

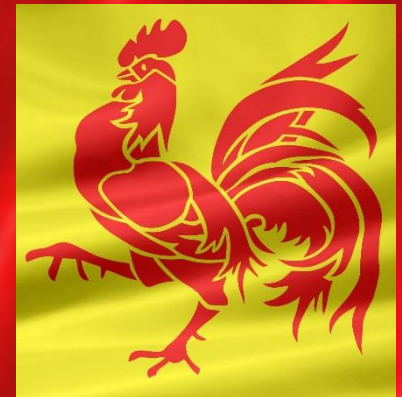
Université  
de Liège



# PCR for Diagnosis in Mycology



Marie-Pierre Hayette  
University Hospital of Liège  
NRC Mycosis  
*SBMHA symposium 2015*



# Table of content

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- Molecular method applied to biological samples
  - PANFUNGAL
  - SPECIFIC
- Molecular methods applied on cultures
- Conclusions

## PCR-based assays on clinical samples

*Which samples?*

Blood plasma  
or serum?

BAL, CSF,..

Tissue  
biopsies

*What kind of extraction method?*

Extraction  
method?

Extraction  
method ?

Extraction  
method ?

*What kind of qPCR method?*

Quantitative/  
qualitative?

Quantitative/  
qualitative?

Quantitative/  
qualitative?

*Which targets?*

Multicopy rRNA genes (18S, 28S, ITS regions)

## Molecular methods applied to cultures

PCR for  
identification

PCR for detection  
of resistance

PCR  
+ gene sequencing

# Pan-fungal PCR

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- Only in patients at high risk of fungal infection
- In sterile fluids or biopsies (with or without positive microscopy)
- Multiplex PCR assays or panfungal with high conserved regions of fungal DNA.
- In house tests or commercial kits

# Panfungal PCR: *in house assays*

**Targets ITS1 and ITS2**

**Gene sequencing**

Retrospective study

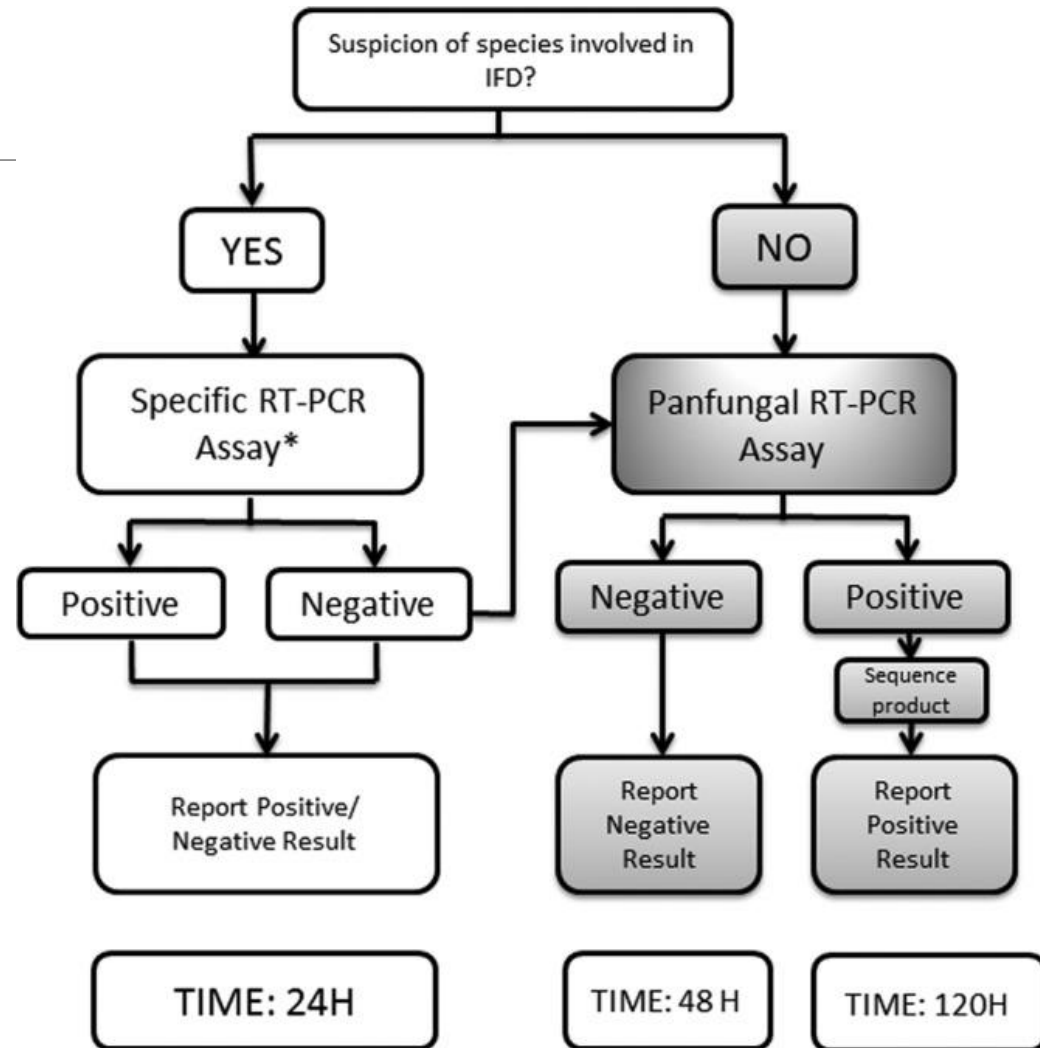
151 biopsies (fresh and paraffin-embedded tissues)

132 patients with proven IFI

**Sensitivity**

**89% (overall)**

**80% in case of negative culture results**



# Panfungal PCR:

## *commercial kits*

Kit provider	Type of targets	Validated samples	Technology
<b>Septi-fast<sup>®</sup> Test MGRADE</b> (Roche, Switzerland)	19 bacteria+6 fungi (Af+5 Candida sp.)	Whole blood	qPCR
<b>Vyoo<sup>®</sup> Test (Analytik, Germany)</b>	34 bacteria + 7 fungi (Af, 6 Candida)	Whole blood	Conventional PCR
<b>RenDx<sup>®</sup> Fungiplex assay</b> (Renishaw, UK)	<b>Aspergillus sp (3 species)+ Candida sp+ C. glabrata and C. krusei</b>	Blood, serum or plasma	PCR+ raman technology
<b>Sepsitest<sup>®</sup> (Molzym, US, CE IVD kit)</b>	Panbacterial + panfungal	Blood+biopsies+ liquids	PCR+gene sequencing (16S or 18S)
<b>Provit<sup>®</sup> (Mobidiag, Finland)</b>	60 bacteria+13 fungi+bacterial resistance gene	Blood cultures	Microarray
<b>Magicplex™ Sepsis Real-time Test, Seegene, Korea</b>	85 bacteria+6 fungi+bacterial resistance genes	Whole blood	Multiplex qPCR

# Clinical evaluation of commercial nucleic acid amplification tests in patients with suspected sepsis

Lars Ljungström<sup>1</sup>, Helena Enroth<sup>2</sup>, Berndt EB Claesson<sup>2</sup>, Ida Ovemyr<sup>3</sup>, Jesper Karlsson<sup>2</sup>, Berit Fröberg<sup>2</sup>, Anna-Karin Brodin<sup>2</sup>, Anna-Karin Pernestig<sup>3</sup>, Gunnar Jacobsson<sup>1</sup>, Rune Andersson<sup>4</sup> and Diana Karlsson<sup>3\*</sup>

Comparison of Magicplex sepsis  
(multiplex PCR) vs Provit kit (microarray on  
blood cultures  
382 episodes of sepsis  
Whole blood/blood cultures

## Results : % of positivity

Blood cultures as reference  
(11%)  
Magicplex (9,7%)  
Provit (8,4%)

Conclusion

Low performance of both tests

# Biomarkers and Molecular Analysis to Improve Bloodstream Infection Diagnostics in an Emergency Care Unit

Anne J. M. Loonen<sup>1,6</sup>, Cornelis P. C. de Jager<sup>2</sup>, Janna Tosserams<sup>2</sup>, Ron Kusters<sup>3</sup>, Mirrian Hilbink<sup>4</sup>, Peter C. Wever<sup>5</sup>, Adriaan J. C. van den Brule<sup>1,6\*</sup>

Comparison of SepsiT<sub>est</sub> (Molz<sub>ym</sub>)  
Vs Magicplex sepsis (multiplex PCR) vs  
Blood cultures

125 samples analysed  
Emergency department

## SEN. SPE. PPV. NPV.

SepsiT<sub>est</sub> 11%, 96%, 43%, 80%,  
Magicplex 37%, 77%, 30%, 82%

Conclusion

Poor performance of both tests



# PCR coupled with ESI-Mass Spectrometry

## Broad Assay Menu and Sample Types

ASSAY	COVERAGE	SAMPLE TYPE
BAC BSI BAC SFT	750+ Bacteria, Candida and 4 Antibiotic Resistance Markers: mecA, vanA, vanB and kpc	5ml EDTA whole blood Sterile fluids and tissues
BAC LRT	Identical coverage as BAC BSI and BAC SFT with additional <b>semi-quantitative threshold</b>	BAL and ETA
Fungal	200+ fungi	BAL and Isolates
Viral IC	130+ viruses in 13 reporting groups	Plasma



CE Marked. For In Vitro Diagnostic Use.

# Principle

## PCR/ESI-MS: Polymerase chain reaction/electrospray ionization-mass spectrometry

### NUCLEIC ACID EXTRACTION



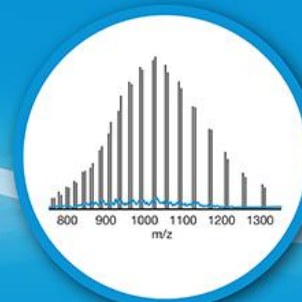
Pathogen nucleic acids extracted from clinical samples

### AMPLIFICATION



Broad-range primers bind to conserved regions in bacteria, viruses or fungi  
Variable pathogen-specific regions flanked by the conserved regions are amplified

### MASS SPECTROMETRY



ESI-MS analysis provides information on the base composition of the amplicons

### PATHOGEN DATABASE ANALYSIS



Algorithms used to compare base composition of amplicons against pathogen database

SAMPLE

'Same-shift' pathogen identification is an achievable goal

RESULT

# Evaluation of the Broad-Range PCR/ESI-MS Technology in Blood Specimens for the Molecular Diagnosis of Bloodstream Infections

Elena Jordana-Lluch<sup>1,2,5</sup>, Montserrat Giménez<sup>1,2</sup>, M<sup>a</sup> Dolores Quesada<sup>1</sup>, Belén Rivaya<sup>1</sup>, Clara Marcó<sup>1</sup>, M<sup>a</sup> Jesús Domínguez<sup>3</sup>, Fernando Arméstar<sup>4</sup>, Elisa Martró<sup>1,5,6\*</sup>, Vicente Ausina<sup>1,2,5</sup>

410 Whole blood specimen  
Emergency unit and ICU  
patients  
“suspicion of sepsis”

Table 1. Agreement between methods according to the two gold standards used by microorganisms isolated by conventional microbiological methods and detected by IRIDICA.

	Global		Emergency Room		Intensive Care Unit	
	BC gold standard	Clinical infection criterion	BC gold standard	Clinical infection criterion	BC gold standard	Clinical infection criterion
Matched positives ( <i>n</i> )	176	217 <sup>a</sup>	147	152 <sup>a</sup>	29	65 <sup>a</sup>
Matched negatives ( <i>n</i> )	143	143	n.a.	n.a.	143	143
IRIDICA overcalls ( <i>n</i> )	80	39	21	16	59	23
IRIDICA misses ( <i>n</i> )	64	64	56	56	8	8
Overall agreement (%)	68.9	77.8	n.a.	n.a.	72.0	87.0
Positive agreement (%)	73.3	77.2	72.4	73.1	78.4 <sup>b</sup>	89.0 <sup>b</sup>
Negative agreement (%)	64.1	78.6	n.a.	n.a.	70.8 <sup>b</sup>	86.1 <sup>b</sup>

## CONCLUSION

Sensitivity, specificity, positive and negative predictive values compared with blood culture were 83.3%, 78.6%, 33.9% and 97.3% respectively, and 90.5%, 87.2%, 64.4% and 97.3% respectively, in comparison with the clinical infection criterion.

# More recent concept: the syndromic approach

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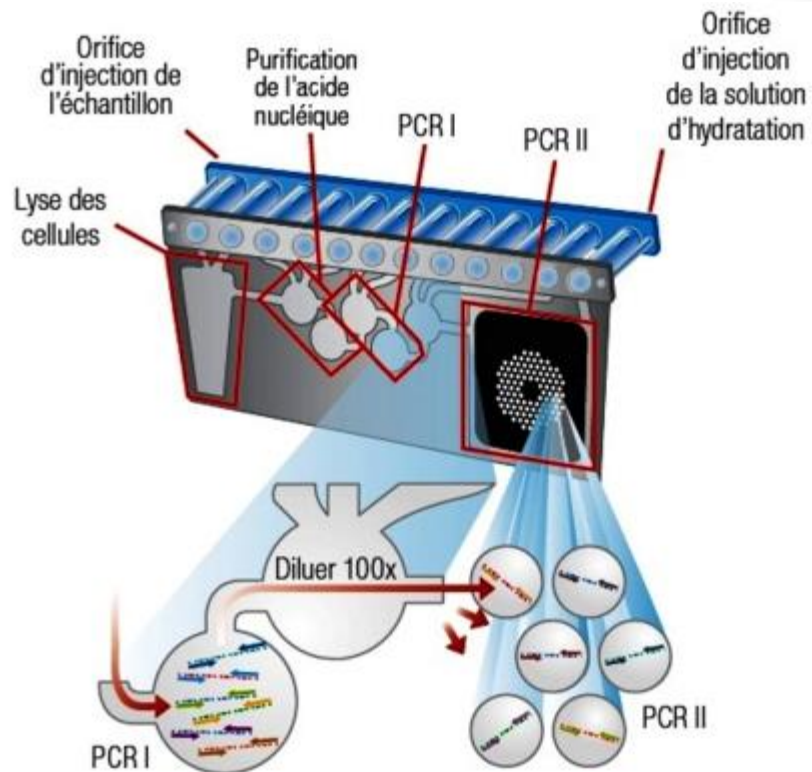
- Application of fungal diagnosis: meningitis
  - Very few fungal targets: only in CSF with *Cryptococcus* for Film array (BioFire, bioMérieux)
  - Genmark (not yet)

# ePlex, GENMARK

*award winning*

## Technology

Click [here](#) to learn more about GenMark's proprietary eSensor® technology



## BIOFIRE, bioMérieux



# Molecular methods=targeted PCR-based assays

	Pneumocystis	Candida	Aspergillus	Mucorales
Used in routine?	Yes	Rarely	Rarely	In development
Which technology?	Real-time PCR (quantitative or qualitative)			
Commercial kits?	Yes mostly in house	Yes	Yes	In development
Kit providers	FTD <i>P. jirovecii</i> (fast-track Diagnostics)	Sacace Renishaw	Myconostics Fast-Track PathoNostics Renishaw	In house only



# Extraction methods

## MolYsis™ Isolation (Bacteria & Fungi)

### Benefits:

- Removal of human DNA
- Removal of PCR inhibitors
- Sensitivity increase up to factor 40,000
- Broad-range lysis of Gram-positive, Gram-negative bacteria and fungi
- DNA-free reagents
- Low hands-on-time



**MolYsis™** is a pre-analytical tool serving the molecular detection of bacteraemia, fungemia and other infections. It constitutes a unique system for the targeted isolation of microbial DNA from clinical samples. By the selective lysis of human cells and the degradation of non-target human DNA, PCR analysis of pathogens is extremely enhanced through increased sensitivity and specificity.

Complete automation



## Aspergillus PCR: One Step Closer to Standardization<sup>▽†</sup>

P. Lewis White,<sup>1\*</sup> Stéphane Bretagne,<sup>2</sup> Lena Klingspor,<sup>3</sup> Willem J. G. Melchers,<sup>4</sup> Elaine McCulloch,<sup>5</sup> Bettina Schulz,<sup>6</sup> Niklas Finnstrom,<sup>7</sup> Carlo Mengoli,<sup>8</sup> Rosemary A. Barnes,<sup>9</sup> J. Peter Donnelly,<sup>4</sup> and Juergen Loeffler<sup>10</sup> on behalf of the European *Aspergillus* PCR Initiative

NPHS Microbiology, Cardiff, United Kingdom<sup>1</sup>; Henri Mondor Hospital, Créteil, France<sup>2</sup>; Karolinska University Hospital, Stockholm, Sweden<sup>3</sup>; Radboud University Nijmegen Medical Centre, Nijmegen, Netherlands<sup>4</sup>; Royal Hospital for Sick Children, Glasgow, United Kingdom<sup>5</sup>; Charité Hospital, Berlin, Germany<sup>6</sup>; Cepheid AB, Toulouse, France<sup>7</sup>; University of Padua, Padua, Italy<sup>8</sup>; Cardiff University, UHW, Cardiff, United Kingdom<sup>9</sup>; and Wuerzburg University, Wuerzburg, Germany<sup>10</sup>

Received 8 September 2009/Returned for modification 6 November 2009/Accepted 4 February 2010

## Multicenter Comparison of Serum and Whole-Blood Specimens for Detection of *Aspergillus* DNA in High-Risk Hematological Patients

Jan Springer,<sup>a</sup> C. O. Morton,<sup>b\*</sup> Michael Perry,<sup>c</sup> Werner J. Heinz,<sup>a</sup> Melinda Paholcsek,<sup>\*\*</sup> Mona Alzheimer,<sup>a</sup> T. R. Rogers,<sup>b</sup> Rosemary A. Barnes,<sup>d</sup> Hermann Einsele,<sup>a</sup> Juergen Loeffler,<sup>a</sup> P. Lewis White<sup>c</sup>

## Clinical Performance of *Aspergillus* PCR for Testing Serum and Plasma: a Study by the European *Aspergillus* PCR Initiative

P. Lewis White,<sup>a</sup> Rosemary A. Barnes,<sup>b</sup> Jan Springer,<sup>c</sup> Lena Klingspor,<sup>d</sup> Manuel Cuenca-Estrella,<sup>e</sup> C. Oliver Morton,<sup>f</sup> Katrien Lagrou,<sup>g</sup> Stéphane Bretagne,<sup>h</sup> Willem J. G. Melchers,<sup>i</sup> Carlo Mengoli,<sup>j</sup> J. Peter Donnelly,<sup>i</sup> Werner J. Heinz,<sup>c</sup> Juergen Loeffler,<sup>c</sup> the EAPCRI

Public Health Wales Microbiology Cardiff, Cardiff, United Kingdom<sup>a</sup>; Cardiff University, UHW, Cardiff, United Kingdom<sup>b</sup>; University of Würzburg Medical Center, Würzburg, Germany<sup>c</sup>; Karolinska University Hospital, Stockholm, Sweden<sup>d</sup>; Spanish National Centre for Microbiology, Instituto de Salud Carlos III, Madrid, Spain<sup>e</sup>; University of Western Sydney, Sydney, NSW, Australia<sup>f</sup>; National Reference Centre for Mycosis, University Hospitals Leuven, Leuven, Belgium<sup>g</sup>; Department of Microbiology and Immunology, Paris Diderot University, Saint Louis Hospital-APHP, Paris, France<sup>h</sup>; Radboud University Nijmegen Medical Centre, Nijmegen, The Netherlands<sup>i</sup>; University of Padua, Padua, Italy<sup>j</sup>



# Clinical Performance of *Aspergillus* PCR for Testing Serum and Plasma: a Study by the European *Aspergillus* PCR Initiative

P. Lewis White,<sup>a</sup> Rosemary A. Barnes,<sup>b</sup> Jan Springer,<sup>c</sup> Lena Klingspor,<sup>d</sup> Manuel Cuenca-Estrella,<sup>e</sup> C. Oliver Morton,<sup>f</sup> Katrien Lagrou,<sup>g</sup> Stéphane Bretagne,<sup>h</sup> Willem J. G. Melchers,<sup>i</sup> Carlo Mengoli,<sup>j</sup> J. Peter Donnelly,<sup>i</sup> Werner J. Heinz,<sup>c</sup> Juergen Loeffler,<sup>c</sup> the EAPCRI

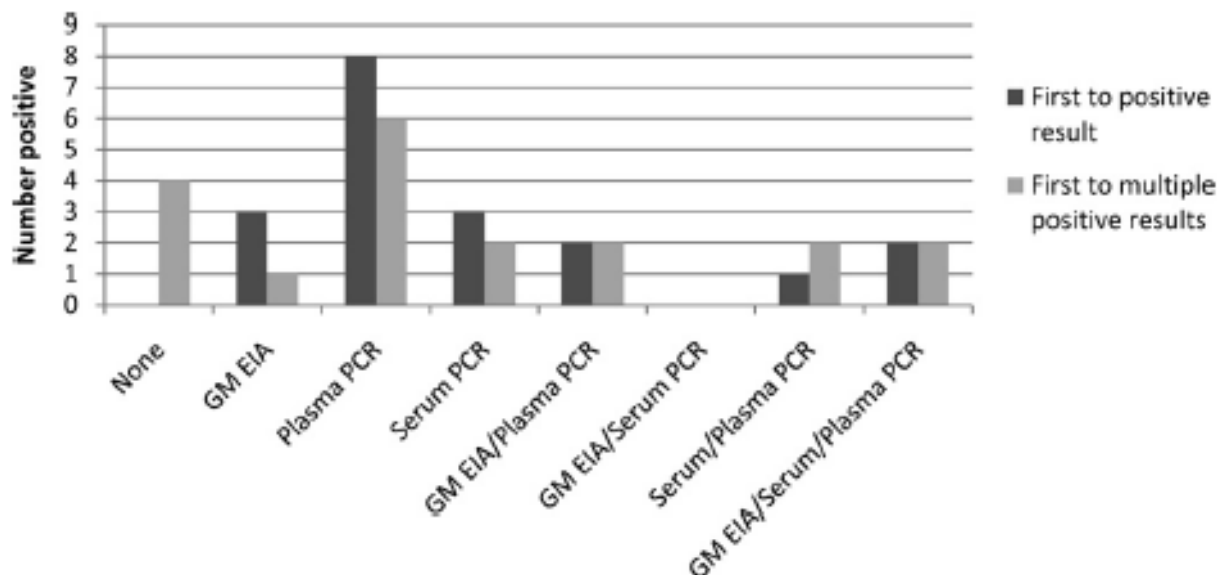
Comparative evaluation of *Aspergillus* PCR on  
serum and plasma  
In population of haematological patients  
Proven/probable IA  
In two centers with standardized qPCR  
Automated DNA extraction

Free DNA is more abundant in plasma than in serum

Poor agreement between plasma PCR and GM in serum 85%

Clinical performance PCR plasma: 95% vs 68% serum

Clinical threshold  $\geq 2$  PCR positive tests



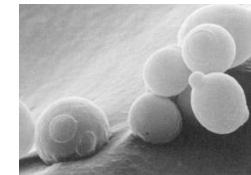
## Validation of a New *Aspergillus* Real-Time PCR Assay for Direct Detection of *Aspergillus* and Azole Resistance of *Aspergillus fumigatus* on Bronchoalveolar Lavage Fluid

Ga-Lai M. Chong,<sup>a</sup> Wendy W. J. van de Sande,<sup>b</sup> Gijs J. H. Dingemans,<sup>c</sup> Giel R. Gaajetaan,<sup>c</sup> Alleke G. Vonk,<sup>b</sup> Marie-Pierre Hayette,<sup>d</sup> Dennis W. E. van Tegelen,<sup>c</sup> Guus F. M. Simons,<sup>c</sup> Bart J. A. Rijnders<sup>a</sup>

37 BAL from hematology patients +40 ICU  
Population classified as IA, probable, unclassifiable  
GM LBA  $\geq 1$   
PCR AsperGenius®  
Identification+ detection of azole resistance

(TR<sub>34</sub>/L98H, TR<sub>46</sub>/T289A, Y121F mutations on CYP51A gene)

PCR <i>Aspergillus</i>	Hematology	ICU
Sensitivity	88.9%	80%
Specificity	89.3%	93.3%
PPV	72.7%	80%
NPV	96.2%	93.3%

**TABLE 2.** Summary of recommendations by *Candida* disease, specimen and test evaluated

Disease	Specimen	Test	Recommendation	Level of evidence
Candidaemia	Blood Serum	Blood culture	Essential investigation <sup>a</sup>	NA
		Mannan/anti-mannan	Recommended	II
		B-D-glucan	Recommended	II
		Other antibodies	No recommendation	No data
		Septifast PCR kit	No recommendation	No data
		In-house PCR	No recommendation	No data
Invasive candidiasis	Blood Serum	Blood culture	Essential investigation	NA
		Mannan/anti-mannan	No recommendation	No data
		B-D-glucan	Recommended	II
		Septifast PCR kit	No recommendation	No data
		In-house PCR	No recommendation	No data
	Tissue and sterile body fluids	Direct microscopy and histopathology	Essential investigation	NA
		Culture	Essential investigation	NA
Chronic disseminated candidiasis	Blood Serum	Immuno-histochemistry	No recommendation	No data
		Tissue PCR	No recommendation	No data
		<i>In situ</i> hybridization	No recommendation	No data
		Blood culture	Essential investigation	NA
		Mannan/anti-mannan	Recommended	II
		B-D-glucan	Recommended	II
	Tissue and sterile body fluids	Septifast PCR kit	No recommendation	No data
		In-house PCR	No recommendation	No data
		Direct microscopy and histopathology	Essential investigation	NA
		Culture	Essential investigation	NA
		Immuno-histochemistry	No recommendation	No data
		Tissue PCR	No recommendation	No data
Oropharyngeal and oesophageal candidiasis	Swab	<i>In situ</i> hybridization	No recommendation	No data
		Culture	Essential investigation	NA
		In-house PCR	No recommendation	No data
	Biopsy <sup>b</sup>	Direct microscopy and histopathology	Essential investigation	NA
		Culture	Essential investigation	NA
		In-house PCR	No recommendation	No data
Vaginal candidiasis	Swab/vaginal secretions	Direct microscopy	Essential investigation	NA
		Culture	Essential investigation	NA
		Commercial tests	Use validated test only	NA
		In-house PCR	No recommendation	No data

# PCR Diagnosis of Invasive Candidiasis: Systematic Review and Meta-Analysis<sup>▽†</sup>

Tomer Avni,<sup>1\*</sup> Leonard Leibovici,<sup>1</sup> and Mical Paul<sup>2</sup>

*Medicine E<sup>1</sup> and Unit of Infectious Diseases,<sup>2</sup> Rabin Medical Center, Beilinson Hospital and Sackler Faculty of Medicine, Tel-Aviv University, Tel Aviv, Israel*

54 studies included

4694 patients → 963 proven/probable or possible invasive candidiasis

Perfect sensitivity+ specificity calculated when ***candidemia+ healthy controls***

***Positivity rates = 85%***

***Vsblood cultures = 38%***

Conclusion of the meta-analysis

- Best results if WB samples, rRNA or P450 genes and if limit of detection is >10 cfu/ml

# Performance of *Candida* Real-time Polymerase Chain Reaction, $\beta$ -D-Glucan Assay, and Blood Cultures in the Diagnosis of Invasive Candidiasis

**M. Hong Nguyen,<sup>1</sup> Mark C. Wissel,<sup>2</sup> Ryan K. Shields,<sup>1</sup> Martin A. Salomoni,<sup>2</sup> Binghua Hao,<sup>1</sup> Ellen G. Press,<sup>1</sup> Ryan M. Shields,<sup>2</sup> Shaoji Cheng,<sup>1</sup> Dimitra Mitsani,<sup>1</sup> Aniket Vadnerkar,<sup>1</sup> Fernanda P. Silveira,<sup>1</sup> Steven B. Kleiboeker,<sup>2</sup> and Cornelius J. Clancy<sup>1,3</sup>**

<sup>1</sup>Department of Medicine, University of Pittsburgh, Pennsylvania; <sup>2</sup>Viracor-IBT Laboratories, Lee's Summit, Missouri; and <sup>3</sup>Department of Medicine, VA Pittsburgh Healthcare System, Pennsylvania

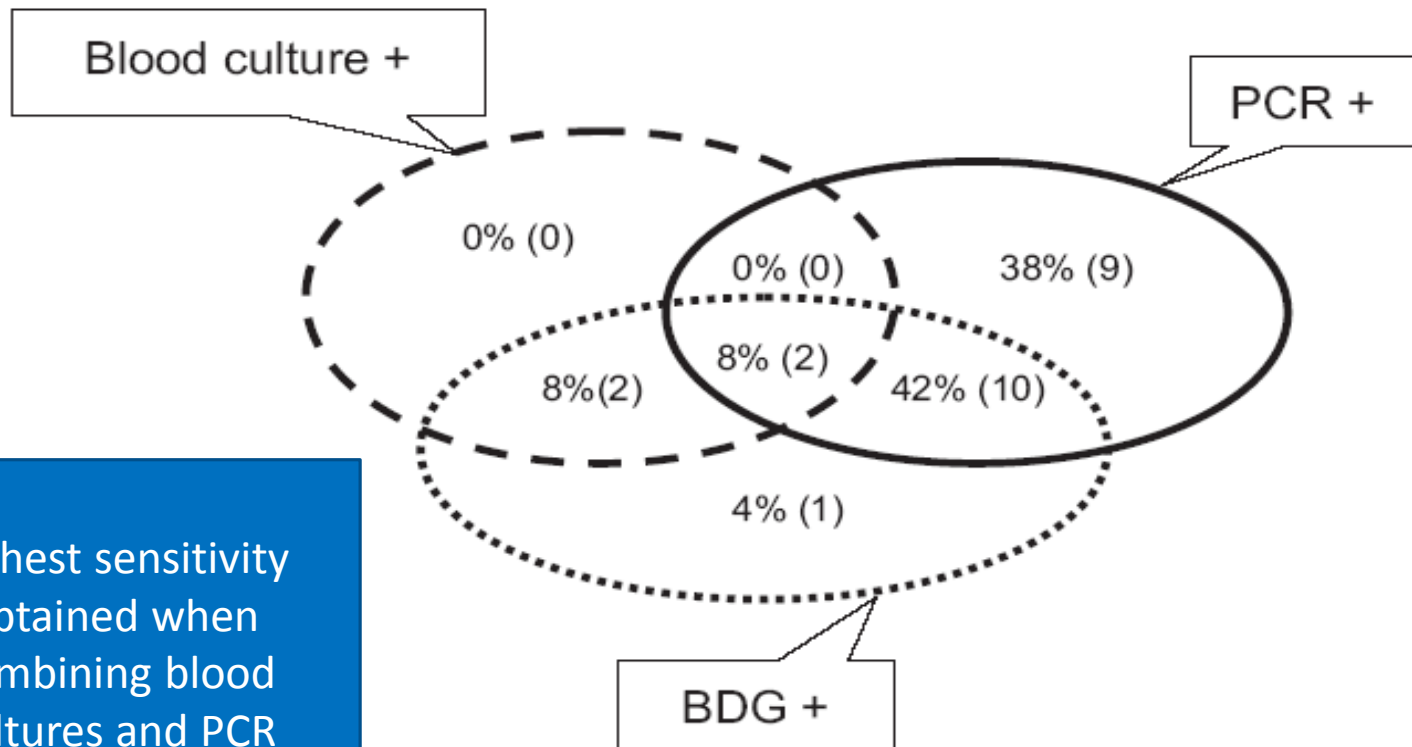
Fungitell® serum or plasma  
Population with invasive candidiasis and controls  
RT-PCR ITS1 & ITS2 on WB, plasma et/or serum  
C. albicans+C. tropicalis,  
C. glabrata+C.krusei, C. parapsilosis complex

Prospective study  
April 2009-2011  
Pittsburg, USA

# Performance of *Candida* Real-time Polymerase Chain Reaction, $\beta$ -D-Glucan Assay, and Blood Cultures in the Diagnosis of Invasive Candidiasis

Assay	Invasive Candidiasis (n = 55)	Candidemia <sup>a</sup> (n = 22)	Deep-Seated Candidiasis <sup>a,b</sup> (n = 38)	Intra-abdominal Candidiasis (n = 34)
PCR <sup>c</sup>				
Sensitivity	80% (44/55)	59% (13/22)	89% (34/38)	88% (30/34)
Specificity	70% (51/73)			
BDG (positive $\geq 80$ pmol/mL)				
Sensitivity	56% (31/55)	68% (15/22)	53% (20/38)	56% (19/34)
Specificity	73% (53/73)			
BDG (positive $\geq 60$ pmol/mL)				
Sensitivity	69% (38/55)	81% (18/22)	66% (25/38)	65% (22/34)
Specificity	63% (46/73)			
P value <sup>d</sup>				
PCR vs BDG (positive $\geq 80$ pmol/mL)	.03	.77	.004	.0015
PCR vs BDG (positive $\geq 60$ pmol/mL)	.31	.23	.04	.06

# PCR performance



Highest sensitivity  
obtained when  
combining blood  
cultures and PCR

# Evaluation of a Commercially Developed Semiautomated PCR–Surface-Enhanced Raman Scattering Assay for Diagnosis of Invasive Fungal Disease

P. Lewis White,<sup>a</sup> Samantha J. Hibblitts,<sup>b</sup> Michael D. Perry,<sup>a</sup> Julie Green,<sup>c</sup> Emma Stirling,<sup>c</sup> Luke Woodford,<sup>c</sup> Graeme McNay,<sup>c</sup> Ross Stevenson,<sup>c</sup> Rosemary A. Barnes<sup>b</sup>

Public Health Wales, Microbiology Cardiff, UHW, Cardiff, United Kingdom<sup>a</sup>; School of Medicine, Cardiff University, UHW, Cardiff, United Kingdom<sup>b</sup>; Rentshaw Diagnostics Ltd., Glasgow, United Kingdom<sup>c</sup>

In vitro testing of spiked blood samples and clinical assessment on stored blood samples.

Sensitivity: 1 -10cfu/ml

Only 28 samples

Sensitivity 80%

- better in case of candidemia)
- Sensitivity vary with sample type =serum<whole blood<plasma

Specificity 87,5%

Conclusion

Nee of further evalutation in a prospective study



# Real-time PCR *Pneumocystis jirovecii*

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## Meta-analysis (*Summah H.. Chin Med J 2013*)

- 10 studies were included 1990-2010
- Sensitivity 98%
- Specificity 94%
- Better performance on BAL than other samples

## RT-PCR semi-quantitative: definition of cut-off<sup>®</sup> (*Fillaux J. et Berry A. Methods Mol Biol. 2013;943:159-70*)

- **Ct ≥ 28** : *probable colonisation*
- **22 ≤ Ct ≤ 28** : possible PcP
- **Ct < 22**: PcP

# Proposed algorithm for PCP diagnosis

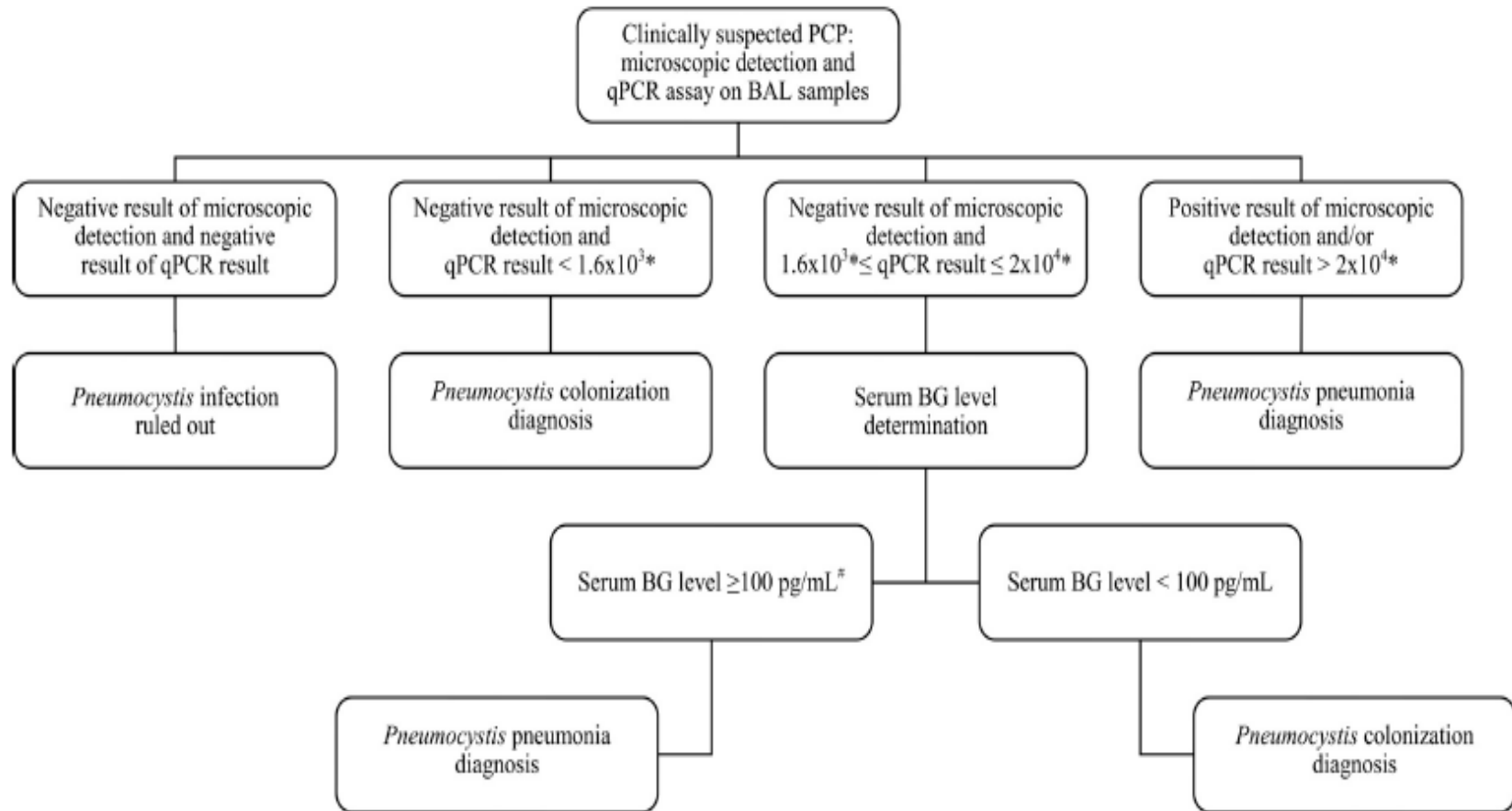
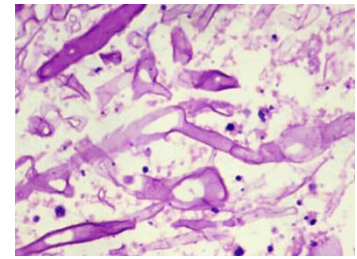


FIG 5 Flow diagram for the diagnosis of *Pneumocystis* infections in patients with clinically suspected *Pneumocystis* pneumonia. Microscopic examination using methanol-Giemsa staining and an immunofluorescence assay (Bio-Rad, Marnes la Coquette, France) and a qPCR assay targeting the mitochondrial large subunit rRNA gene of *P. jirovecii* were used. BAL, bronchoalveolar lavage fluid; BG, (1→3)-β-D-glucan; PCP, *Pneumocystis* pneumonia; \*, results expressed in DNA copies/μl; #, without factors that interfere with (1→3)-β-D-glucan level determinations, particularly concurrent invasive fungal infections.

# Quantitative Polymerase Chain Reaction Detection of Circulating DNA in Serum for Early Diagnosis of Mucormycosis in Immunocompromised Patients



**Laurence Millon,<sup>1,2</sup> Fabrice Larosa,<sup>3</sup> Quentin Lepiller,<sup>2,4</sup> Faezeh Legrand,<sup>3</sup> Steffi Rocchi,<sup>1</sup> Etienne Daguindau,<sup>3</sup>  
Emeline Scherer,<sup>1,2</sup> Anne-Pauline Bellanger,<sup>1,2</sup> Joel Leroy,<sup>5</sup> and Frederic Grenouillet<sup>1,2</sup>**

<sup>1</sup>CNRS-Université de Franche-Comté, UMR 6249 Chrono-environnement, and Departments of <sup>2</sup>Parasitology-Mycology, <sup>3</sup>Clinical Hematology, <sup>4</sup>Virology,  
and <sup>5</sup>Infectious Diseases, University Hospital, Besançon, France

10 Patients with confirmed mucormycosis

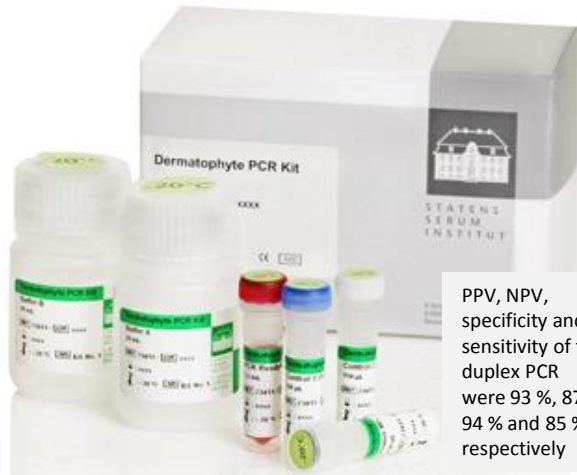
Combinaison of 3 quantitative RT-PCR, 18S RNA genes

Targets: Mucor/Rhizopus, Lichtheimia, Rhizomucor

9/10 PCR positive serum

3-68 days before diagnosis confirmation

# Superficial mycosis: commercial PCR kits for detection/identification of dermatophytes



PPV, NPV, specificity and sensitivity of the duplex PCR were 93 %, 87 %, 94 % and 85 %, respectively

Dermatophytes + *T. rubrum*

**IDEXX**  
**BioResearch**

95% sensitivity  
99 specificity

Microsporium spp.  
*M. canis*  
*M. gypseum*  
*M. ferrugineum*  
*M. audouinii*  
Trichophyton spp.  
*T. mentagrophytes*  
*T. tonsurans* • *T. rubrum*  
*T. megninii*  
*T. violaceum*  
*T. schoenleinii*

*T. mentagrophytes* complex,  
*T. tonsurans*,  
*T. violaceum*,  
*T. rubrum*  
*M. canis*  
*M. audouinii*,  
*M. ferrugineum*

**fast-track** >>  
**DIAGNOSTICS**

FTD Infectious Disease Detection Kit  
- Multiplex Real-time PCR testing -

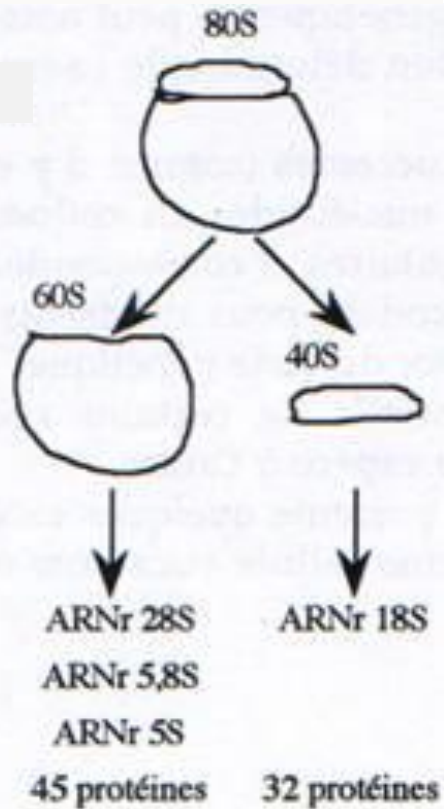


*T. rubrum*,  
*T. mentagrophytes*  
*T. interdigitale*  
*M. canis*  
*M. audouinii*  
*T. violaceum*  
*T. soudanense*  
*Candida albicans*  
*E. floccosum*

Specificity and sensitivity of the PCR are 85 %, 87 %, 94 % and 100 %, respectively

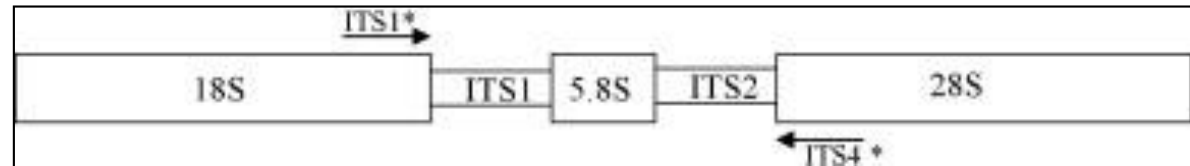
# Molecular sequencing

## EUCARYOTES



## ITS/ARNr

- Internal Transcribed Spacer
- About 1500 pb
- rARN multicopy gene



Chronic granulomatosis  
Multiples hepatic lesions  
-hepatic puncture: mucorales  
-positive blood culutre: mucorales  
ITS sequencing: *Mortierella wolfii*



# ITS primers: choose the right one

Primer name	Sequence (5'-3')	Reference
ITS1 (f)	TCCGTAGGTGAACCTGCGG	White <i>et al.</i> 1990
ITS1F (f)	CTTGGTCATTTAGAGGAAGTAA	Gardes et Bruns 1993
ITS2 (r)	GCTGCGTTCTTCATCGATGC	White <i>et al.</i> 1990
ITS3 (f)	GCATCGATGAAGAACGCAGC	White <i>et al.</i> 1990
ITS4 (r)	TCCTCCGCTTATTGATATGC	White <i>et al.</i> 1990
ITS86F (f)	GTGAATCATCGAATCTTTGAA	Turenne <i>et al.</i> 1999
SR6R (f)	AAGTAAAAGTCGTAACAAGG	Gräser <i>et al.</i> 1999
LR1 (r)	GGTTGGTTTCTTTTCCT	Gräser <i>et al.</i> 1999

# Conclusion

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Need to define the place of fungal PCR and the methods to use for PCR in invasive fungal diagnosis: *Aspergillus* / *Candida* / *mucorales* → screening blood?

Need to have available PCR assays easy to use, not expensive and fully automated

Get the best DNA extraction method and you have >50% to succeed.





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