13.1)**	232 (17.1)**	31 (9.1)**
<i>C. krusei</i>	153 (2.5) §	27 (2)	3 (0.9) §
<i>C. lusitaniae</i>	67 (1.1)	0	0
<i>C. dubliniensis</i>	3 (0.05)	16 (1.2)	0
<i>C. guilliermondii</i>	48 (0.8)	8 (0.6)	2 (0.6)
<i>C. kefyr</i>	10 (0.16)	6 (0.4)	0
<i>C. famata</i>	10 (0.16)	3 (0.22)	0
<i>C. lipolytica</i>	4 (0.07)	3 (0.22)	0
<i>C. rugosa</i>	5 (0.08)	2 (0.15)	1 (0.3)
<i>C. sake</i>	2 (0.03)	2 (0.15)	0
<i>C. pelliculosa</i>	6 (0.1)	2 (0.15)	0
<i>C. inconspicua</i>	3 (0.05)	1 (0.07)	0
<i>C. lambica</i>	1 (0.02)	1 (0.07)	0
<i>C. norvegensis</i>	1 (0.02)	1 (0.07)	0
<i>C. zeylanoides</i>	1 (0.02)	1 (0.07)	0

**Less
fluconazole
prophylaxis
in Taiwan**



The Distribution of Body Sites

Source	Urine	Sputum	Blood	Wound	Ascites	Tip	Pus	others	Total
Total	1810	684	754	169	142	166	130	403	4258
%	42.5	16.1	17.7	4.0	3.3	3.9	3.1	9.5	100
2014	520	158	240	34	38	35	45	98	1168
%	44.5	13.5	20.5	2.9	3.3	3.0	3.9	8.4	100
2010	500	149	224	42	33	54	30	98	1130
%	44.2	13.2	19.8	3.7	2.9	4.8	2.7	8.7	100
2006	416	166	160	46	38	27	41	121	1015
%	41	16.4	15.8	4.5	3.7	2.7	4	11.9	100
2002	374	211	130	47	33	50	14	86	945
%	39.6	22.3	13.7	5	3.5	5.3	1.5	9.1	100

**2/3 of isolates were from urine, sputum, and blood.
Isolates from blood increased since 2010.**



Distribution of Species from Blood

	<i>C. albicans</i>		<i>C. tropicalis</i>		<i>C. glabrata</i>		<i>C. parapsilosis</i>		Others		Total
	No.	%	No.	%	No.	%	No.	%	No.	%	No.
2014											
blood	100	41.7	46	19.2	43	17.9	27	11.3	24	10	240
Subtotal	392	ND	294	25.2	325	27.8	58	5	99	8.5	1168
2010											
blood	103	46.2	41	18.4	22	9.9	35	15.7	22	9.9	223
Subtotal	423	ND	270	24	262	23.2	87	7.7	85	7.5	1127
2006											
blood	79	50.6	31	19.9	20	12.8	9	5.8	17	10.9	156
Subtotal	419	ND	248	24.4	211	20.8	62	6.1	75	7.4	1015
2002											
blood	48	42.9	29	25.9	13	11.6	19	17	3	2.7	112
Subtotal	406	ND	244	25.8	187	19.8	63	6.7	45	4.8	945

Susceptibility to Fluconazole

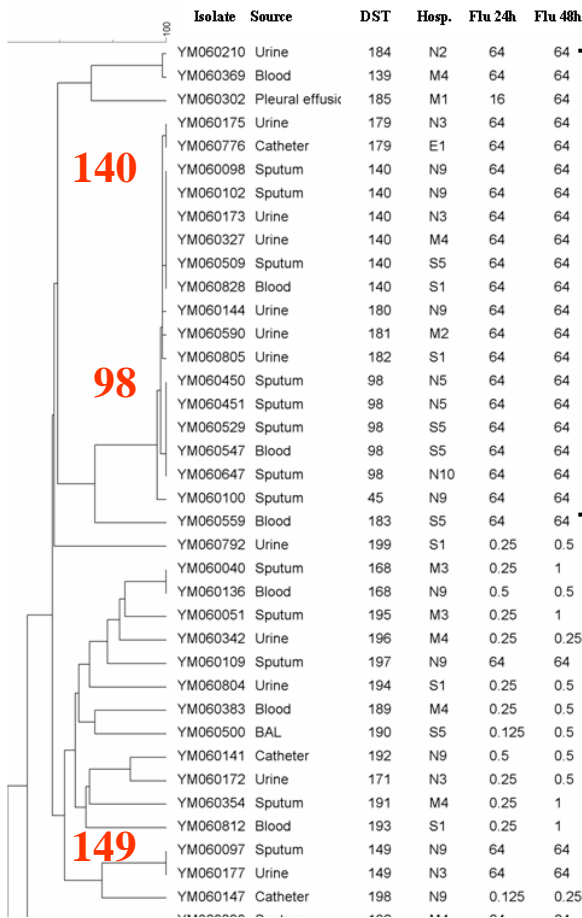
MICs (mg/l)							
At 48 h	<i>C. albicans</i>	<i>C. tropicalis</i>	<i>C. glabrata</i>	<i>C. parapsilosis</i>	<i>C. krusei</i>	Others	Total
2006 ≤ 8	407 (97.1)	108 (43.9)	169 (80.1)	61 (98.4)	0	11 (91.7)	756 (78.4)
16-32	2 (0.5)	6 (2.4)	33 (15.6)	1 (1.6)	0	1 (8.3)	43 (4.5)
≥ 64	10 (2.4)	132 (53.7)	9 (4.3)	0	14 (100)	0	165 (17.1)
Subtotal	419	246	211	62	14	12	964
2002 ≤ 8	373 (94.4)	235 (96.3)	93 (49.7)	63 (100)	1 (11.1)	7 (63.6)	772 (84.9)
16-32	17 (4.3)	9 (3.7)	91 (48.7)	0	1 (11.1)	2 (18.2)	120 (13.2)
≥ 64	5 (1.3)	0	3 (1.6)	0	7 (77.8)	2 (18.2)	17 (1.9)
Subtotal	395	244	187	63	9	11	909
1999 ≤ 8	221 (93.2)	129 (79.1)	121 (77.6)	50 (98)	0	13 (86.7)	534 (84.5)
16-32	7 (3)	10 (6.1)	22 (14.1)	1 (2)	3 (30)	2 (13.3)	45 (7.1)
≥ 64	9 (3.8)	24 (14.7)	13 (8.3)	0	7 (70)	0	53 (8.4)
Subtotal	237	163	156	51	10	15	632

Breakpoints before 2010: MICs ≤ 8: Susceptible, 16-32: SDD, ≥ 64: Resistant. Trialing growth issue.

Are *C. tropicalis* isolates with azole reduced-susceptibility genetically related?



Two Closely Related Fluconazole-Resistant *Candida tropicalis* Clones Circulating in Taiwan from 1999 to 2006



Resistant

18 isolates of DST140 were isolated from **10 different hospitals** localized in all four geographic regions in Taiwan.

7 isolates of DST98 were also isolated from **4 different hospitals** located in both north and south Taiwan.

3 isolates of DST149 from **3 different hospitals** in north.

MLST: Multilocus sequence type



Azole Susceptibilities of *Candida* sp.

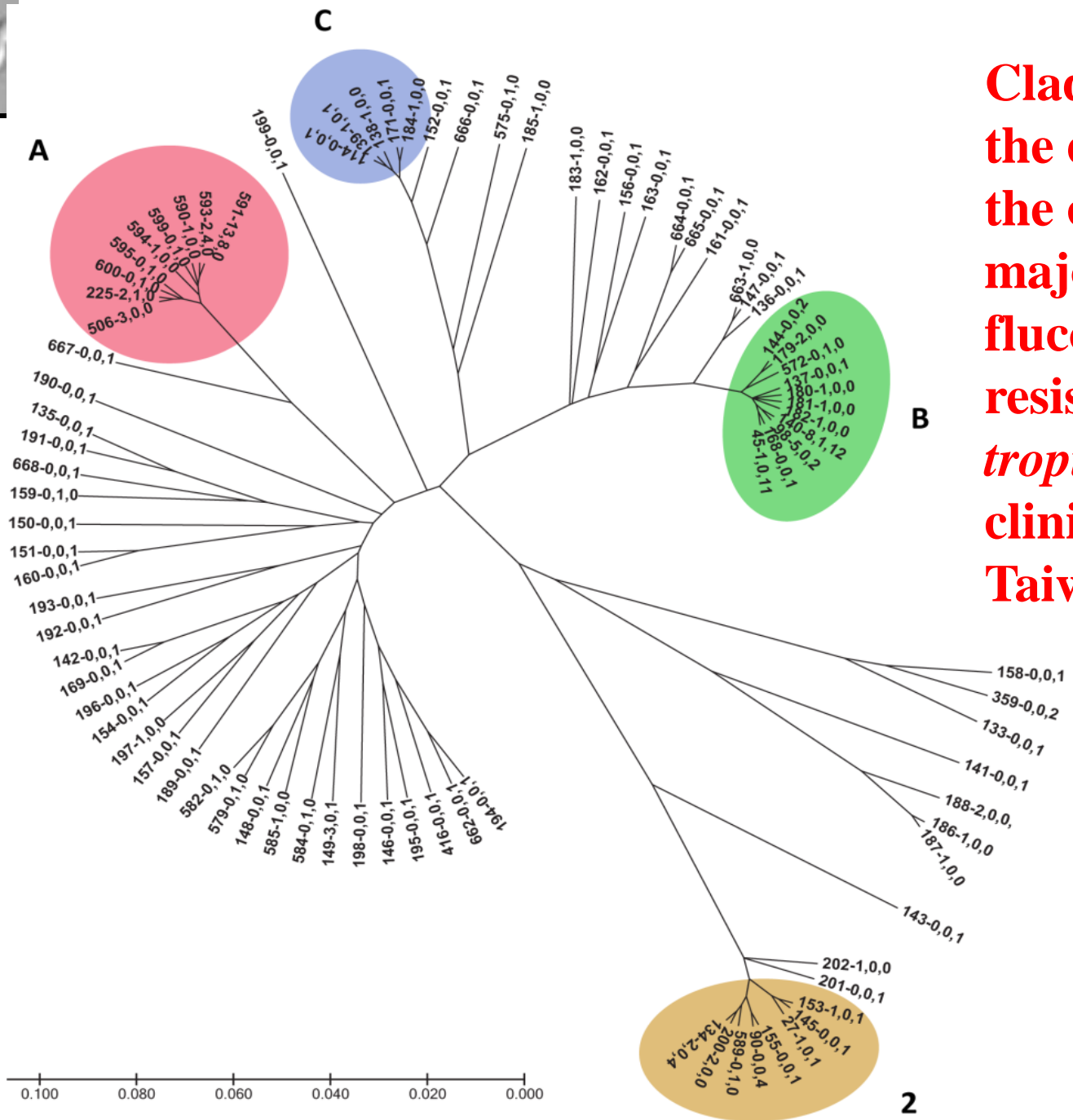
		<i>C. albicans</i>	<i>C. tropicalis</i>	<i>C. glabrata</i>	<i>C. parapsilosis</i>	<i>C. krusei</i>	Others	Total
2010	Total	422	270	258	87	18	28	1083
2014	Total	397	294	308	58	11	31	1099
MIC (mg/L)		Fluconazole						
2010	≤ 2	418	269	232	86	1	23	1029
	4	2*	1*	15	0	0	5*	23
	8	0	0	0	1 [#]	9	0	10
	16	2 [#]	0	7	0	8	0	17
	32	0	0	2*	0	0	0	2
	≥ 64	0	0	2 [#]	0	0	0	2
2014	≤ 2	396	247	160	58	0	28	889
	4	0	22*	125	0	1	2*	150
	8	0	2 [#]	7	0	4	0	13
	16	0	0	5	0	5	1 [#]	11
	32	0	18 [#]	7*	0	0	0	25
	≥ 64	1 [#]	5 [#]	4 [#]	0	1	0	11
		Voriconazole						
2010	≤ 0.125	419	270	248	87	18	28	1070
	0.25	0	0	8	0	0	0	8
	1	3 [#]	0	2	0	0	0	5
2014	≤ 0.125	395	267	293	58	10	30	1053
	0.25	1	3	4	0	0	1	9
	0.5	0	1	7	0	1	0	9
	1	0	19 [#]	4	0	0	0	23
	2	0	2 [#]	0	0	0	0	2
	> 8	1 [#]	2 [#]	0	0	0	0	3

MIC, minimum inhibitory concentration; *susceptible-dose dependent; #resistant





Clade A replaced the dominance of the clade B as the major clone of fluconazole-resistant *C. tropicalis* in clinical settings in Taiwan.



2





Summary of Five TSARYs

There is an increased rate of **mixed cultures**.

Urine (42%) was the most common source, followed by sputum (18%), **blood (15%)** etc.

The distribution of *Candida* species was similar.

Candida albicans is the predominant species accounting for half of invasive infections.

Candida tropicalis was one of the most frequently isolated non-*albicans Candida* sp. and had high fluconazole resistant rate.

The prevalence of candidemia caused by *Candida glabrata* increased in 2014.

Two closely related fluconazole resistant *C. tropicalis* clones (DST98 and DST140) circulate in Taiwan from 1999 to 2006.

An Emerging Azole-Resistant *Candida tropicalis* DST591 Clone in Taiwan

Majority of isolates are susceptible to amphotericin B (97.5% to 99.9%) and anidulafungin (99.6% in 2010).





How Can These Closely Related Clones Widely Distribute?

Same azole drugs are used in **humans, animals, and plants** because the same groups of fungi cause disease in humans, animals and plants. **About 80% of plant infections are due to fungi.** The implications of this for development of resistance are unknown.

These closely related clones may be transmitted by different means instead of person to person.

Animals?

Foods? (Fruits, orange juice, cider)

Contaminated Medical supply?

Environments: Water? **Soil?**

Dr. Lee at NHUE had recovered 56 *C. tropicalis* isolates from different soil samples from 2006 to 2008.





Approximately 1/3 of 56 *Candida tropicalis* from Soil with Reduced-Susceptibility to Fluconazole

MIC (mg/l)	North		Middle		South		Total
	Tao yuan	Hsin chu	Chang hua	Miaoli	Tai nan	Kao hsiung	
0.125	0	1	0	0	0	0	1
0.25	1	0	0	0	0	1	2
0.5	6	2	0	2	14	3	27
1	0	2	0	0	0	3	5
2	1	0	0	0	0	0	1
4	1	0	0	0	0	0	1
8	1	0	0	0	0	0	1
≥ 64	7	5	1	3	1	1	18
Total	17	10	1	5	15	8	56



Candida tropicalis from Soil and Humans Were Genetically Closely Related

DST	North		Middle		South		Total
	Tao yuan	Hsin chu	Chang hua	Miao li	Tai nan	Kao hsiung	
140	0	0	0	1	1	1	3
149	1	5	3	0	0	0	9
Other 5	5	0	0	0	0	0	5
Total	7	5	3	1	1	1	18

18 of 56 (32%) *C. tropicalis* with reduced-susceptibility to fluconazole and 12 of 18 (67%) were DST140 or DST149, which were found in isolates from human.



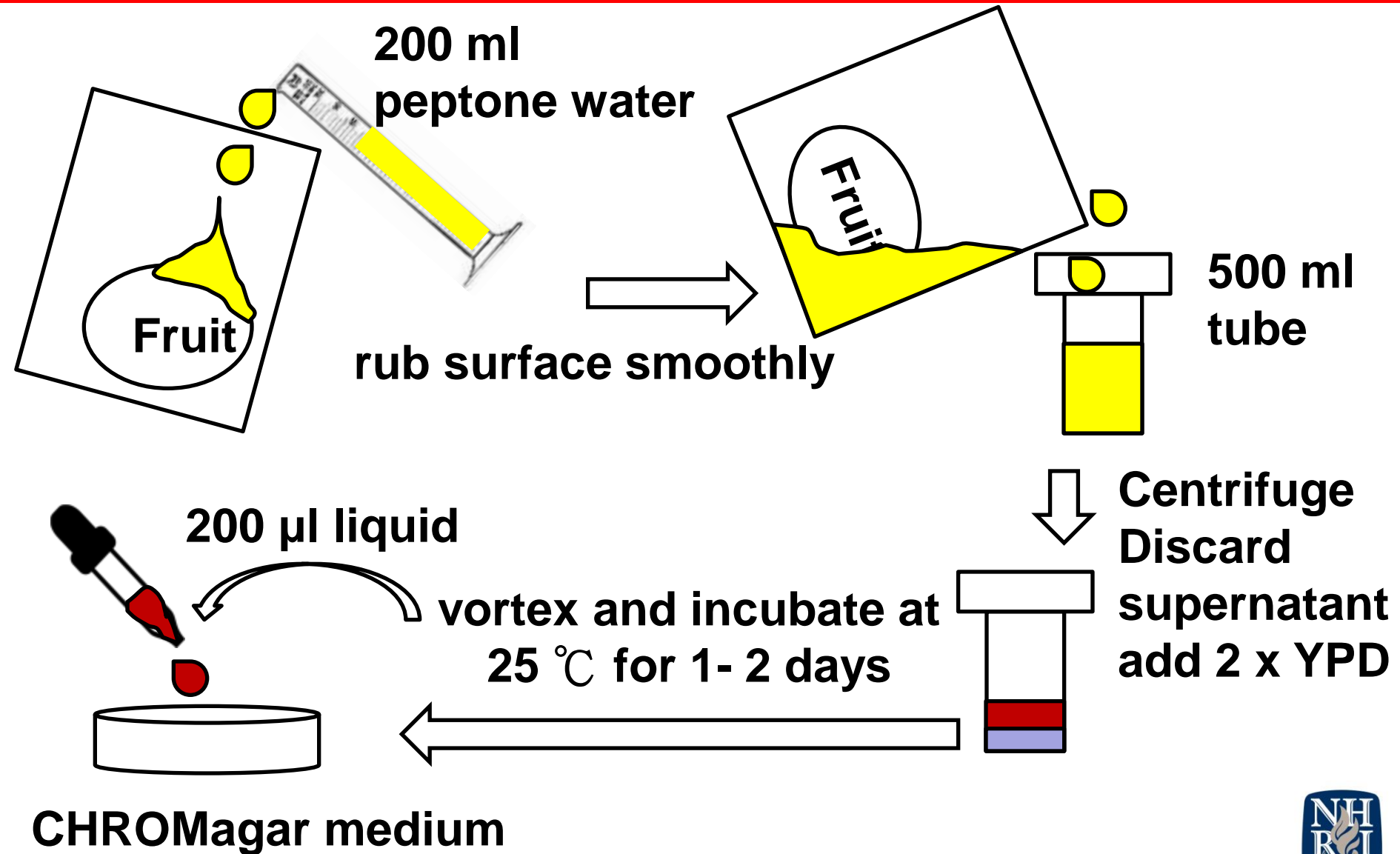


**How can pathogenic yeasts in soil
transmitted to humans?**

**How about the Yeasts from
Surfaces of Fruits
in Supermarkets in Taiwan?**



Isolation of Yeasts from Fruits



Distribution of Yeast Species



Type of fruit (number of total fruit sampled/number of sampling)

species	Type of fruit (number of total fruit sampled/number of sampling)																						
	Total	Mango (36/6)	Melon (19/7)	Pear (31/7)	Citrus (129/9)	Grape (900/5)	Tomato (246/4)	Lemon (26/2)	Peach (33/2)	Star fruit (9/3)	Wax apple (18/2)	Jujube (73/2)	Banana (11/2)	Guava (10/2)	Coconut (3/1)	Kiwi (10/1)	Loquat (30/1)	Papaya (4/2)	Plum (100/1)	Salted olive (24/1)	Shakya (4/1)	Persimmon (3/1)	Salted plum (26/1)
Total	184	22	21	20	18	12	12	8	8	8	8	7	5	5	4	4	4	4	4	3	3	2	2
<i>Pichia kluyveri</i>	24	4	4	1	3	0	2	0	1	2	1	1	0	2	0	0	0	0	1	1	1	0	0
<i>Candida fermentati*</i>	17	1	2	3	2	0	1	0	1	0	1	1	1	1	1	1	1	0	0	0	1	0	0
<i>Hanseniaspora opuntiae</i>	10	3	2	1	2	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Hanseniaspora uvarum</i>	10	1	2	0	1	1	0	1	1	0	0	1	0	0	0	0	0	0	0	1	0	0	1
<i>Candida quercitrusa*</i>	9	0	1	2	2	0	0	0	1	0	1	1	0	0	0	0	0	0	0	0	1	0	0
<i>Candida jamata*</i>	7	1	0	2	1	0	0	0	0	1	0	0	1	0	0	0	0	1	0	0	0	0	0
<i>Hanseniaspora thailandica</i>	7	2	0	0	1	1	0	0	0	1	0	0	0	0	0	0	1	1	0	0	0	0	0
<i>Sporidiobolus parosaeus</i>	7	1	2	0	0	1	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Lodderomyces elongisporus*</i>	6	1	2	0	0	0	0	1	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0
<i>Candida oleophila</i>	5	0	0	0	1	2	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	0	0
<i>Candida natalensis</i>	4	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1
<i>Candida railenensis</i>	4	0	0	0	2	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida sorboxylosa</i>	4	0	0	0	0	1	0	0	0	2	0	0	0	0	0	0	0	0	1	0	0	0	0
<i>Rhodotorula mucilaginosa*</i>	4	1	0	0	0	1	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Aureobasidium melanogenum*</i>	3	1	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida krusei*</i>	3	1	1	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida pulcherrima*</i>	3	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
<i>Candida tropicalis*</i>	3	0	1	0	0	0	0	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0
<i>Debaryomyces nepalensis*</i>	3	1	0	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Issatchenkia occidentalis</i>	3	0	0	0	0	0	0	0	0	0	2	0	1	0	0	0	0	0	0	0	0	0	0
<i>Rhodospiridium paludigenum</i>	3	0	2	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Rhodotorula glutinis*</i>	3	0	0	1	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
<i>Candida catenulata*</i>	2	0	0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida conglobata*</i>	2	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
<i>Candida orthopsilosis*</i>	2	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida valida*</i>	2	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Cryptococcus flavescens*</i>	2	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
<i>Issatchenkia terricola*</i>	2	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Pichia aff. Fermentans Y153</i>	2	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0
<i>Rhodotorula diobovata</i>	2	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Trichosporon asahii*</i>	2	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0
<i>Aureobasidium pullulans*</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0
<i>Candida akabanensis</i>	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida diversa</i>	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida guilliermondii var. membranifaciens*</i>	1	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida guilliermondii*</i>	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida intermedia*</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
<i>Candida lipolytica*</i>	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida nonsorbohypha</i>	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida parapsilosis*</i>	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida pelliculosa*</i>	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Candida xylopoeci</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
<i>Cyberlindnera xylosilytica</i>	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Hanseniaspora guilliermondii</i>	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
<i>Pichia mexicana*</i>	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Pichia pipieri</i>	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
<i>Pseudozyma fusiformata</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
<i>Pseudozyma hubeiensis</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
<i>Rhodospiridium babjevae</i>	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Rhodotorula dairenensis*</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
<i>Saccharomyces bulderi</i>	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Saccharomyces cerevisiae*</i>	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Torulopsis delbrueckii</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0
<i>Trichosporon jirovecii*</i>	1	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Wickerhamomyces pipieri</i>	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0

*Has been reported to cause diseases in humans





Distribution of Yeast Species

22 different types of fruits

184 isolates,

55 species

29 species, including *Candida famata*, *Candida fermentati*, *Candida guilliermondii*, *Candida intermedia*, *Candida krusei*, *Candida orthopsilosis*, *Candida parapsilosis*, *Candida pelliculosa*, *Candida tropicalis*, and others have been reported to cause diseases in humans.





Species	MIC (mg/l)												Total
	fluconazole						triadimenol						
	2	4	8	16	32	64	2	4	8	16	32	64	
Total	59	11	5	3	7	30	65	19	19	8	1	3	115
<i>Pichia kluyveri</i>	0	0	0	1	6	17	5	5	12	2	0	0	24
<i>Candida fermentati</i> *	15	1	1	0	0	0	5	10	1	1	0	0	17
<i>Candida quercitrusa</i> *	6	0	2	0	0	0	7	1	0	0	0	0	8
<i>Lodderomyces elongisporus</i> *	6	0	0	0	0	0	6	0	0	0	0	0	6
<i>Candida famata</i> *	4	0	0	0	0	0	4	0	0	0	0	0	4
<i>Rhodotorula mucilaginosa</i> *	0	0	0	0	0	4	0	0	2	1	0	1	4
<i>Candida natalensis</i>	3	0	0	0	0	0	2	1	0	0	0	0	3
<i>Candida krusei</i> *	0	0	1	2	0	0	3	0	0	0	0	0	3
<i>Candida pulcherrima</i> *	3	0	0	0	0	0	3	0	0	0	0	0	3
<i>Candida tropicalis</i> *	2	1	0	0	0	0	2	0	0	1	0	0	3
<i>Issatchenkia occidentalis</i>	0	0	0	0	0	3	0	0	0	3	0	0	3
<i>Candida oleophila</i>	2	0	0	0	0	0	2	0	0	0	0	0	2
<i>Candida railenensis</i>	1	1	0	0	0	0	2	0	0	0	0	0	2
<i>Candida sorboxylosa</i>	0	2	0	0	0	0	2	0	0	0	0	0	2
<i>Debaryomyces nepalensis</i> *	1	1	0	0	0	0	2	0	0	0	0	0	2
<i>Rhodotorula glutinis</i> *	0	0	0	0	1	1	0	0	1	0	1	0	2
<i>Candida catenulata</i> *	2	0	0	0	0	0	2	0	0	0	0	0	2
<i>Candida conglobata</i> *	2	0	0	0	0	0	1	1	0	0	0	0	2
<i>Candida orthopsilosis</i> *	2	0	0	0	0	0	2	0	0	0	0	0	2
<i>Rhodotorula diobovata</i>	0	0	0	0	0	2	0	0	1	0	0	1	2
<i>Aureobasidium melanogenum</i> *	0	0	1	0	0	0	1	0	0	0	0	0	1
<i>Rhodosporidium paludigenum</i>	0	0	0	0	0	1	1	0	0	0	0	0	1
<i>Cryptococcus flavescens</i> *	0	1	0	0	0	0	1	0	0	0	0	0	1
<i>Trichosporon asahii</i> *	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Candida akabanensis</i>	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Candida guilliermondii</i> var. <i>membranifaciens</i> *	0	1	0	0	0	0	1	0	0	0	0	0	1
<i>Candida guilliermondii</i> *	1	0	0	0	0	0	0	0	1	0	0	0	1
<i>Candida intermedia</i> *	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Candida nonsorbophila</i>	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Candida parapsilosis</i> *	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Candida pelliculosa</i> *	0	1	0	0	0	0	1	0	0	0	0	0	1
<i>Candida xylopoeci</i>	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Pichia mexicana</i> *	1	0	0	0	0	0	0	1	0	0	0	0	1
<i>Pichia pijperi</i>	0	1	0	0	0	0	1	0	0	0	0	0	1
<i>Pseudozyma fusiformata</i>	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Rhodosporidium babjevae</i>	0	0	0	0	0	1	0	0	0	0	0	1	1
<i>Rhodotorula dairenensis</i> *	0	0	0	0	0	1	0	0	1	0	0	0	1
<i>Saccharomyces bulderi</i>	1	0	0	0	0	0	1	0	0	0	0	0	1
<i>Saccharomyces cerevisiae</i> *	0	1	0	0	0	0	1	0	0	0	0	0	1

Drug Susceptibility

C. krusei, all *Rhodotorula* and *Rhodosporidium* species were resistant to fluconazole. One each of *C. tropicalis* isolate was belonged to diploid sequence type (DST)149 and DST225, genotypes also detected in isolates from humans.



Antifungal Susceptibilities of 112 *Candida* spp. at 48-h

Species / MIC (mg/L)	Amphotericin B		Fluconazole			Triadimenol			Total
	≤ 2	≥ 4	≤ 8	16-32	≥ 64	≤ 8	16-32	≥ 64	
<i>Candida guilliermondii</i>	16	0	16	0	0	6	9	1	16
<i>Candida pulcherrima</i>	16	0	15	1	0	15	1	0	16
<i>Candida famata</i>	8	1	8	1	0	9	0	0	9
<i>Candida quercitrusa</i>	6	0	6	0	0	6	0	0	6
<i>Candida catenulata</i>	5	0	5	0	0	5	0	0	5
<i>Candida fermentati</i>	5	0	4	0	1	3	2	0	5
<i>Candida intermedia</i>	5	0	5	0	0	5	0	0	5
<i>Candida sphaerica</i>	3	2	4	0	1	3	2	0	5
<i>Candida parapsilosis</i>	3	0	3	0	0	3	0	0	3
<i>Candida tropicalis</i>	3	0	1	1	1 (F91)	1	1	1 (F91)	3
<i>Candida conglobata</i>	2	0	2	0	0	2	0	0	2
<i>Candida haemulonii</i>	2	0	2	0	0	2	0	0	2
<i>Candida inconspicua</i>	2	0	0	0	2	0	2	0	2
<i>Candida lusitaniae</i>	2	0	2	0	0	2	0	0	2
<i>Candida orthopsilosis</i>	2	0	2	0	0	2	0	0	2
<i>Candida sorboxylosa</i>	2	0	2	0	0	2	0	0	2
<i>Candida zeylanoides</i>	2	0	2	0	0	2	0	0	2
Others	0	0	0	0	0	22	3	0	25
Total	109	3	102	4	6	90	20	2	112

MIC: minimum inhibitory concentration



Fruits as the vehicle of drug resistant pathogenic yeasts

Lo, et al. J Infect. 75, 254-262.

WHY?





Annual Pesticides Use in Taiwan

	Year		
	2005	2007	2009
Tons			
Azole	106	151	144
Fungicides	2254	2306	2210
Total	9228	9492	8589

Similar type of azole drugs used in medical and agriculture needs more attention.



Summary

1. The importance of drug resistant *Candida auris*
2. The importance of **mix yeast infections**
3. The prevalence of candidemia caused by *C. glabrata* increased in 2014.
4. The important of *C. tropicalis* in Taiwan
 - a. *C. tropicalis* isolates from soils and fruits are genetically closely related to those from humans.
 - b. Major clone of fluconazole-resistant *C. tropicalis* in clinical settings in Taiwan has switched from DST140/98 to DST591.
 - c. How genetically related *C. tropicalis* clones coexist in environments and humans?



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Scientific Advisory Board

MIRL Steering Committee

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MOST



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感染症研究組
國家衛生研究院

我們的足跡



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etc





I am lucky to have their help.



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I am luck to have their help.

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**Thank You
for Your
Attention**

