

Classification and new diagnostic methods for invasive fungal infections

Stéphane Bretagne

National Reference Center for Mycoses & Antifungals
CNRS URA3012



INSTITUT PASTEUR



Tunis 24th May 2012

Classification Invasive Fungal Diseases

- **Proven IFD**
- **Probable IFD**
- **Possible IFD**
- **EORTC/MSG criteria**

Ascioglu et al CID 2002; Ben de Pauw et al CID 2008

Classification

Invasive Fungal Diseases

- Proven IFD
 - ◆ Irrespective of host factors or clinical features
 - ◆ Demonstration of fungal elements in diseased tissue
 - ◆ If no culture, conclude to proven mold or yeast IFD
 - Except if histological appearance sufficiently distinctive (e.g. endemic mycoses such as Histoplasmosis, Coccidioidomycosis, Blastomycosis)
 - ◆ Individual IFD entities require culture and identification

Ben de Pauw et al CID 2008

Table 1: Criteria for proven mould infections

| Analysis and specimen | Molds ^a |
|--|---|
| Microscopic analysis: sterile material | Histopathologic, cytopathologic, or direct microscopic examination ^b of a specimen obtained by needle aspiration or biopsy in which hyphae or melanized yeast-like forms are seen accompanied by evidence of associated tissue damage |
| Culture | |
| Sterile material | Recovery of a mold or "black yeast" by culture of a specimen obtained by a sterile procedure from a normally sterile and clinically or radiologically abnormal site consistent with an infectious disease process, excluding bronchoalveolar lavage fluid, a cranial sinus cavity specimen, and urine |
| Blood | Blood culture that yields a mold ^d (e.g., <i>Fusarium</i> species) in the context of a compatible infectious disease process |
| Serological analysis: CSF | Not applicable |

Ben de Pauw et al CID 2008

Table 1: Criteria for proven mould infections

| Analysis and specimen | Molds ^a |
|--|---|
| Microscopic analysis: sterile material | Histopathologic, cytopathologic, or direct microscopic examination ^b Biopsies and culture by a sterile procedure which hyphae or melanized yeast-like forms are seen accompanied by evidence of associated tissue damage |
| Culture | |
| Sterile material | Recovery of a mold or "black yeast" by culture of a specimen obtained by a sterile procedure from a normally sterile and clinical site ^c |
| | ^d Recovery of <i>Aspergillus</i> species from blood invariably represents contamination |
| Blood | Blood culture that yields a mold ^d (e.g., <i>Fusarium</i> species) in the context of a compatible infectious disease process |
| Serological analysis: CSF | Not applicable |

Ben de Pauw et al CID 2008

Table 1: Criteria for proven yeast infections

| Analysis and specimen | Yeasts ^a |
|--|--|
| Microscopic analysis: sterile material | Histopathologic, cytopathologic, or direct microscopic examination ^b of a specimen obtained by needle aspiration or biopsy from a normally sterile site (other than mucous membranes) showing yeast cells—for example, <i>Cryptococcus</i> species indicated by encapsulated budding yeasts or <i>Candida</i> species showing pseudo-hyphae or true hyphae ^c |
| Culture | |
| Sterile material | Recovery of a yeast by culture of a sample obtained by a sterile procedure (including a freshly placed [<24 h ago] drain) from a normally sterile site showing a clinical or radiological abnormality consistent with an infectious disease process |
| Blood | Blood culture that yields yeast (e.g., <i>Cryptococcus</i> or <i>Candida</i> species) or yeast-like fungi (e.g., <i>Trichosporon</i> species) |
| Serological analysis: CSF | Cryptococcal antigen in CSF indicates disseminated cryptococcosis |

Ben de Pauw et al CID 2008

Table 1: Criteria for proven yeast infections

| Analysis and specimen | Yeasts ^a |
|--|---|
| Microscopic analysis: sterile material | <p>Histopathologic, cytopathologic, or direct microscopic examination^b of a specimen obtained by needle aspiration or biopsy from a mucous membrane) showing yeast cells—for example, <i>Cryptococcus</i> species indicated by encapsulated budding yeasts or <i>Candida</i> species showing pseudo-hyphae or true hyphae^c</p> <p>Biopsies and culture</p> |
| <p>Culture</p> <p>Sterile material</p> | <p>Recovery of a yeast by culture of a sample obtained by a sterile procedure (including a freshly placed [<24 h ago] drain) from a normally sterile site showing a clinical or radiological abnormality consistent with an infectious disease process</p> <p>Blood or sterile material culture</p> |
| Blood | <p>Blood culture that yields yeast (e.g., <i>Cryptococcus</i> or <i>Candida</i> species) or yeast-like fungi (e.g., <i>Trichosporon</i> species)</p> |
| Serological analysis: CSF | <p>Cryptococcal antigen in CSF indicates disseminated cryptococcosis</p> |

Ben de Pauw et al CID 2008

Table 1: Criteria for proven IFD foot notes

^a If culture is available append the identification at the genus or species level from the culture results

^b Tissue and cells submitted for histopathologic or cytopathologic studies should be stained by Grocott-Gomori methenamine silver stain or by periodic acid Schiff stain to facilitate inspection of fungal structures. Whenever possible, wet mounts of specimens from foci related to IFD should be stained with a fluorescent dye (e.g. calcofluor or blankophor)

Direct examination using a fluorescent dye

- « Aspergillus » like
- « Mucormycosis » like

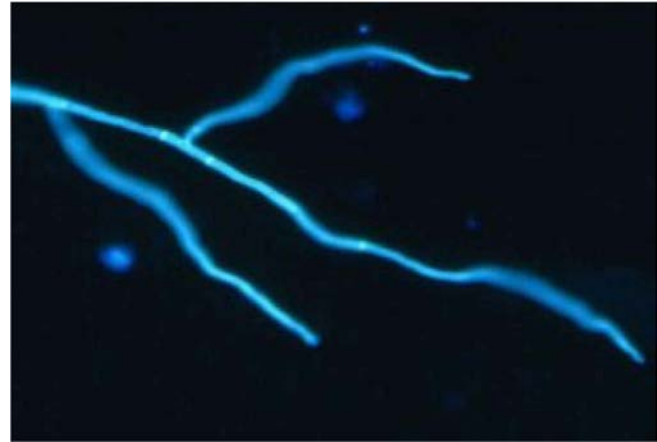


Table 2: Criteria for probable IFD

- Host factors
 - ◆ Neutropenia (< 500 neutrophils /mm³ for > 10 days) related to the onset of IFD
 - ◆ Allo-HSCT
 - ◆ Steroids (>0.3 mg/kg/d for >3 weeks)
 - ◆ Immunosuppressants (cyclosporine, anti-TNF, monoclonal Ab, nucleoside analogues) for the last 90 days
 - ◆ Inherited severe immunodeficiency

Table 2: Criteria for probable IFD

- Host factors **What about ICU patients?**
 - ◆ Neutropenia (< 500 neutrophils /mm³ for > 10 days) related to the onset of IFD
 - ◆ Allo-HSCT
 - ◆ Steroids (>0.3 mg/kg/d for >3 weeks)
 - ◆ Immunosuppressants (cyclosporine, anti-TNF, monoclonal Ab, nucleoside analogues) for the last 90 days
 - ◆ Inherited severe immunodeficiency

Table 2: Criteria for probable IFD

Clinical criteria^b

Lower respiratory tract fungal disease^c

The presence of 1 of the following 3 signs on CT:

- Dense, well-circumscribed lesions(s) with or without a halo sign
- Air-crescent sign
- Cavity

Tracheobronchitis

Tracheobronchial ulceration, nodule, pseudomembrane, plaque, or eschar seen on bronchoscopic analysis

Sinonasal infection

Imaging showing sinusitis plus at least 1 of the following 3 signs:

- Acute localized pain (including pain radiating to the eye)
- Nasal ulcer with black eschar
- Extension from the paranasal sinus across bony barriers, including into the orbit

CNS infection

1 of the following 2 signs:

- Focal lesions on imaging
- Meningeal enhancement on MRI or CT

Disseminated candidiasis^d

At least 1 of the following 2 entities after an episode of candidemia within the previous 2 weeks:

- Small, target-like abscesses (bull's-eye lesions) in liver or spleen
- Progressive retinal exudates on ophthalmologic examination

Table 2: Clinical criteria for probable IFD

Clinical criteria^b

Lower respiratory tract fungal disease^c

The presence of 1 of the following 3 signs on CT:

- Dense, well-circumscribed lesions(s) with or without a halo sign
- Air-crescent sign
- Cavity

Tracheobronchitis

Tracheobronchial ulceration, nodule, pseudomembrane, plaque, or eschar seen on bronchoscopic analysis

Sinonasal infection

Imaging showing sinusitis plus at least 1 of the following 3 signs:

- Acute localized pain (including pain radiating to the eye)
- Nasal ulcer with black eschar
- Extension from the paranasal sinus across bony barriers, including into the orbit

CNS infection

1 of the following 2 signs:

- Focal lesions on imaging
- Meningeal enhancement on MRI or CT

Disseminated candidiasis^d

At least 1 of the following 2 entities after an episode of candidemia within the previous 2 weeks:

- Small, target-like abscesses (bull's-eye lesions) in liver or spleen
- Progressive retinal exudates on ophthalmologic examination

Table 2: Criteria for probable IFD

Mycological criteria

Direct test (cytology, direct microscopy, or culture)

Mold in sputum, bronchoalveolar lavage fluid, bronchial brush, or sinus aspirate samples, indicated by 1 of the following:

Presence of fungal elements indicating a mold

Recovery by culture of a mold (e.g., *Aspergillus*, *Fusarium*, *Zygomycetes*, or *Scedosporium* species)

Indirect tests (detection of antigen or cell-wall constituents)^e

Aspergillosis

Galactomannan antigen detected in plasma, serum, bronchoalveolar lavage fluid, or CSF

Invasive fungal disease other than cryptococcosis and zygomycoses

β -D-glucan detected in serum

Table 2: Criteria for probable IFD

Mycological criteria

Direct test (cytology, direct microscopy and culture)

Cytology, direct microscopy and culture

Mold in sputum

Indicated by 1 of the following:

Presence of fungal elements indicating a mold

Recovery by culture of a mold (e.g., *Aspergillus*, *Fusarium*, *Zygomycetes*, or *Scedosporium* species)

Indirect tests (detection of antigen or cell-wall constituents)^e

Aspergillosis

Galactomannan antigen detected in plasma, serum, bronchoalveolar lavage fluid, or CSF

Invasive fungal disease other than cryptococcosis and zygomycoses

β -D-glucan detected in serum

Indirect tests: galactomannan and β -D-glucan

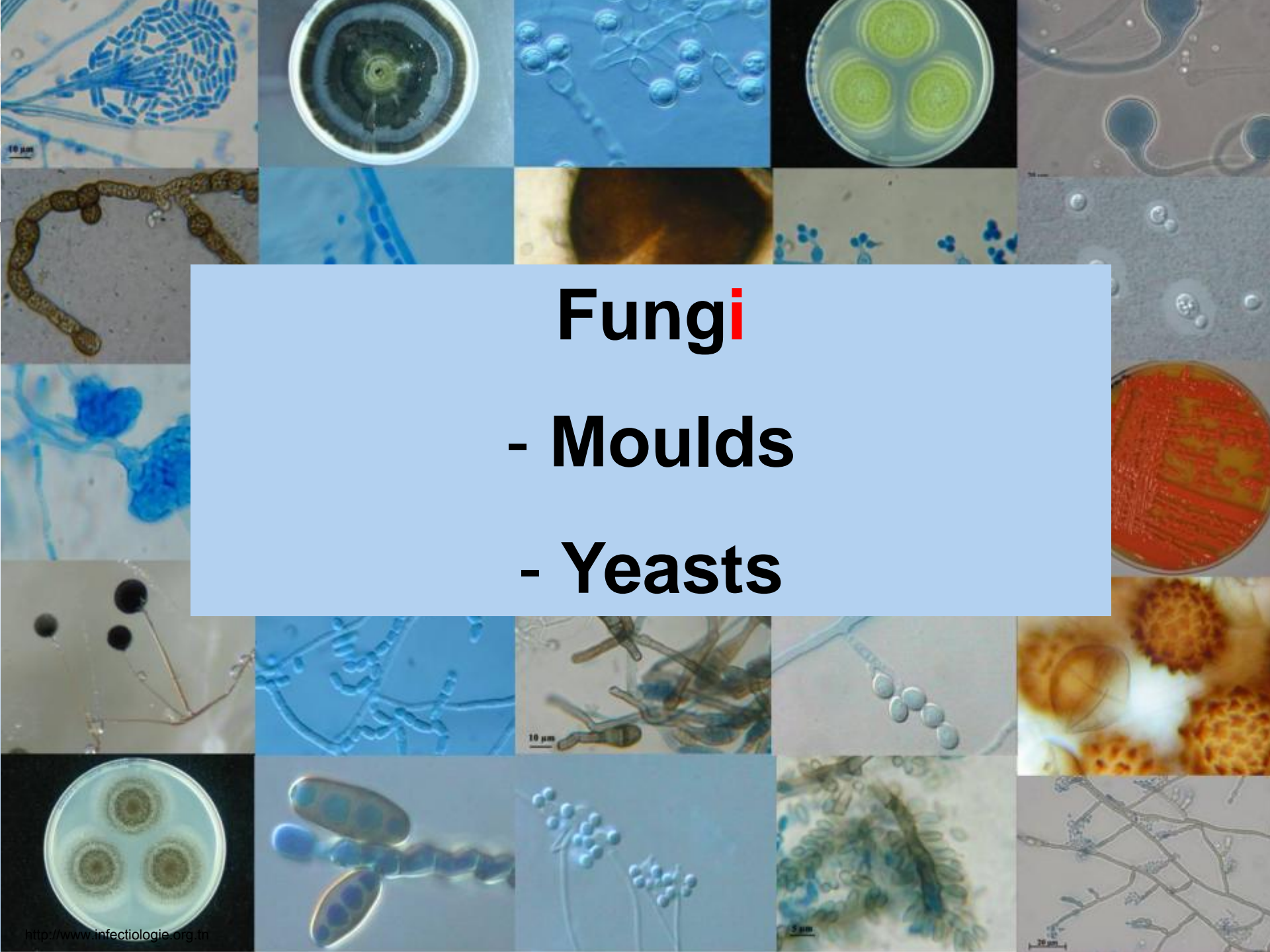
Definition: probable or possible IFD

PROBABLE =
HOST FACTOR
+ CLINICAL CRITERION
+ MYCOLOGICAL CRITERION

POSSIBLE =
HOST FACTOR
+ CLINICAL CRITERION

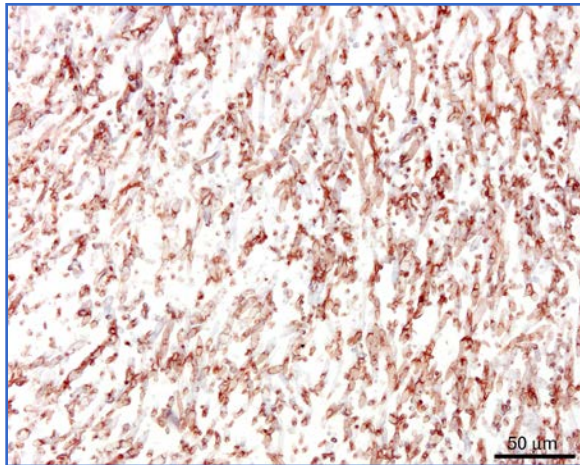
“New” diagnostic tools



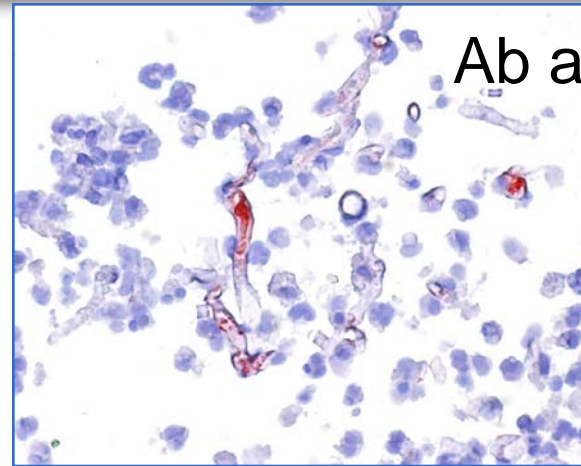


Fungi
- Moulds
- Yeasts

Biopsies: Immunostaining

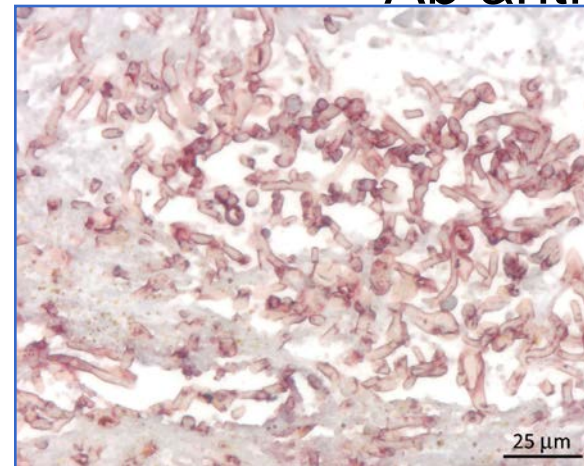


Ab anti-*Aspergillus*



Ab anti-*Rhizomucor*

Ab anti-*Candida*



G. Jouvion, F. Chrétien
Institut Pasteur
Human Histopathology & Animal Models
Infection and Epidemiology Department

Molecular ID from biopsies

- Common practice
 - ◆ DNA extraction
 - ◆ Amplification of ITS regions
- Pitfalls
 - ◆ Poor DNA quality from formalin fixed tissues (ask for -80° C)
 - ◆ At least 40% of false identification in public data base (GenBank)
 - ◆ Hybridization of primers with human DNA (hence low sensitivity)

- ITS1 (TCCGTAGGTGAACCTGCGG)
- 19 nucleotides
- Identities = 19/19 (100%), Gaps = 0/19 (0%)
- Query 1 TCCGTAGGTGAACCTGCGG
- | | | | | | | | | | | | | | | | | | | |
- Sbjct 110917 TCCGTAGGTGAACCTGCGG

- > Homo sapiens unplaced genomic contig,
alternate assembly HuRef

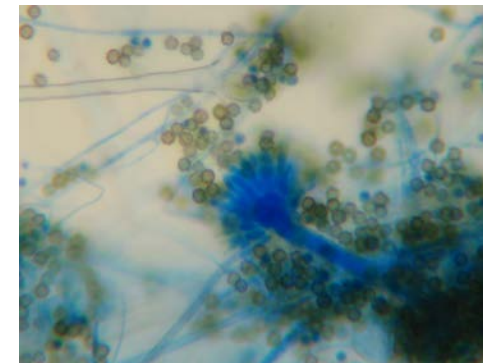
White T, et al In PCR-protocols a guide to methods and applications. Academic press: 1990:315-322

Importance of fungal culture for species identification

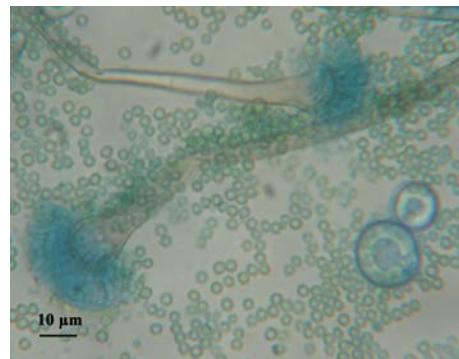
| Species | % (n=246) |
|----------------------|-----------|
| <i>A. fumigatus</i> | 85 |
| <i>A. flavus</i> | 4 |
| <i>A. nidulans</i> | 3 |
| <i>A. terreus</i> | 2 |
| <i>A. niger</i> | 4 |
| <i>A. ustus</i> | 0.5 |
| <i>A. versicolor</i> | 0.5 |
| Others | 1 |



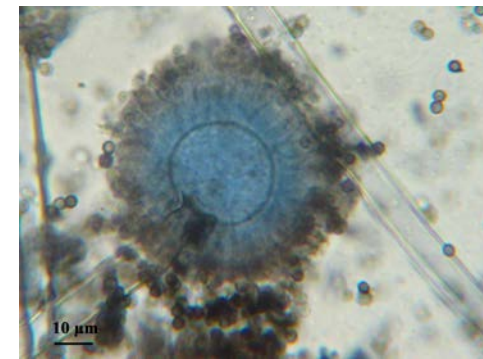
A. terreus



A. ustus



A. nidulans



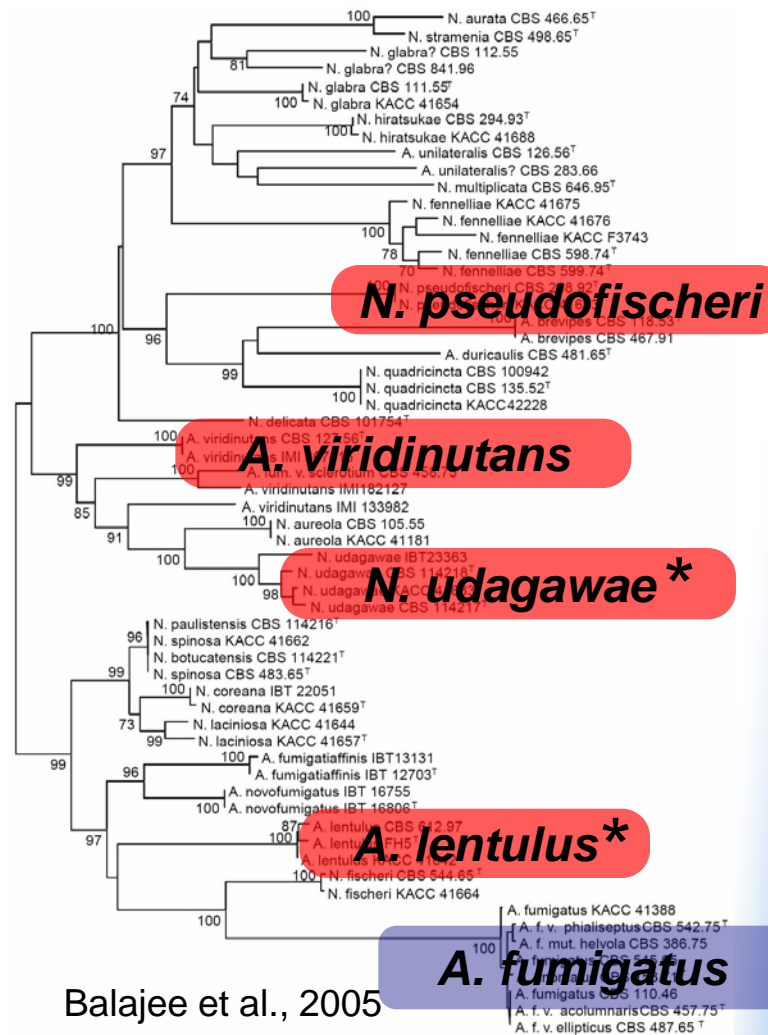
A. niger

Lortholary et al CMI 2011

Aspergilli taxonomy

Innate
resistance to
azoles

Innate susceptibility to azoles



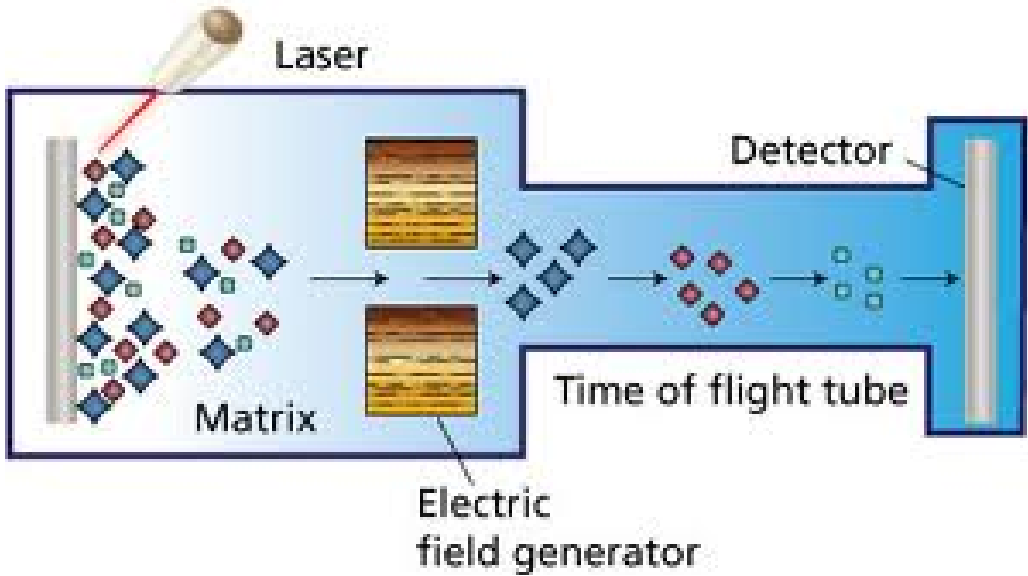
Balajee et al., 2005

Innate resistance

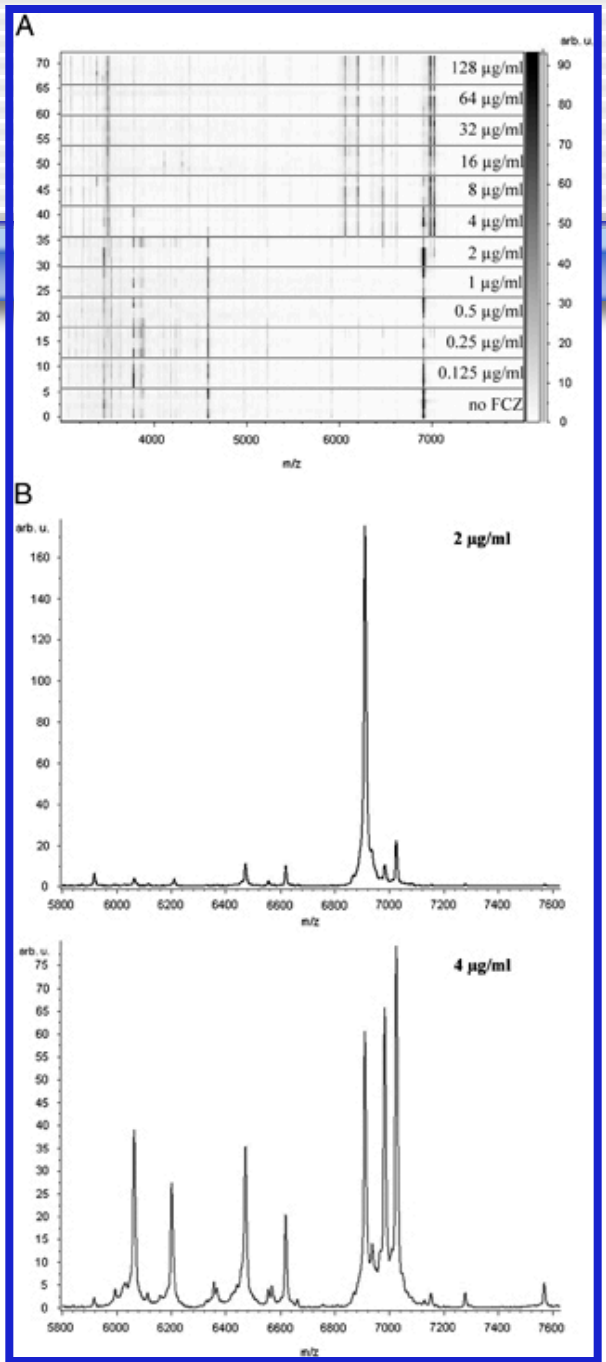
| species | Section | AmB | Azoles | Candines |
|---------------------------|-----------------|---------------------|---------------------|---------------------|
| <i>A. fumigatus</i> | <i>Fumigati</i> | Resistant (Red) | Susceptible (Green) | Susceptible (Green) |
| <i>A. lentulus</i> | | Resistant (Red) | Resistant (Red) | Resistant (Red) |
| <i>A. fumigatiaffinis</i> | | Resistant (Red) | Susceptible (Green) | Susceptible (Green) |
| <i>A. viridinutans</i> | | Susceptible (Green) | Resistant (Red) | Susceptible (Green) |
| <i>A. fumisynnematus</i> | | Susceptible (Green) | Susceptible (Green) | Susceptible (Green) |
| <i>N. fischeri</i> | | Susceptible (Green) | Susceptible (Green) | Susceptible (Green) |
| <i>N. pseudofischeri</i> | | Susceptible (Green) | Resistant (Red) | Susceptible (Green) |
| <i>N. udagawae</i> | | Resistant (Red) | Resistant (Red) | Susceptible (Green) |
| <i>N. fennelliae</i> | | Susceptible (Green) | Susceptible (Green) | Susceptible (Green) |
| <i>N. hiratsukae</i> | | Susceptible (Green) | Susceptible (Green) | Susceptible (Green) |
| <i>N. spinosa</i> | | Susceptible (Green) | Susceptible (Green) | Susceptible (Green) |

Alcazar-Fuoli et al. 2008, Balajee et al. 2006

MALDI-TOF



Phenotypic method

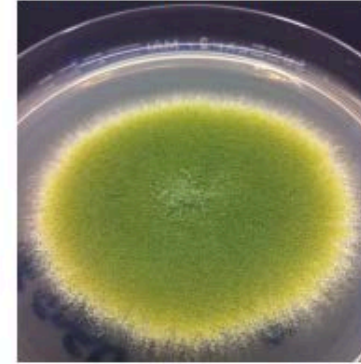


After 24h incubation

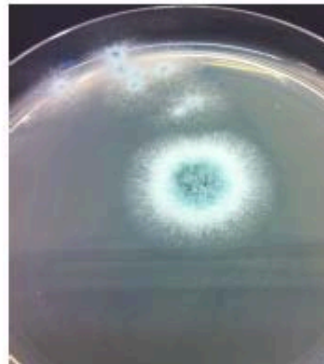
After 48h incubation

After 5 days incubation

Aspergillus flavus



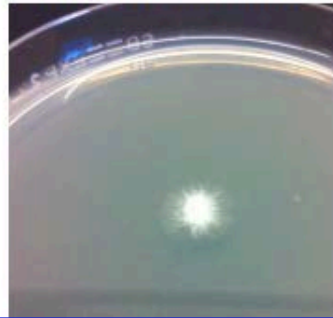
Aspergillus fumigatus



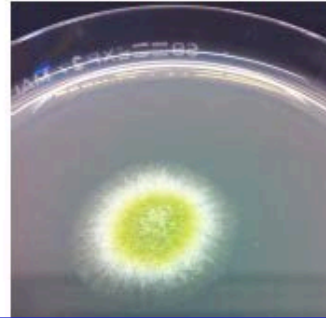
Schrenzel ECCMID 2012, London

Aspergillus flavus

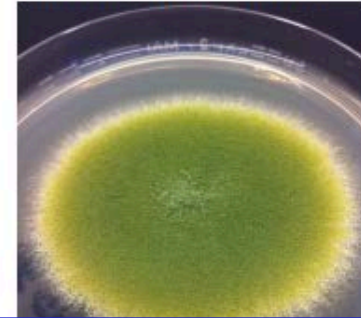
After 24h incubation



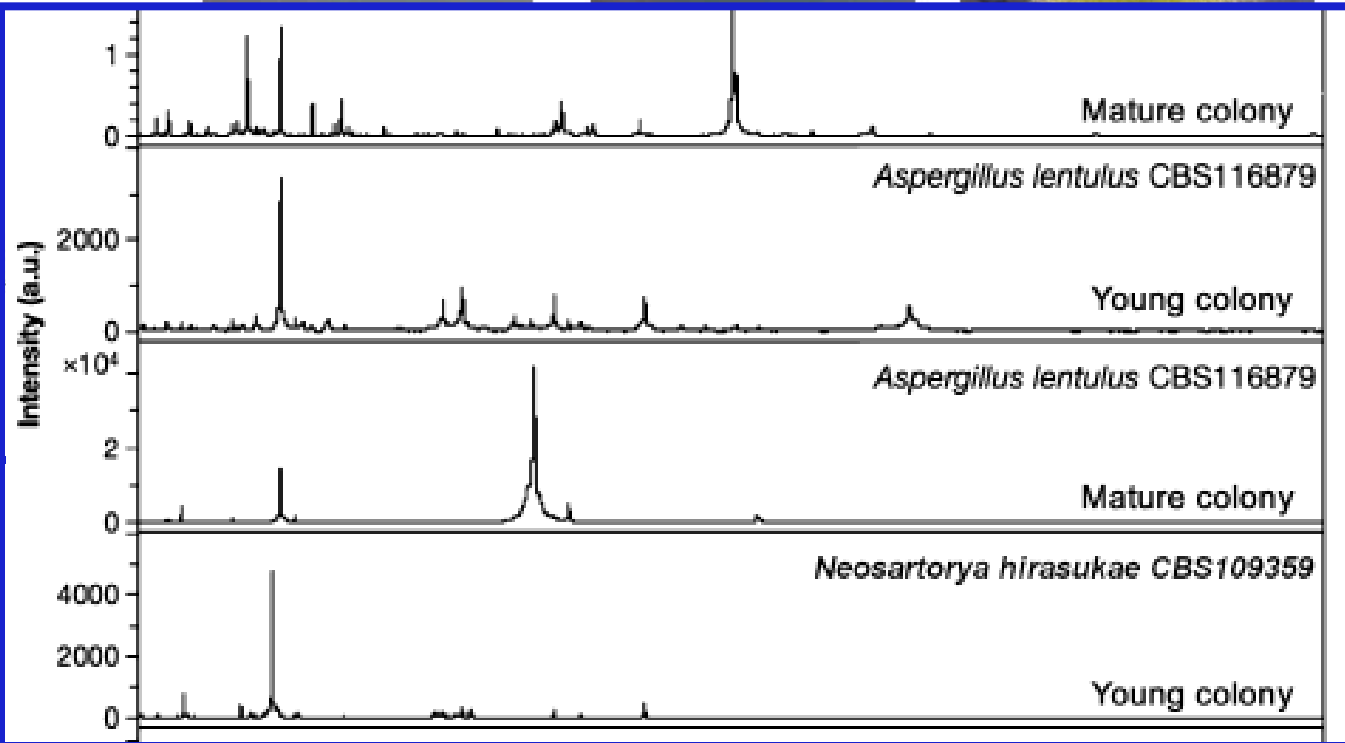
After 48h incubation



After 5 days incubation



Aspergillus



Design specific age-dependent data banks

MALDI-TOF



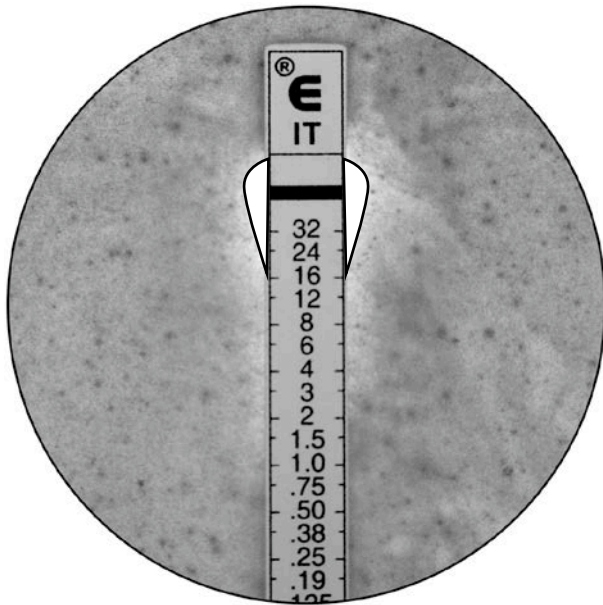
- Easier for yeast ID
- Direct ID in positive blood culture



Ferroni et al JCM 2010; Spanu et al JCM 2012

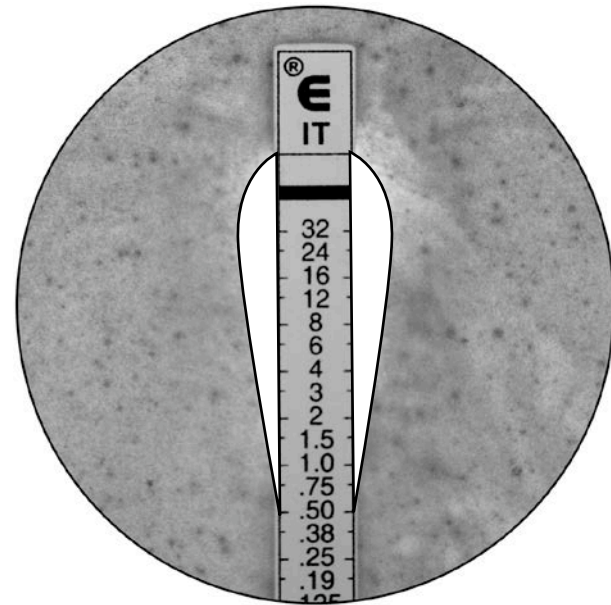
MIC (Etest®) for molds

Resistant



CMI = 16 mg/l

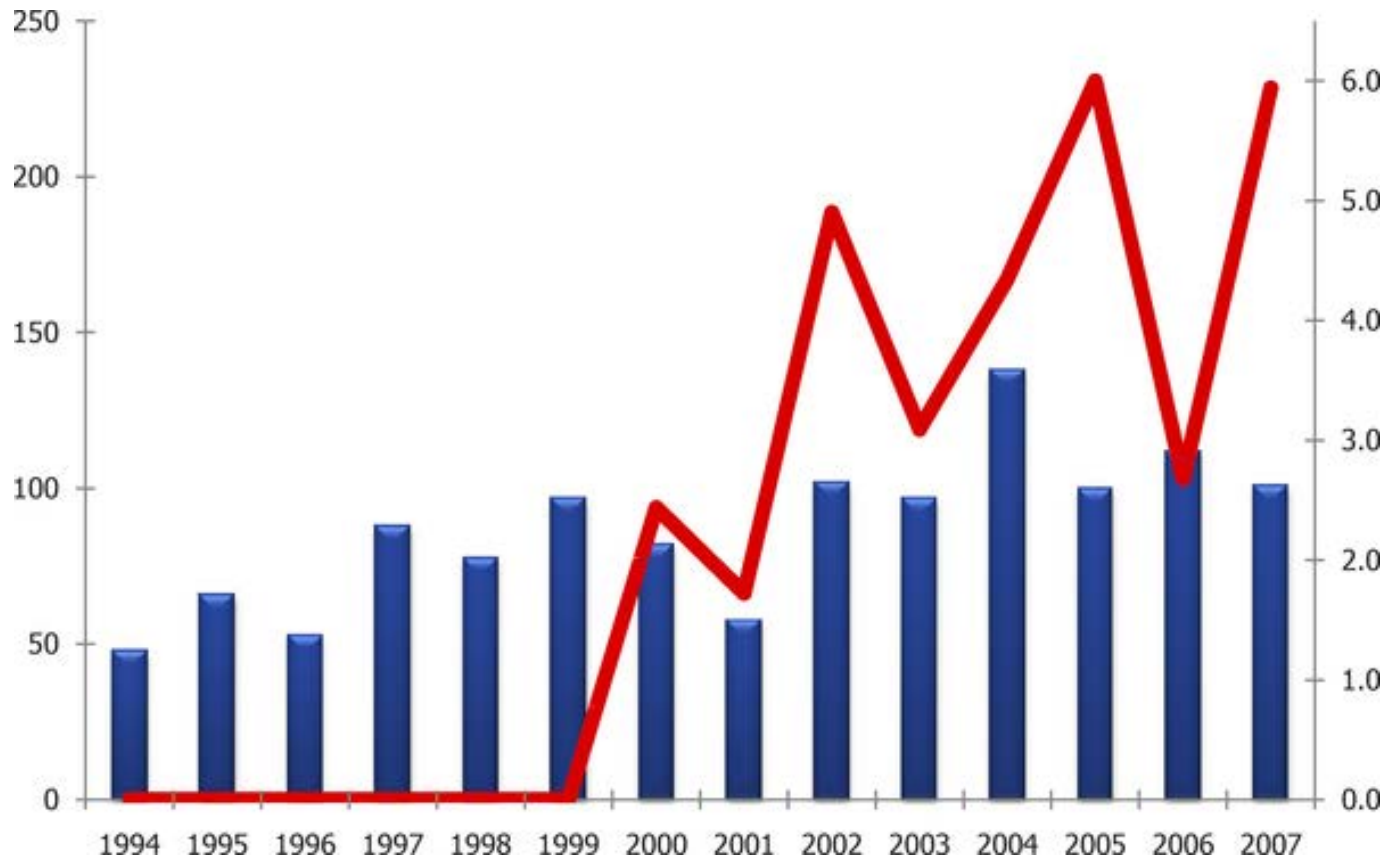
Susceptible



CMI = 0.5 mg/l

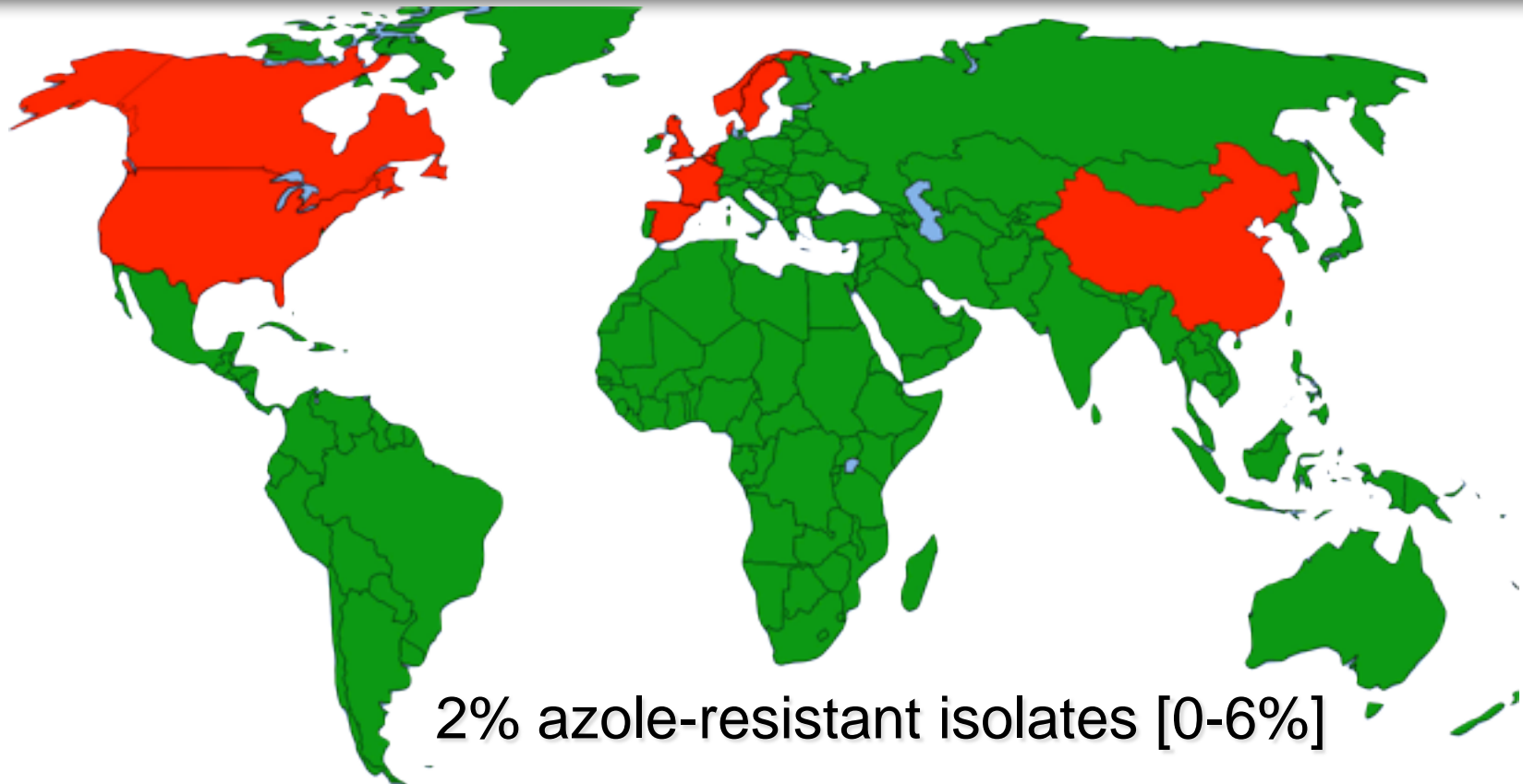
EUCAST values: Arendrup et al Clin Microbiol Inf March 2012

Increasing acquired resistance in *A. fumigatus*



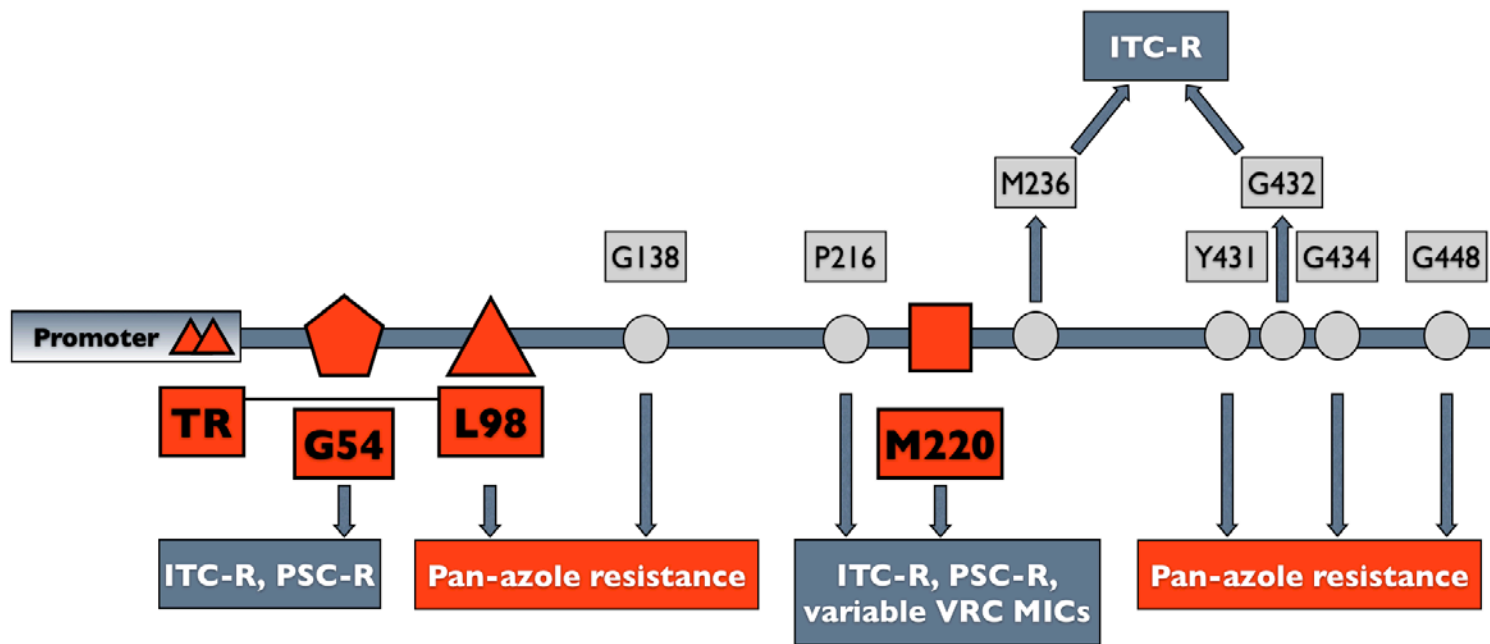
Snelders et al, Plos Medicine 2008

Epidemiological data



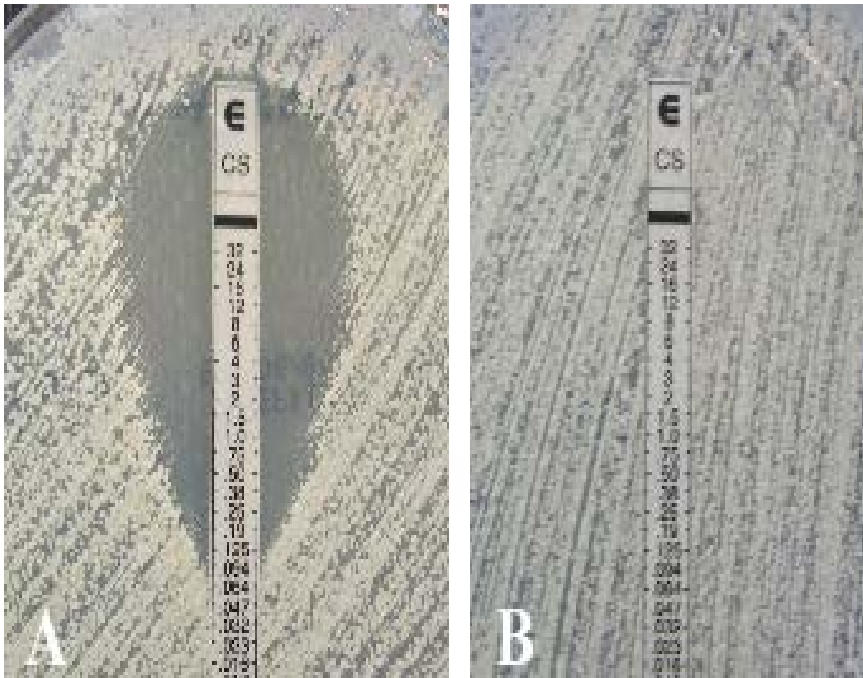
*Alanio et al. J. Antimicrob. Chemother. 2010; Bueid et al. J. Antimicrob. Chemother. 2010
Howard et al. Med Mycol. 2011, Lockhart et al. AAC. 2011; Van der Linden et al. EID 2011*

Resistance mechanisms in *A. fumigatus*

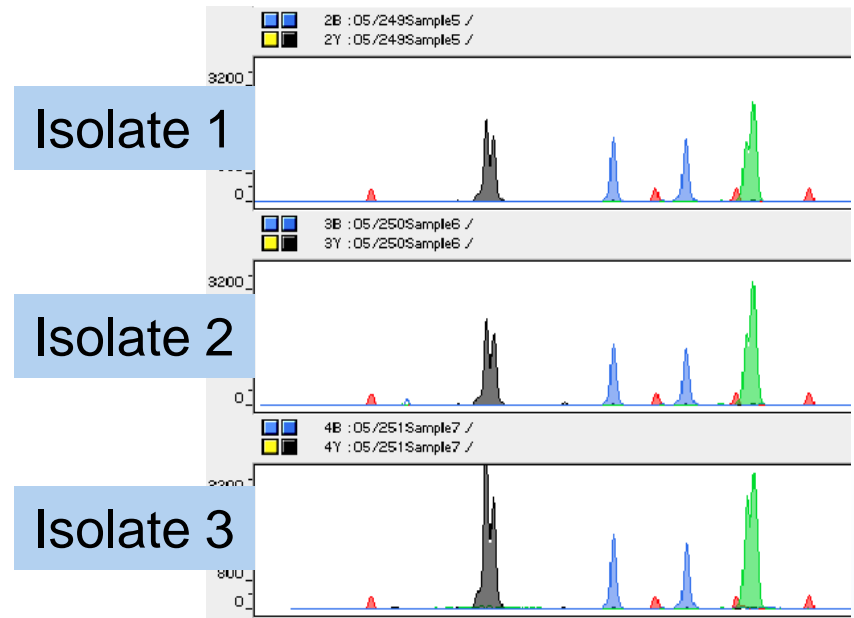


Alanio et al. *Curr Fungal Infect Rep* 2011; Howard et al. *Med Mycol* 2011; Denning et al. *CID* 2011

MIC (Etest®) for yeasts



Acquired resistance under antifungal pressure



Microsatellite genotyping

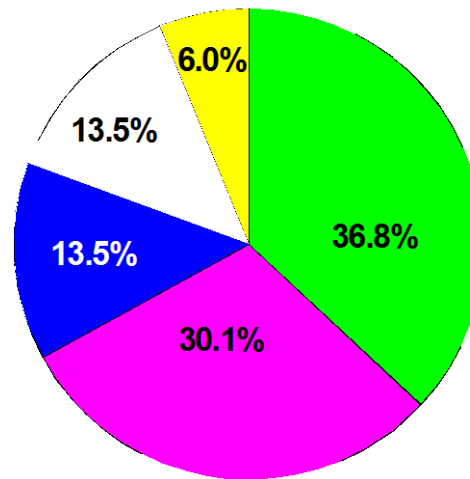
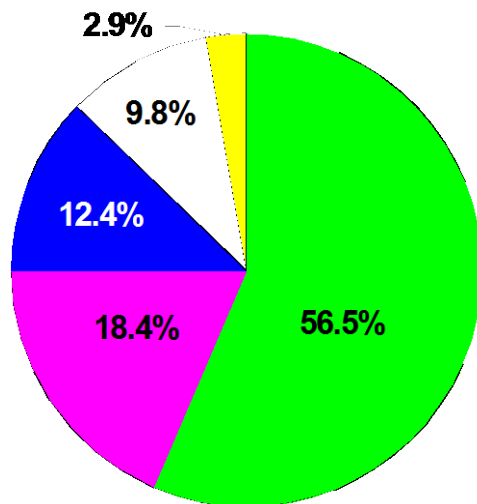
Baixench et al JAC 2007

Recent (within 30 previous days) exposure

None recorded (n=1821)

Recorded (n=133)

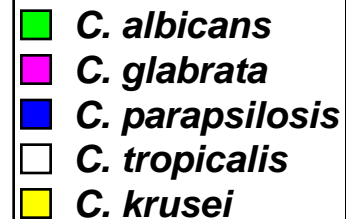
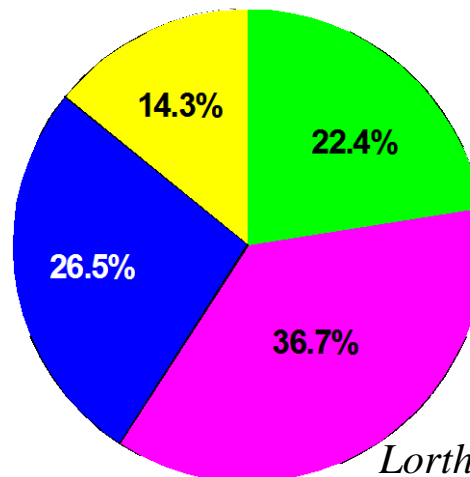
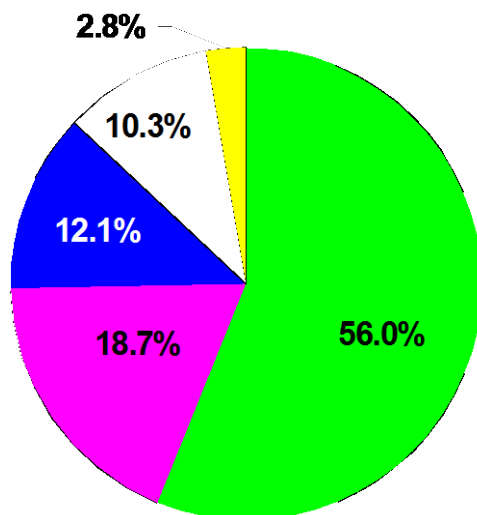
Fluconazole
(P=0.001)



None recorded (n=1905)

Recorded (n=49)

Caspofungin
(P<0.001)



Current biomarkers

- Ag
 - ◆ GM
 - ◆ β -D-glucan
 - ◆ Mn
 - ◆ GMX cryptococcus
 - ◆ *Histoplasma* sp.
- DNA
 - ◆ *Aspergillus*
 - ◆ *Candida*

Galactomannan: ELISA (BioRad)

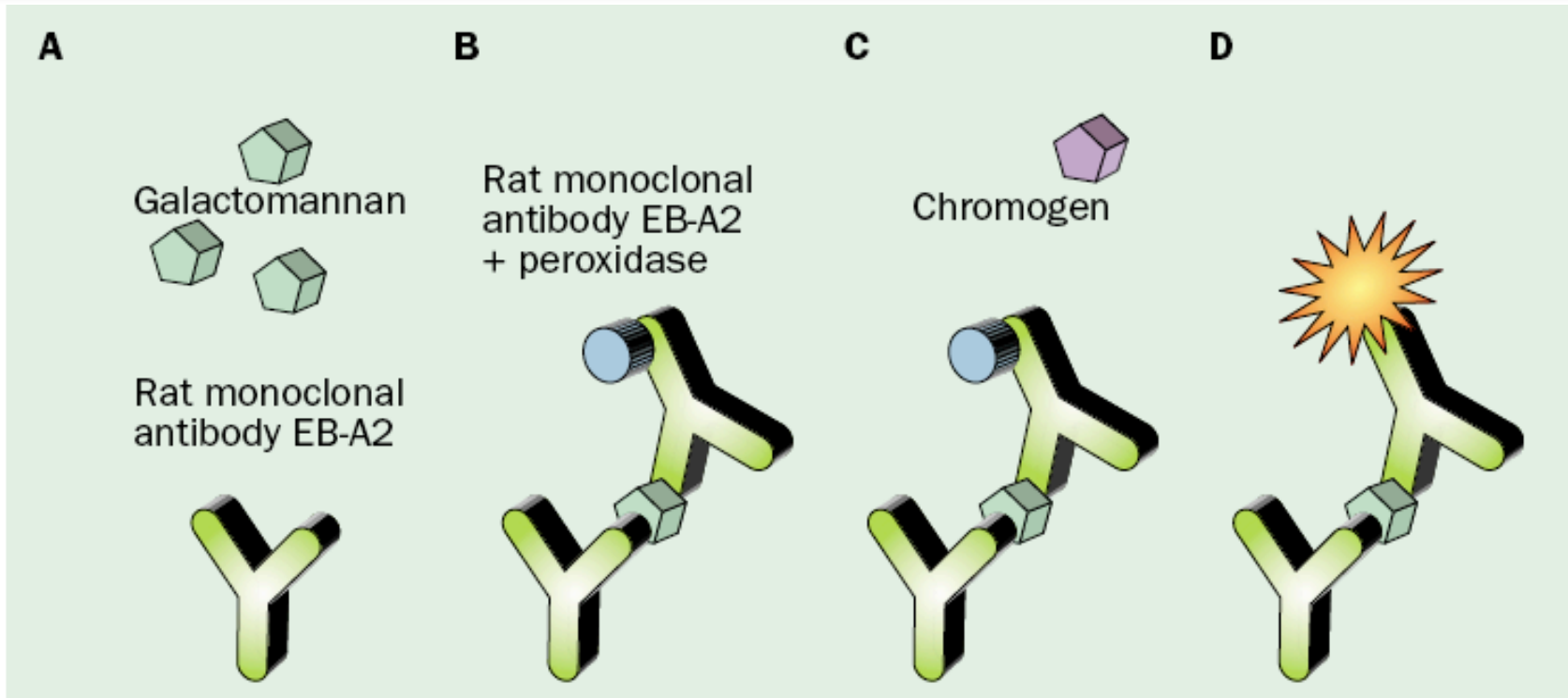
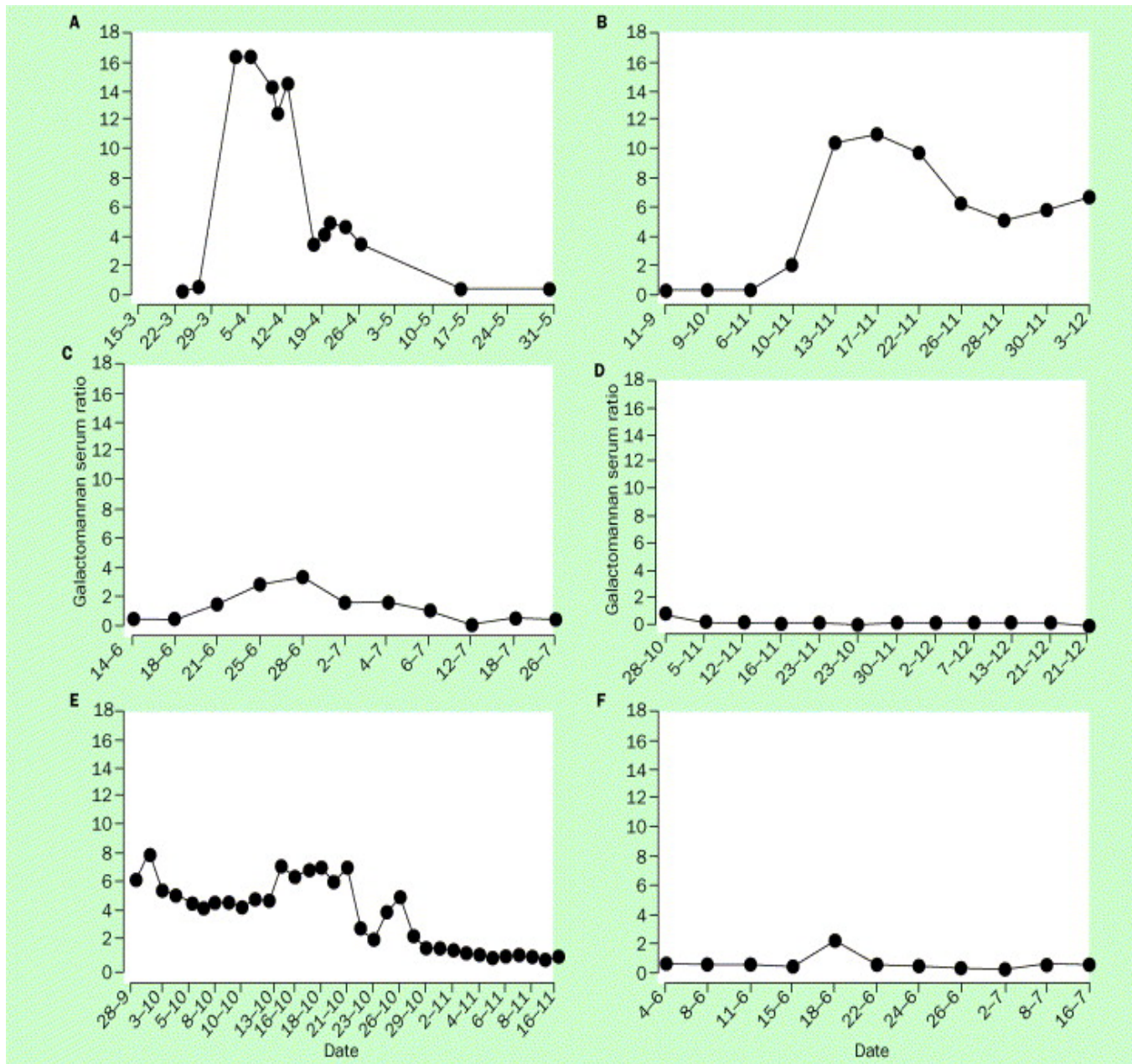


Figure 1. The *Platelia Aspergillus* ELISA technique. A serum ratio is calculated by dividing the optical density of the patient's serum sample by the mean optical density of two threshold control samples that contain 1 $\mu\text{g/L}$ of galactomannan.

Mennink-Kersten et al, *Lancet Inf Dis*, 2004



Mennink-Kersten et al, *Lancet Inf Dis*, 2004

GM recommendations for strategy in adults (ECIL3)

Prospective monitoring of serum* is a feasible approach in adult neutropenic patients undergoing intensive chemotherapy for leukemia or receiving an allogeneic stem cell transplantation for the early diagnosis of invasive aspergillosis (AII)

GM monitoring is recommended every three to four days in admitted patients (AII)

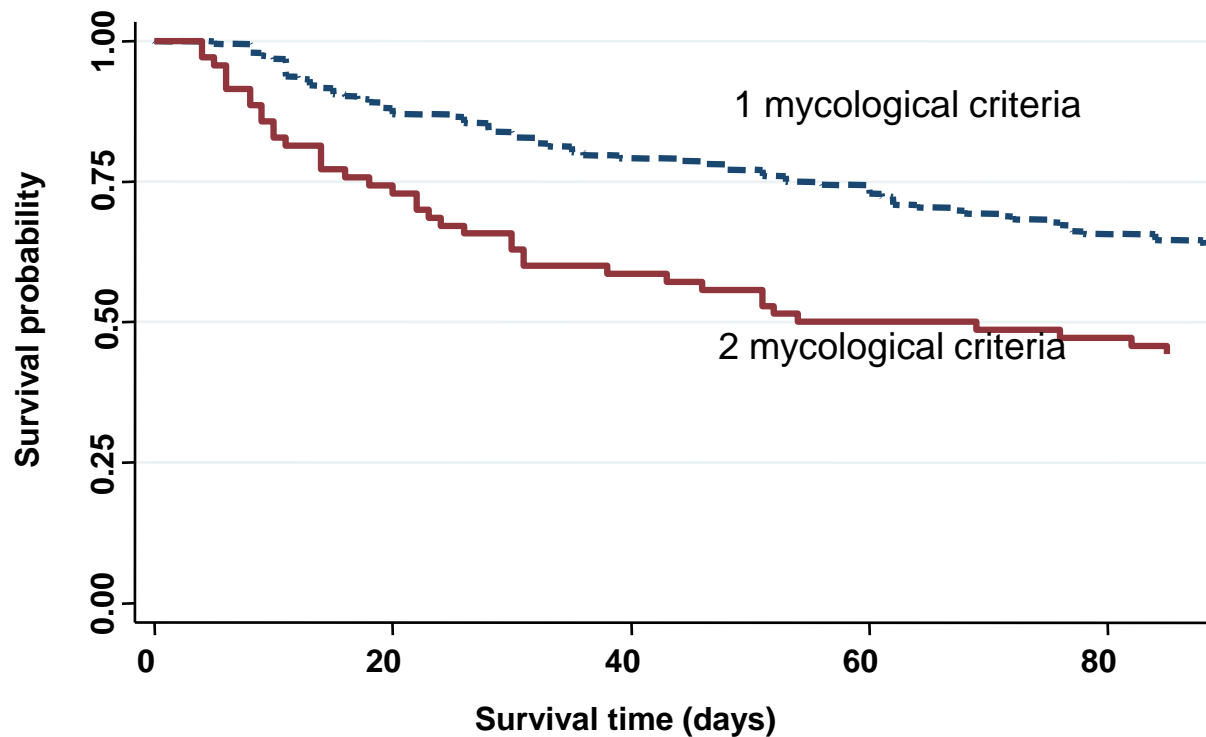
In GM positive patients persistent GM antigenemia during therapy is a poor prognostic sign and should prompt a reassessment of the management of the patient (BII)

A diagnostic driven strategy that incorporates GM monitoring should be combined with high resolution CT imaging, appropriate clinical and microbiological evaluation to early diagnose invasive aspergillosis. A single positive GM index of ≥ 0.7 or 2 consecutive samples of ≥ 0.5 should prompt a diagnostic work-up (AII)

* Plasma may also be used (CIII)

*Marchetti et al BMT 2011
European Conference on Infections in Leukaemia*

Prognosis value of combined diagnostic means



*National Reference Centre for Mycology and Antifungals
Lortholary et al CMI 2011*

Factors that influence GM performance

Table 2 Biological and epidemiological factors that influence the performance of GM detection in invasive aspergillosis³

| <i>Biological factors</i> | <i>Epidemiological factors</i> |
|--|---|
| Site of infection | Patient population |
| <i>Aspergillus</i> species causing infection | Sampling strategy |
| Microenvironment at the site of infection: nutrients, oxygen level, pH | Definition of a positive result |
| Exposure to antifungal agents | Definition of an IFD |
| Molecular structure of released galactomannan | Prevalence of IFD |
| Underlying condition/neutropenia/level of immunosuppression | Cutoff for positivity |
| Renal clearance, hepatic metabolism | Laboratory experience |
| Circulating galactomannan antibodies | Nutritional factors (galactomannan-containing food) |
| Storage of clinical sample | Treatment with semi-synthetic β -lactam antibiotics |
| Pre-analytical treatment procedure | |

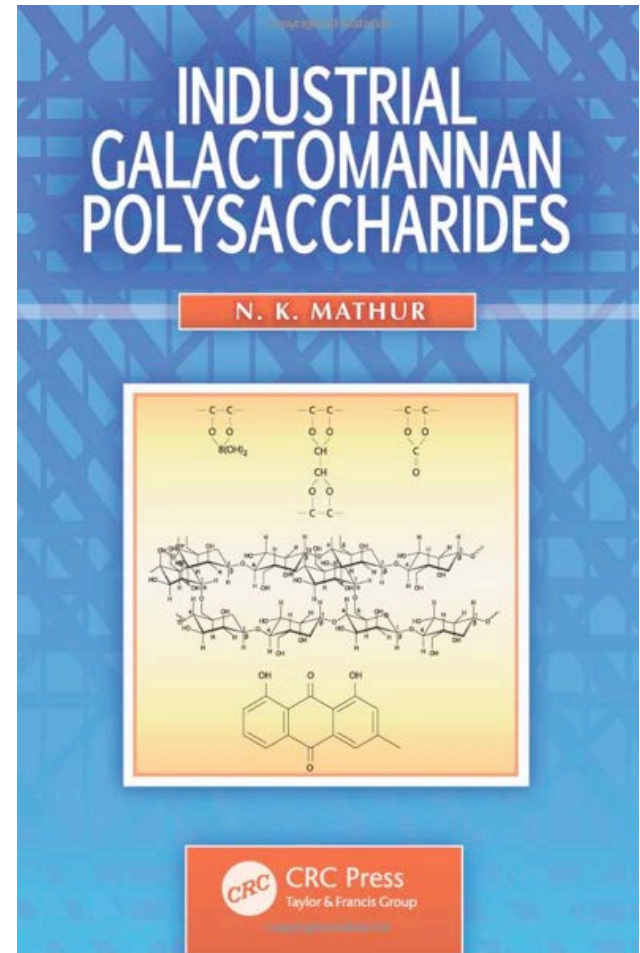
Factors that influence GM performance

Table 2 Biological and epidemiological factors that influence the performance of GM detection in invasive aspergillosis³

| <i>Biological factors</i> | <i>Epidemiological factors</i> |
|--|---|
| Site of infection | Patient population |
| <i>Aspergillus</i> species causing infection | Sampling strategy |
| Microenvironment at the site of infection: nutrients, oxygen level, pH | Definition of a positive result |
| Exposure to antifungal agents | Definition of an IFD |
| Molecular structure of released galactomannan | Prevalence of IFD |
| Underlying condition/neutropenia/level of immunosuppression | Cutoff for positivity |
| Renal clearance, hepatic metabolism | Laboratory experience |
| Circulating galactomannan antibodies | Nutritional factors (galactomannan-containing food) |
| Storage of clinical sample | Treatment with semi-synthetic β -lactam antibiotics |
| Pre-analytical treatment procedure | |

Galactomannan

Food-processing using galactomannan to modify food texture



Factors that influence GM performance

Table 2 Biological and epidemiological factors that influence the performance of GM detection in invasive aspergillosis³

| <i>Biological factors</i> | <i>Epidemiological factors</i> |
|--|---|
| Site of infection | Patient population |
| <i>Aspergillus</i> species causing infection | Sampling strategy |
| Microenvironment at the site of infection: nutrients, oxygen level, pH | Definition of a positive result |
| Exposure to antifungal agents | Definition of an IFD |
| Molecular structure of released galactomannan | Prevalence of IFD |
| Underlying condition/neutropenia/level of immunosuppression | Cutoff for positivity |
| Renal clearance, hepatic metabolism | Laboratory experience |
| Circulating galactomannan antibodies | Nutritional factors (galactomannan-containing food) |
| Storage of clinical sample | Treatment with semi-synthetic β -lactam antibiotics |
| Pre-analytical treatment procedure | |

GM specificity

- Numerous GM-producing fungal species

Fungus

Acremonium species
Alternaria alternata
Botrytis tulipae
Cladosporium cladosporioides
Cladosporium herbarum
Cryptococcus neoformans
Fusarium oxysporum
(but not *Fusarium solani*)
Geotricum capitatum
Paecilomyces variotii
Penicillium chrysogenum
Penicillium digitatum
Penicillium marneffeii
Rhodotorula rubra
Trichophyton interdigitalis
Trichophyton rubrum
Wallemia sebi
Wangiella (Exophiala) dermatitidis

Aquino, VR et al, *Mycopathologia*, 2007

When facing a positive GM result

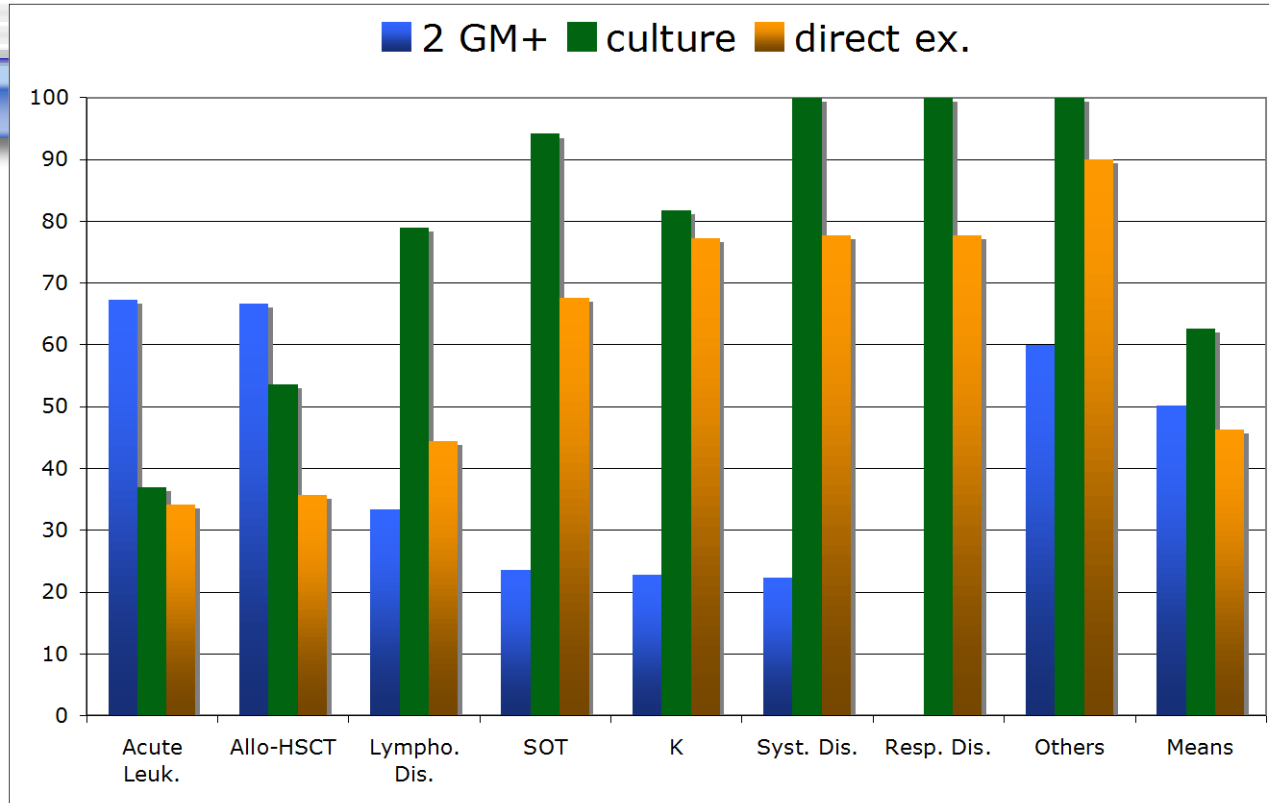
- Disease
 - ◆ To gather and to analyze EORTC/MSG criteria
- In parallel, explore the other possible sources
 - ◆ Antibiotics (test batches if necessary)
 - ◆ Other species than *Aspergillus* spp.
 - ◆ Mucites, gastrointestinal diseases
 - ◆ Intravenous products

Factors that influence GM performance

Table 2 Biological and epidemiological factors that influence the performance of GM detection in invasive aspergillosis³

| <i>Biological factors</i> | <i>Epidemiological factors</i> |
|--|---|
| Site of infection | Patient population |
| <i>Aspergillus</i> species causing infection | Sampling strategy |
| Microenvironment at the site of infection: nutrients, oxygen level, pH | Definition of a positive result |
| Exposure to antifungal agents | Definition of an IFD |
| Molecular structure of released galactomannan | Prevalence of IFD |
| Underlying condition/neutropenia, level of immunosuppression | Cutoff for positivity |
| Renal clearance, hepatic metabolism | Laboratory experience |
| Circulating galactomannan antibodies | Nutritional factors (galactomannan-containing food) |
| Storage of clinical sample | Treatment with semi-synthetic β -lactam antibiotics |
| Pre-analytical treatment procedure | |

Diagnostic means



Decrease of GM yield in Lymphoprolif. Dis. and other categories¹

- poor performances in SOT²

- neutropenia/steroid ratio³

- diagnostic attitude for not performing culture when GM+?

(1) Lortholary et al CMI 2011 (2) Pfeiffer et al, CID, 2006 ; (3) Cordonnier et al, CMI 2009

Galactomannan indices in neutropenia (group 1; PMN < 100/mm³) vs non-neutropenia (groups 2+3; PMN ≥ 100/mm³)

| Galactomannan index | Group 1 (n = 17) PNN < 100 | Groups 2 + 3 (n = 81) PNN ≥ 100 | p value |
|---|----------------------------------|---------------------------------------|-------------------|
| GM index ≥ 1 | 8 (44.4%) | 8 (9.9%) | .001 ^b |
| GM index ≥ 0.7 | 8 (44.4%) | 12 (14.8%) | .009 ^b |
| GM index ≥ 0.5 | 11 (61.1%) | 15 (18.52%) | .001 ^b |
| GM index, mean ± SD | 1.71 ± 1.99 | .44 ± .75 | .01 ^a |
| Steroid administration | | | |
| Yes | 4.95 ± .64 | .39 ± .44 | .001 ^a |
| No | .72 ± .79 | .47 ± .88 | .19 ^a |
| Potentially GM contaminated antibiotic(s) | | | |
| Yes | 2.6 ± 2.97 | .64 ± 1.11 | .76 ^a |
| No | 1.44 ± 1.66 | .33 ± .41 | .003 ^a |
| Anti-mold therapy | | | |
| Yes | .76 ± .99 | .43 ± .66 | .694 ^a |
| No | 2.1 ± 2.20 | .44 ± .77 | .006 ^a |

NOTE. Data are no (%) of episodes

^a p value of the non parametric Kruskal Wallis t est

^b p value of Fischer's exact test

<http://www.infectiologie.org.tn>

C. Cordonnier et al, CMI 2009; Balloy et al Infect Immunity 2005; Stergiopoulou et al Am J Clin Pathol 2007

Galactomannan indices in neutropenia (group 1; PMN < 100/mm³) vs non-neutropenia (groups 2+3; PMN ≥ 100/mm³)

| Galactomannan index | Group 1 (n = 17) PNN < 100 | Groups 2 + 3 (n = 81) PNN ≥ 100 | p value |
|--|----------------------------------|---------------------------------------|-------------------|
| GM index ≥ 1 | 8 (44.4%) | 8 (9.9%) | .001 ^b |
| GM index ≥ 0.7 | 8 (44.4%) | 12 (14.8%) | .009 ^b |
| GM index ≥ 0.5 | 11 (61.1%) | 15 (18.52%) | .001 ^b |
| GM index, mean ± SD | 1.71 ± 1.99 | .44 ± .75 | .01 ^a |
| Steroid administration | | | |
| Yes | 4.95 ± .64 | .39 ± .44 | .001 ^a |
| No | .72 ± .79 | .47 ± .88 | .19 ^a |
| Potentially GM contaminated antibiotic(s) | | | |
| Yes | 2.6 ± 2.97 | .64 ± 1.11 | .76 ^a |
| No | 1.44 ± 1.66 | .33 ± .41 | .003 ^a |
| Anti-mold therapy | | | |
| Yes | .76 ± .99 | .43 ± .66 | .694 ^a |
| No | 2.1 ± 2.20 | .44 ± .77 | .006 ^a |

NOTE. Data are no (%) of episodes

^a p value of the non parametric Kruskal Wallis t est

^b p value of Fischer's exact test

<http://www.infectiologie.org.tn>

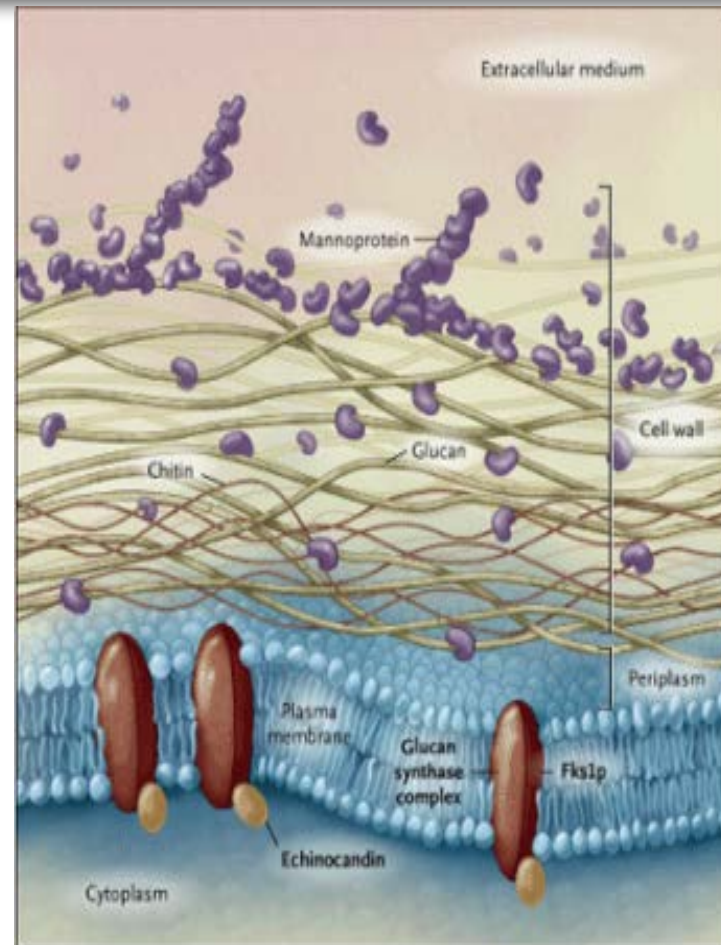
C. Cordonnier et al, CMI 2009; Balloy et al Infect Immunity 2005; Stergiopoulou et al Am J Clin Pathol 2007

GM in BAL

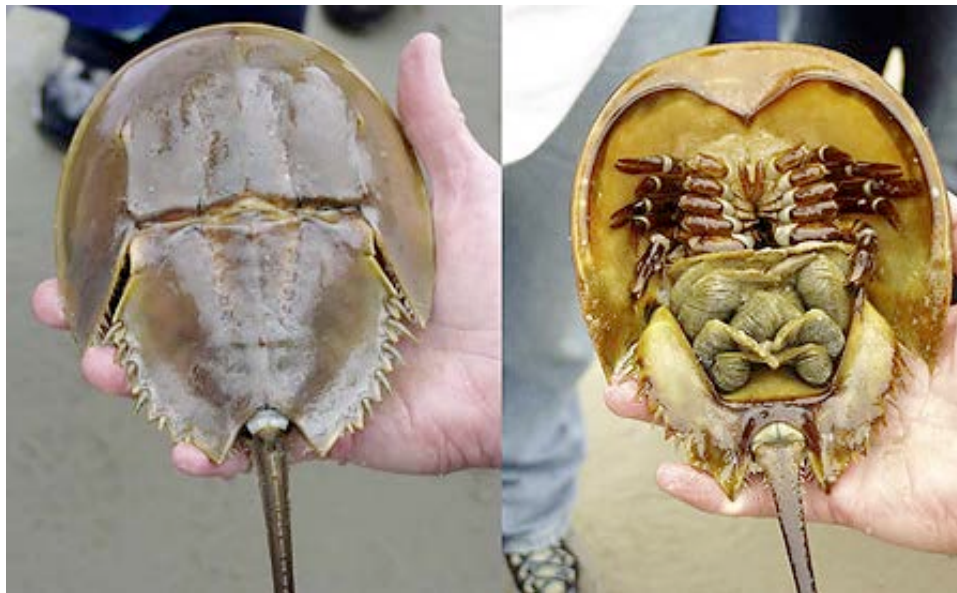
- GM > 0.5 23/26 1st BAL
 - ♦ BAL procedure
 - ♦ 2 x 20 ml
- 6 false GM+ (13%)
 - ♦ Standards?
 - ♦ Colonization or infection?
- 15/26 positive direct examination or positive culture (58%)
 - ♦ direct examination or positive culture
- GM > 0.5 42% serum samples
 - ♦ Useful for follow-up
 - ♦ Less disputable meaning
- Reproducibility? Transfer in other centres?

(1,3)beta-D-Glucan

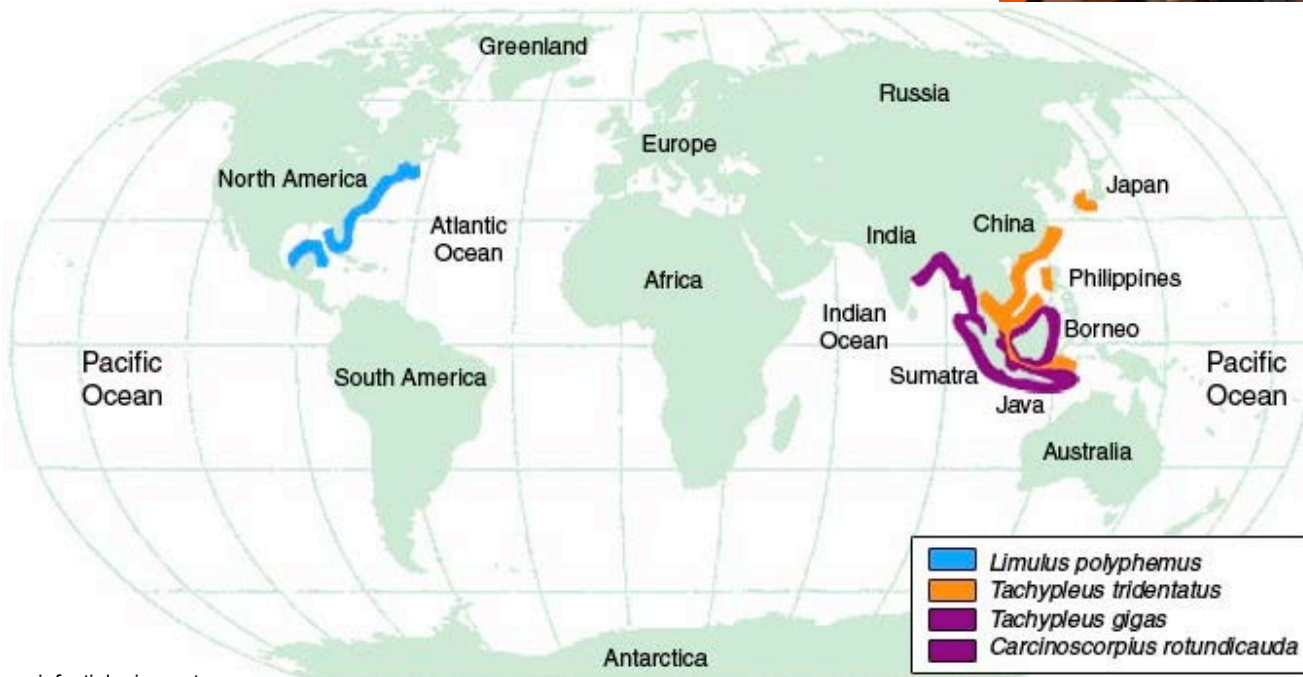
- Therapeutic target of echinocandins
- Ag common to most of the fungal species (excepted *Cryptococcus* spp and Mucorales)
 - ◆ *Candida*, *Saccharomyces*, *Aspergillus*, *Fusarium*, *Acremonium*, ...
 - ◆ *Pneumocystis jirovecii*



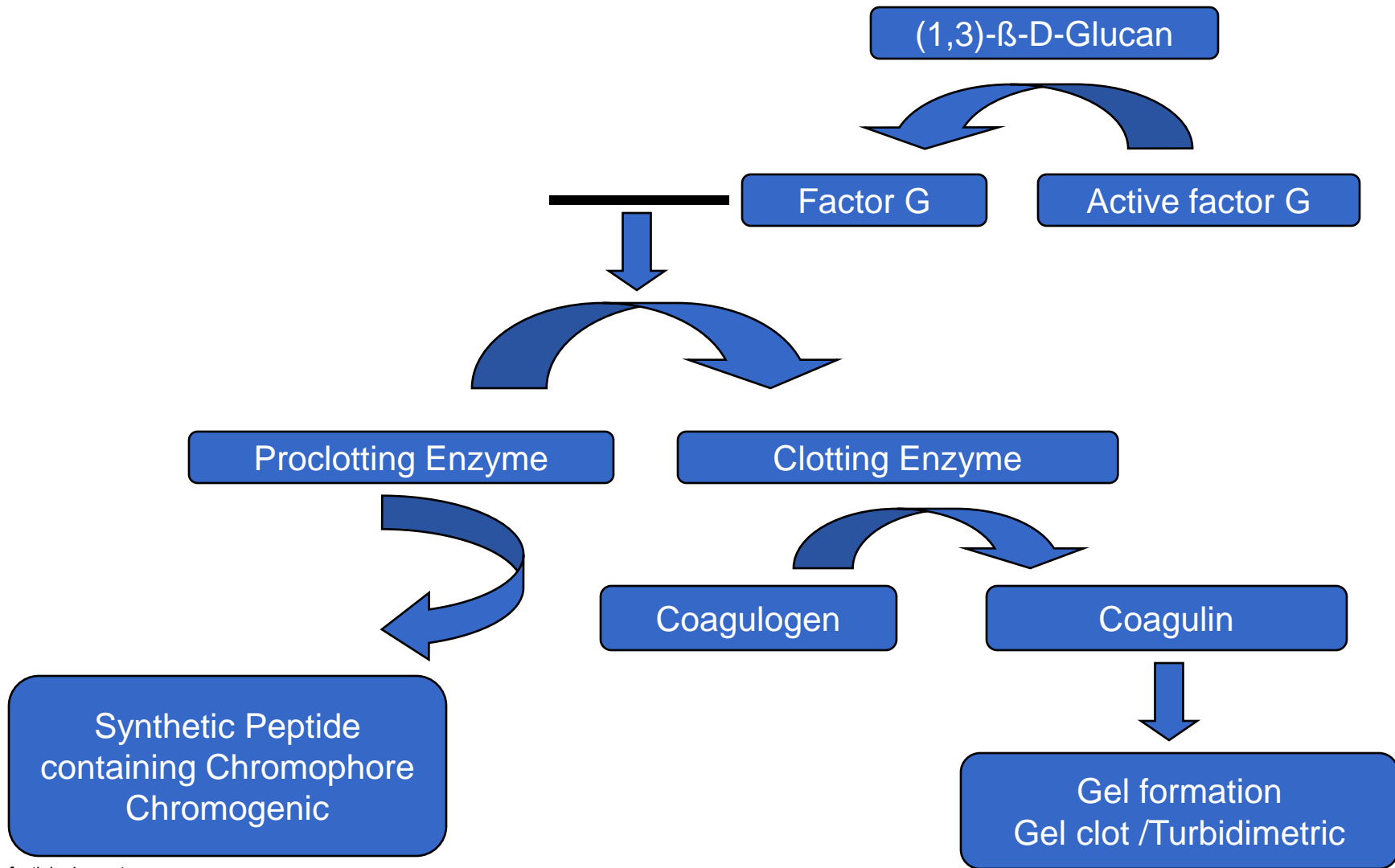
Bennet, NEJM, 2006



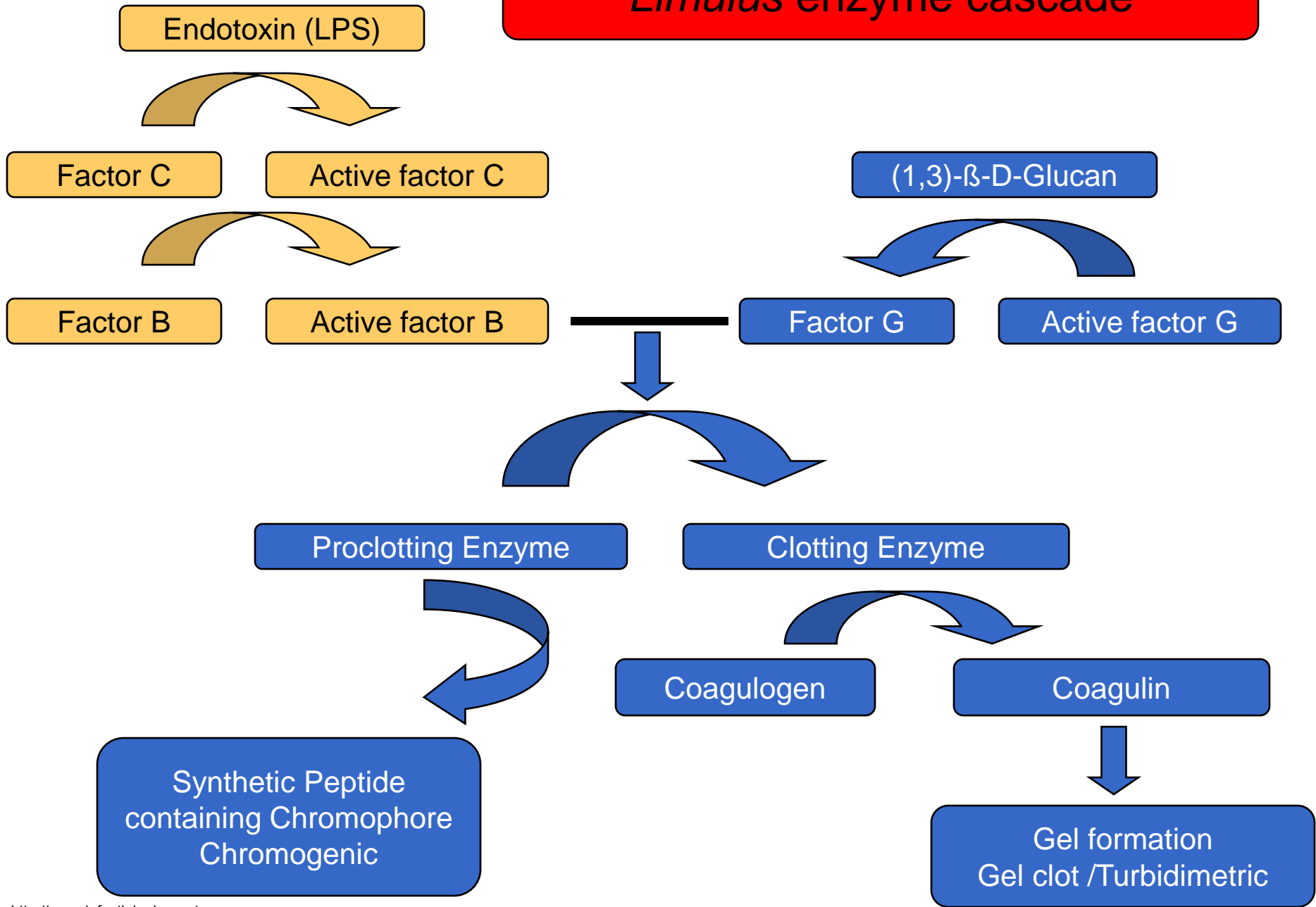
“horse-shoe crab”



Limulus enzyme cascade



Limulus enzyme cascade



4 different commercial kits

Table 3. Comparison of 4 commercial kits for the serum (1→3)- β -D-glucan (β -glucan) assay.

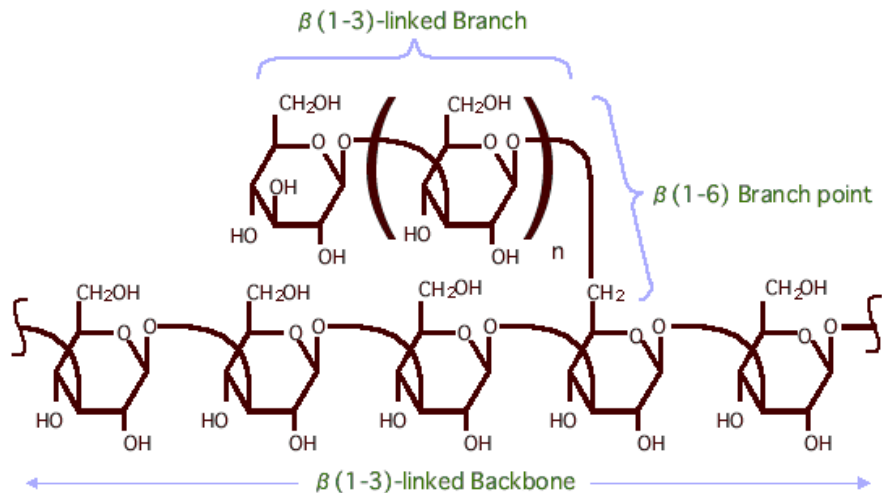
| Variable | Fungitec G-Test MK | β -glucan Test Wako | B-G Star | Fungitell |
|--------------------------|-------------------------------|---------------------------|-------------------------------|---------------------------|
| Manufacturer | Seikagaku Corporation | Wako Pure Chemical | Maruha Corporation | Associates of Cape Cod |
| Country | Japan | Japan | Japan | USA |
| Approval year | 1995 | 1996 | 2001 | 2004 |
| Assay method | Kinetic chromogenic | Kinetic turbidimetry | Endpoint chromogenic | Kinetic chromogenic |
| Sample | Serum or plasma | Serum or plasma | Serum or plasma | Serum |
| Pretreatment | Alkali | Dilution and heating | Dilution and heating | Alkali |
| Standard β -glucan | Pachyman | Carboxymethyl-curdlan | Lentinan | Pachyman |
| Origin of lysate | <i>Tachypleus tridentatus</i> | <i>Limulus polyphemus</i> | <i>Tachypleus tridentatus</i> | <i>Limulus polyphemus</i> |
| Cutoff value, pg/mL | 20 | 11 | 11 | 60 or 80 |
| Measurable range, pg/mL | 3.9–500 | 6–600 | 1.2–120 | 31.25–500 |
| Turn-around time, min | 30 | 90 | 30 | 40 |

False positives

| | |
|----------------------|---|
| Treatments | Immunoglobulines Albumin Coagulation factors Antibiotics (piperacillin – tazobactam) Others? (chemotherapies ...) |
| Patient cares | Hemodialysis with cellulose membranes Gauze or other materials that contain glucans Tubes handling |
| Bacterial infections | Gram negative bacteria Some streptococci |
| Patient linked | Mucosal damages (yeast colonization) Hemolytic or lipemic samples ... |

Pickering JW et al, JCM 2005, 43 : 5957-62

Glucan



- β -glucans occur most commonly as cellulose in plants, the bran of cereal grains, the cell wall of baker's yeast, certain fungi, mushrooms and bacteria
- Some forms of beta glucans are useful in human nutrition as texturing agents and as soluble fiber supplements

Conclusions (EORTC / MSG)

- Included in EORTC/MSG criteria

De Pauw et al CID 2008

Candida versus Aspergillus

Table 3. Sensitivity of (1 → 3)-β-D-Glucan (BDG) Testing to Detect Proven or Probable Systemic *Candida* Infection in Comparison with Invasive Aspergillosis As Reported in Different Studies

| Study | Cutoff(pg/mL) | Systemic <i>Candida</i> infections, proportion (%) | Invasive aspergillosis, proportion (%) |
|-------------------------------------|---------------------------|--|--|
| Hachem et al 2009 [27] | 80 (2 consecutive values) | 13/21 (62) | 14/21 (67) |
| Koo et al 2009 [28] | 80 | 26/41 (63) | 24/32 (75) |
| Obayashi et al 2008 [19] | 30 | 3/3 (100) | 28/28 (100) |
| Persat et al 2008 [31] | 80 | 22/26 (85) | 48/70 (69) |
| Senn et al 2008 [32] | 7 (2 consecutive values) | 10/17 (59) | 9/15 (60) |
| Akamatsu et al 2007 [33] | 40 | 7/14 (50) | 5/5 (100) |
| Ostrosky-Zeichner et al 2005 [36] | 80 | 83/107 (78) | 8/10 (80) |
| Odabashi et al 2004 [20] | 80 | 9/11 (82) | 4/4 (100) |
| Mori et al 1997 [43] | 1000 | 11/12 (92) | 4/4 (100) |
| Mitsutake et al 1996 [44] | 60 | 27/32 (84) | 5/5 (100) |
| Miyazaki et al 1995 [45] | 10 | 11/11 (100) | 3/3 (100) |
| Total from all studies ^a | ... | 222/295 (75) | 152/197 (77) |

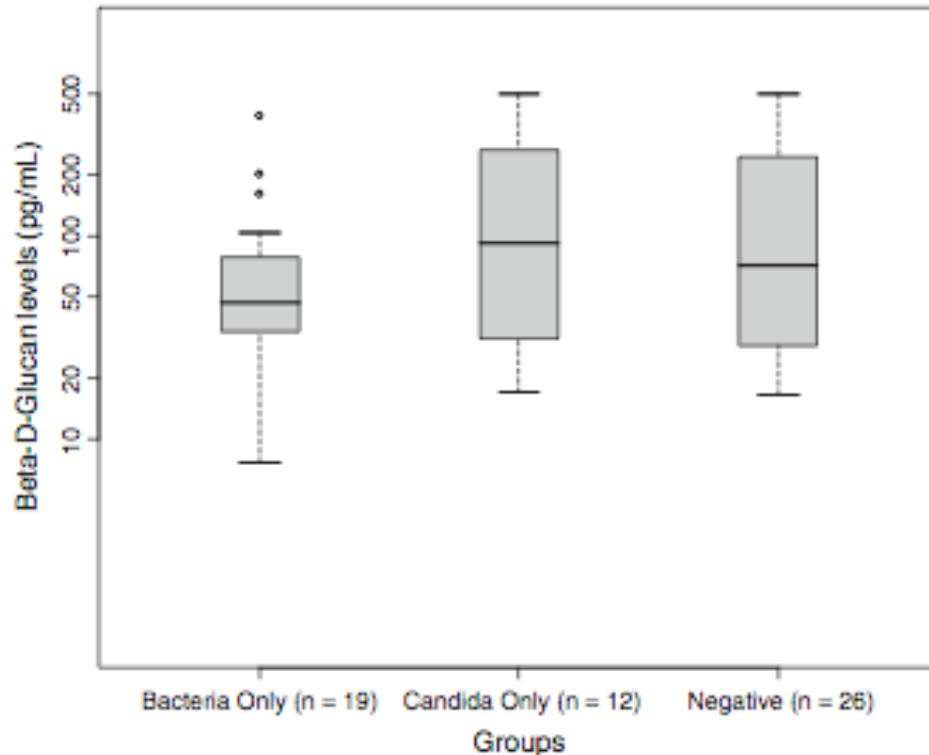
^a Total represents cumulative data

No difference between *Candida* and *Aspergillus* ¹

Performance similar to Galactomannan for invasive aspergillosis ^{2,3}

¹ Karageorgopoulos et al CID 2011; ^{2,3} Leeflang MM et al Cochrane 2008; Pfeiffer CD et al CID 2006.

β -D-glucan in ICU



No discrimination between bacterial sepsis, candida and others

Fig. 2 Results of BG levels by culture group. Box-and-whisker plots of BG levels by culture groups. There were no significant differences across groups ($P = .57$ from the Kruskal-Wallis test).

β -D-glucan metaanalyses

β -D-Glucan Assay for the Diagnosis of Invasive Fungal Infections: A Meta-analysis

Drosos E. Karageorgopoulos,^{1,2} Evidiki K. Vouloumanou,¹ Fotinie Ntziora,^{1,2} Argyris Michalopoulos,^{1,3} Petros I. Rafailidis,^{1,4} and Matthew E. Falagas^{1,4,5}

¹Alfa Institute of Biomedical Sciences; ²Department of Medicine, Laikon General Hospital, and ³Intensive Care Unit and ⁴Department of Medicine, Henry Dunant Hospital, Athens, Greece; and ⁵Department of Medicine, Tufts University School of Medicine, Boston, Massachusetts

β -Glucan Antigenemia Assay for the Diagnosis of Invasive Fungal Infections in Patients With Hematological Malignancies: A Systematic Review and Meta-Analysis of Cohort Studies From the Third European Conference on Infections in Leukemia (ECIL-3)

Frédéric Lamothe,^{1,a} Mario Cruciani,^{2,a} Carlo Mengoli,³ Elio Castagnola,⁴ Olivier Lortholary,^{5,6,7} Malcolm Richardson,⁸ and Oscar Marchetti,¹ on behalf of the Third European Conference on Infections in Leukemia (ECIL-3)

Karageorgopoulos et al CID 2011; Lamothe et al CID 2012

Warning: false+ and false-

- False +
 - ◆ Blood products (immunoglobulines, albumin)
 - ◆ Hemodialysis with cellulose membrane
 - ◆ Antibiotics (amoxicillin-clavulanate, piperacillin-tazobactam)
 - ◆ Bacterial sepsis
 - ◆ Gauze (surgery)
 - ◆ Severe mucitis
- False -
 - ◆ Antifungals (empirical, prophylaxis)
 - ◆ Glucan non-producing fungi (mucorales, *Cryptococcus*)
- **Skilled technicians**
 - ◆ Risk of contamination from the bed-side to the lab
 - ◆ Not easy-to-perform test

Karageorgopoulos et al CID 2011; Marchetti et al ECIL3

β -D-glucan and pneumocystosis

Blood (1 \rightarrow 3)- β -D-Glucan as a Diagnostic Test for HIV-Related *Pneumocystis jirovecii* Pneumonia

Paul E. Sax,¹ Lauren Komarow,² Malcolm A. Finkelman,³ Philip M. Grant,⁴ Janet Andersen,² Eileen Scully,¹ William G. Powderly,⁵ and Andrew R. Zolopa⁴ for the AIDS Clinical Trials Group Study A5164 Team

¹Division of Infectious Diseases, Department of Medicine, Brigham and Women's Hospital and Harvard Medical School, and ²Harvard School of Public Health, Boston, ³Associates of Cape Cod, East Falmouth, Massachusetts; ⁴Division of Infectious Diseases and Department of Medicine, Stanford University School of Medicine, Stanford, California; and ⁵School of Medicine, University College Dublin, Ireland

Sax et al CID 2011

β -D-glucan titers according to microscopy and *P. jirovecii* qPCR BAL results

| | Group 1 Microscopy-positive qPCR-positive BAL n=10 (%) | Group 2 Microscopy-negative qPCR-positive BAL n=26 (%) | Group 3 Microscopy-negative qPCR-negative BAL n=34 (%) | <i>P</i> values |
|--|---|---|---|-----------------|
| Number with β -glucan >500 pg/mL (%) | 10 (100) | 6 (23) | 2 (6) | <0.0001 |
| Number with β -glucan >80 and \leq 500 pg/mL (%) | 0 | 19 (73) | 8 (24) | <0.0001 |
| Number with β -glucan \leq 80 pg/mL (%) | 0 | 1 (4) | 24 (71) | <0.0001 |

β -D-glucan performance

- Thus, β -glucan specificity decreased from 70.6% to 41.7% depending on whether the microscopy-negative and qPCR-positive BALs were considered as true-positives or false-positives, respectively.

A systematic literature review on the diagnosis of invasive aspergillosis using polymerase chain reaction (PCR) from bronchoalveolar lavage clinical samples

Felipe Francisco Tuon

- 15 articles out of 45
- Criteria:
 - ◆ Data for Ss and Sp
 - ◆ >10 BAL
 - ◆ EORTC criteria
 - ◆ Inclusions of control patients
- No condition on PCR assays and DNA extraction
- Mean Sensitivity: 79% (95% CI: 72.8-83.1)
- Mean Specificity: 94% (95% CI: 92.1-95.0)

Rev Iberoam Micol 2007; 24 89-94

A systematic literature review on the diagnosis of invasive aspergillosis using polymerase chain reaction (PCR) from bronchoalveolar lavage clinical samples

Felipe Francisco Tuon

- 15 articles out of 45
- Criteria:
 - ◆ Data for Ss and Sp
 - ◆ >10 BAL
 - ◆ EORTC criteria
 - ◆ Inclusions of control patients

Aspergillus DNA detected in BAL from 4/11 volunteers (36%)

- Mean Sensitivity: 79% (95% CI: 72.8-83.1)
- Mean Specificity: 94% (95% CI: 92.1-95.0)

Denning et al CID 2011

Use of PCR for diagnosis of invasive aspergillosis: systematic review and meta-analysis

Carlo Mengoli, Mario Cruciani, Rosemary A Barnes, Juergen Loeffler, J Peter Donnelly

- 16 publications (>10,000 samples; 1618 patients)
 - EORTC criteria
 - Prospective design
- Sensitivity: 75% \geq 2+ PCR (95% CI: 54-88)
88% \geq 1+ PCR (95% CI: 75-95)
- Specificity: 87% \geq 2+ PCR (95% CI: 78-93)
75% \geq 1+ PCR (95% CI: 63-84)
- Prerequisite on PCR assays
 - ♦ Previously “validated”
 - ♦ No constraint on PCR itself

C. Mengoli, Lancet Inf Dis, 2009

Use of PCR for diagnosis of invasive aspergillosis: systematic review and meta-analysis

Carlo Mengoli, Mario Cruciani, Rosemary A Barnes, Juergen Loeffler, J Peter Donnelly

| | Sample type | Sample volume | Cell wall disruption* | DNA extraction methods* | PCR method† | Target gene | Appropriate controls | | Minimum samples needed for positive PCR | Methods used (refs) |
|---|------------------------|---------------|-------------------------|-------------------------|-----------------------------------|-------------|----------------------|---|---|---------------------|
| | | | | | | | Negative‡ | Positive§ | | |
| Hebart et al (2000) ²⁸ | Whole blood | 5 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 and 2 | 36 |
| Hebart et al (2000) ²⁹ | Whole blood | 5 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 | 37 |
| Williamson et al (2000) ³⁶ | Serum | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 | 37 |
| Buchheidt et al (2001) ²¹ | Whole blood and sputum | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 | 38 |
| Ferns et al (2002) ²⁵ | Whole blood | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 | 39,40 |
| Raad et al (2002) ³¹ | Whole blood | 10 mL | Zymolyase or Lyticase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 and 2 | 39 |
| Buchheidt et al (2004) ²² | Whole blood and sputum | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 and 2 | 38 |
| Kawazu et al (2004) ³¹ | Plasma | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 and 2 | 41 |
| Lass-Floerl et al (2004) ³² | Whole blood | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 | 37 |
| Halliday et al (2006) ²⁷ | Whole blood | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 2 | 42 |
| Jordanides et al (2005) ³⁰ | Whole blood | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 | 37,42 |
| Scotter et al (2005) ³⁴ | Whole blood | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 1 and 2 | 38 |
| El Mahallawi et al (2006) ³⁴ | Serum | 10 mL | Zymolyase | QIAamp | PCR-ELISA | 18S rRNA | Yes | Serial dilutions | 2 | 36 |
| Florent et al (2006) ³⁶ | Serum | .. | QIAamp DNA Mini Kit | QIAamp | PCR-ELISA | mtDNA | Yes | <i>A. fumigatus</i> DNA (10 conidia) or inhibition control (<i>S. pyogenes</i>) | 2 | 39 |
| White et al (2006) ³⁵ | Whole blood | 2 mL | Mechanical, glass beads | MagNA Pure | RT-PCR with TaqMan and nested PCR | 28S rRNA | Yes | Serial dilutions or cloned PCR products in serial dilutions | 1 | 37 |
| Cesaro et al (2008) ²³ | Whole blood | 3 mL | Zymolyase | QIAamp | RT-PCR with FRET | 18S rRNA | .. | .. | 1 and 2 | 42 |

Whole blood vs serum = 12/16
 Volume = 10 ml-200 µl
 Zymolyase or Lyticase = 13/16
 Nested-PCR = 6/16
 DNA target 18S rRNA = 11/16
 ...
 No requirement for PCR test

A. fumigatus=*Aspergillus fumigatus*. BALF=bronchoalveolar lavage fluid. mtDNA=mitochondrial DNA. SDS=sodium dodecyl sulphate buffer. *S. pyogenes*=*Streptococcus pyogenes*. ..=not reported. *QIAamp, QIAGEN; MagNA Pure, Roche. Zymolyase, ICN or Sigma; Lyticase, Sigma; Phenol-chloroform, Sigma; glass beads, Sigma. †TaqMan uses hydrolysis probe for real-time (RT)-PCR; FRET (fluorescent resonance energy transfer) uses hybridisation probe for RT-PCR. ‡Sterile water, sterile buffer, or blood from healthy individuals. §Serially diluted *Aspergillus* DNA, serially diluted conidia, or internal control (eg, plasmid).

Table 2: Technical details of the PCR methods used in the studies included

Metaanalysis

JOURNAL OF CLINICAL MICROBIOLOGY, Feb. 2011, p. 665–670
0095-1137/11/\$12.00 doi:10.1128/JCM.01602-10
Copyright © 2011, American Society for Microbiology. All Rights Reserved.

Vol. 49, No. 2

PCR Diagnosis of Invasive Candidiasis: Systematic Review and Meta-Analysis^{▽†}

Tomer Avni,^{1*} Leonard Leibovici,¹ and Mical Paul²

- pooled sensitivity = 0.95 (95% CI: 0.88-0.98)
- pooled specificity = 0.92 (95% CI: 0.88-0.95)
- No requirement for the PCR itself

Table 1. MIQE checklist for authors, reviewers, and editors.^a

| Item to check | Importance | Item to check | Importance |
|---|------------|---|----------------|
| Experimental design | | qPCR oligonucleotides | |
| Definition of experimental and control groups | E | Primer sequences | E |
| Number within each group | E | RTPriMerDB identification number | D |
| Assay carried out by the core or investigator's laboratory? | D | Probe sequences | D ^d |
| Acknowledgment of authors' contributions | D | Location and identity of any modifications | E |
| Sample | | Manufacturer of oligonucleotides | |
| Description | E | Purification method | D |
| Volume/mass of sample processed | D | qPCR protocol | E |
| Microdissection or macrodissection | E | Complete reaction conditions | E |
| Processing procedure | E | Reaction volume and amount of dNADNA | E |
| If frozen, how and how quickly? | E | Primer, (probe), Mg ²⁺ , and dNTP concentrations | E |
| qPCR validation | | | |
| Evidence of optimization (from gradients) | | D | E |
| Specificity (gel, sequence, melt, or digest) | | E | D |
| For SYBR Green I, C _q of the NTC | | E | D |
| Calibration curves with slope and y intercept | | E | E |
| PCR efficiency calculated from slope | | E | E |
| CI _s for PCR efficiency or SE | | D | E |
| r ² of calibration curve | | E | E |
| Linear dynamic range | | E | E |
| C _q variation at LOD | | E | E |
| CI _s throughout range | | D | E |
| Evidence for LOD | | E | E |
| If multiplex, efficiency and LOD of each assay | | E | D |
| Location of each primer by exon or intron (if applicable) | E | Software (source, version) | E |
| What splice variants are targeted? | E | C _q or raw data submission with RDM | D |

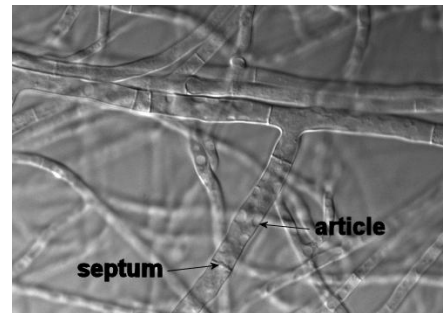
qPCR validation

| | |
|--|---|
| Evidence of optimization (from gradients) | D |
| Specificity (gel, sequence, melt, or digest) | E |
| For SYBR Green I, C _q of the NTC | E |
| Calibration curves with slope and y intercept | E |
| PCR efficiency calculated from slope | E |
| CI _s for PCR efficiency or SE | D |
| r ² of calibration curve | E |
| Linear dynamic range | E |
| C _q variation at LOD | E |
| CI _s throughout range | D |
| Evidence for LOD | E |
| If multiplex, efficiency and LOD of each assay | E |

^a All essential information (E) must be submitted with the manuscript. Desirable information (D) should be submitted if available. If primers are from RTPriMerDB, information on qPCR target, oligonucleotides, protocols, and validation is available from that source.
^b FFPE, formalin-fixed, paraffin-embedded; RIN, RNA integrity number; RQI, RNA quality indicator; GSP, gene-specific priming; dNTP, deoxynucleoside triphosphate.
^c Assessing the absence of DNA with a no-reverse transcription assay is essential when first extracting RNA. Once the sample has been validated as DNA free, inclusion of a no-reverse transcription control is desirable but no longer essential.
^d Disclosure of the probe sequence is highly desirable and strongly encouraged; however, because not all vendors of commercial predesigned assays provide this information, it cannot be an essential requirement. Use of such assays is discouraged.

Why these technical aspects so important?

- Origin of DNA
 - ◆ Conidia, hyphae, cell free DNA?
 - ◆ No rationale for DNA extraction and specimens
 - Fungus itself -> stringent DNA extraction
 - Cell-free DNA -> automated DNA extraction, simple protocols
- Numerous sources of false positives AND false negatives
 - Always very low DNA burdens



EAPCRI*: “Towards an European Standard for *Aspergillus*-PCR”

- P. Donnelly, J. Löffler, L. White
 - ◆ 2006-2007: 24 laboratories (all of them use qPCR)
- PCR amplification methods are very consistent in their performance
 - ◆ 95% of methods detected the predicted 100% threshold
 - ◆ *Aspergillus* gene target, PCR platform does not seem to matter
- Wide variation in the performance of extraction methods
 - ◆ Use of larger volumes of blood correlated with better performance: at least 4 ml EDTA blood should be used
 - ◆ Bead-beating methods performed optimally when testing QC panel
 - ◆ Specimen validation using animal models
 - ◆ Performance in clinical specimens?

*The European *Aspergillus* PCR Initiative

Lewis White et al, JCM 2010 and 2011

Goals for biomarkers

- Two ways of using biomarkers
 - ◆ Diagnostic tools
 - Low sensitivity, meaning for opportunistic diseases
 - ◆ Defining an optimal risk-based strategy minimizing the risks of both invasive fungal diseases and over-treatment

Conclusion

- Direct examination and culture far to be outdated
 - ◆ Identification
 - ◆ MIC
 - ◆ Resistance mechanisms
 - ◆ Prognosis (with GM)
- GM
 - ◆ Define your objectives (Diagnostic or screening)
 - ◆ If screening, prevalence of the disease should be >5%
- Glucan
 - ◆ *Pneumocystis*
- To program regular reappraisals
 - ◆ epidemiological trends, new treatments, new markers ...



Institut Pasteur

<http://www.infectiologie.org.tn>