



ESGMYC

ESCMID STUDY GROUP  
FOR MYCOBACTERIAL  
INFECTIONS

European Society of Clinical Microbiology and Infectious Diseases

Groupe Hospitalier Universitaire  
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université  
PARIS  
DIDEROT

# A year in Clinical Microbiology

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**COI disclosure:** consultancy for bioMérieux, Becton-Dickinson, Hain lifesciences

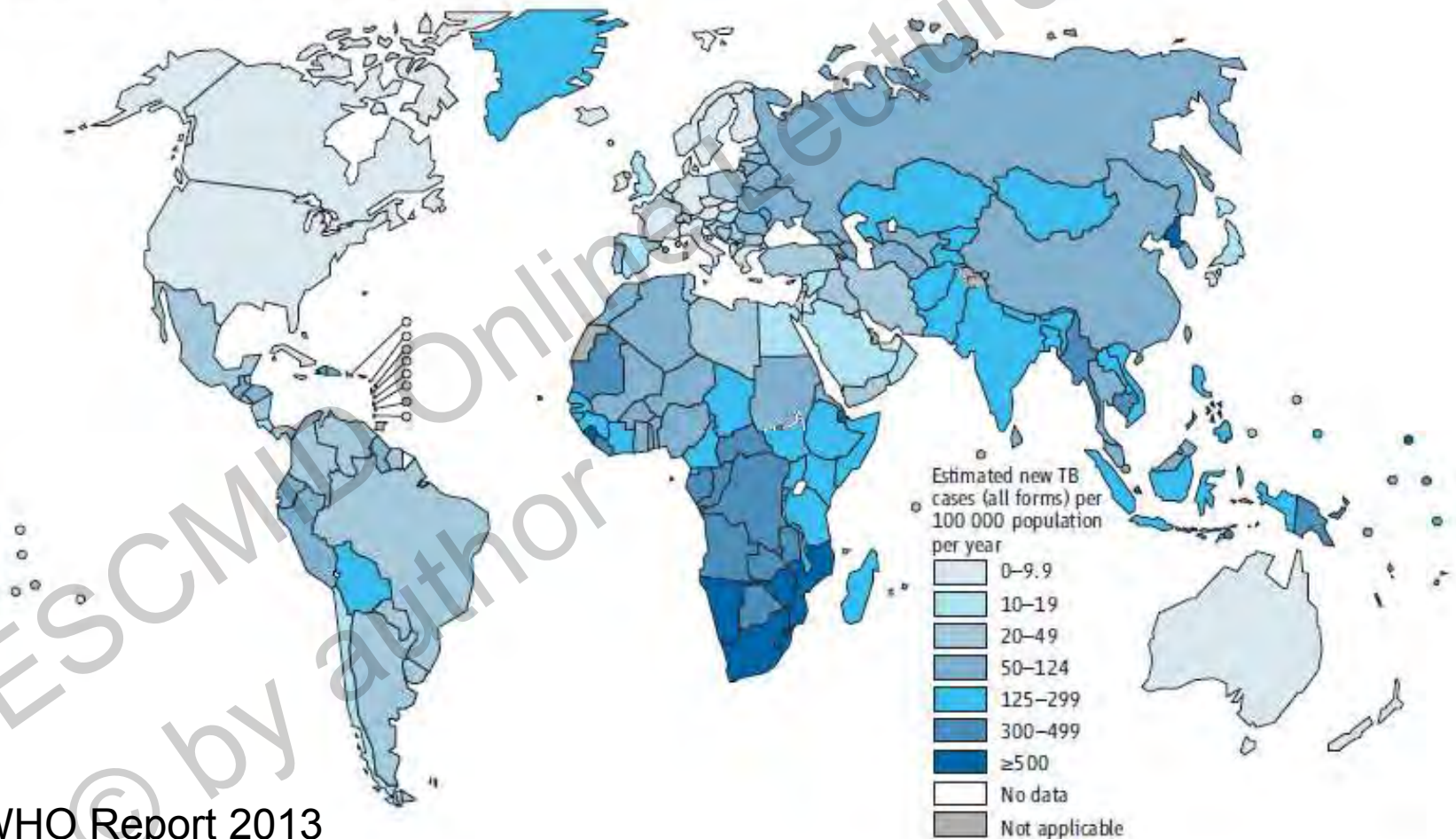
# New anti-tuberculous drugs approved by EMA

- Bedaquiline = TMC-207= R201970  
(Janssen, Johnson and Johnson)
- Delamanid = OPC-67683  
(Otsuka Pharmaceutical, Tokushima)

# 8.6 millions new cases TB in 2012

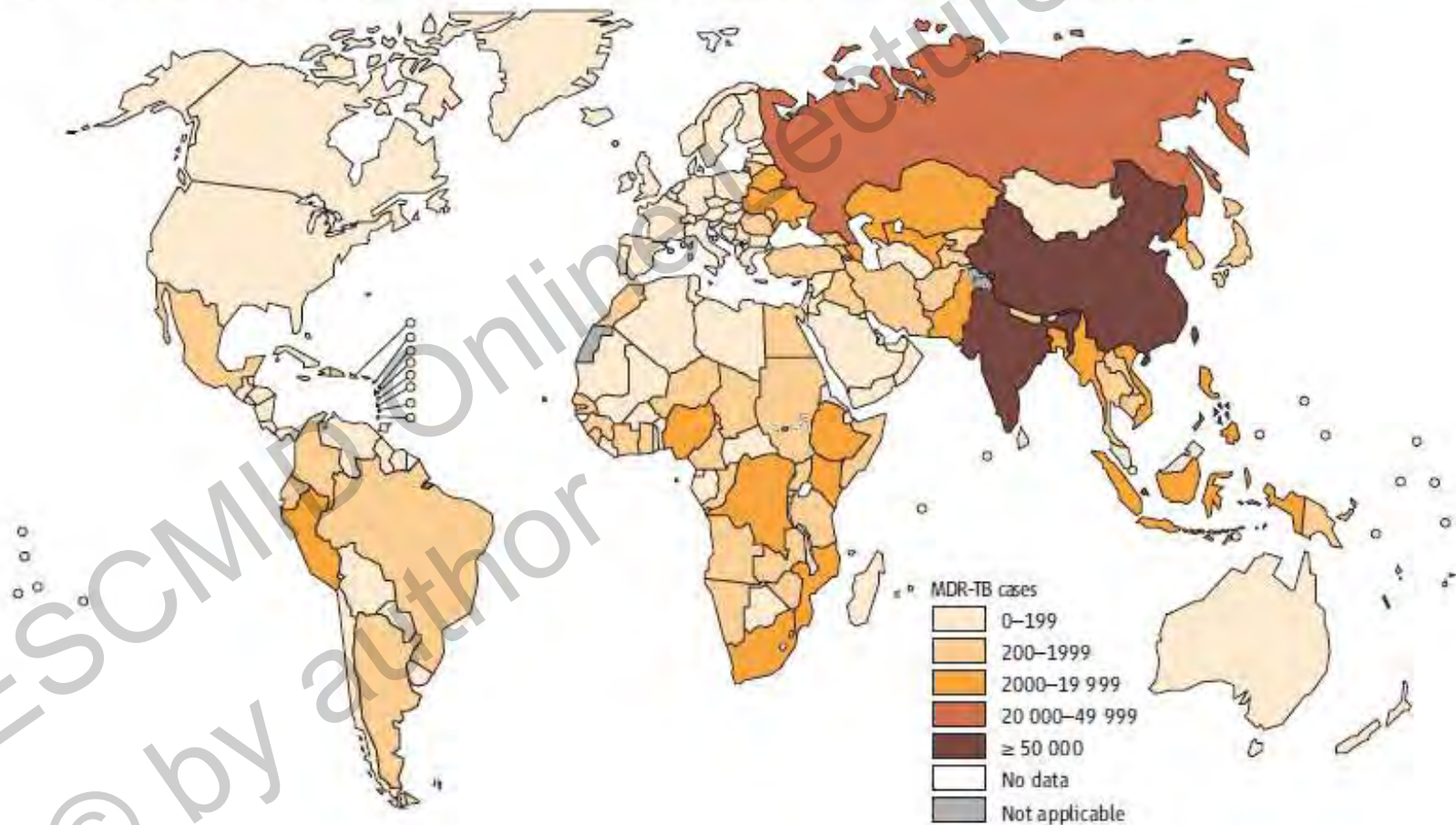
## 1.3 millions of deaths (15%)

Estimated TB incidence rates, 2012



450 000 MDR-TB cases (3.6% new cases  
and 20.2% previously treated cases)  
170 000 deaths (38%)

Number of MDR-TB cases estimated to occur among notified pulmonary TB cases, 2012

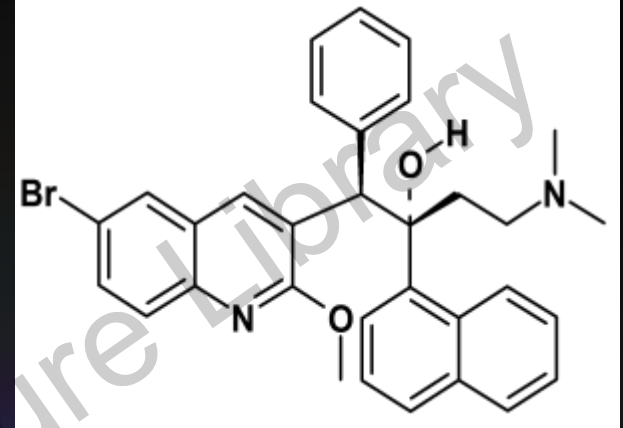


WHO Report 2013

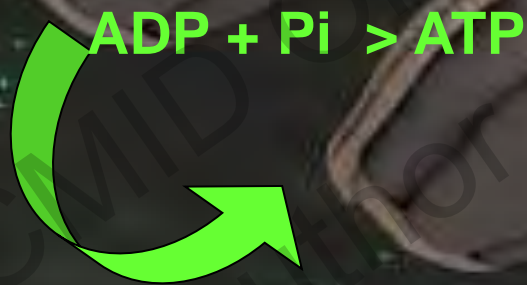


# Bedaquiline

diarylquinoline



Andries K et al.  
Science 2005



**Inhibition of subunit C of  
ATP synthase**

# In vitro activity of bedaquiline on *Mycobacterium tuberculosis*

Strains	N	MIC (mg/L)
fully susceptible	6	0.03-0.12
R rifampicin	1	0.03
R ethambutol	1	0.01
R pyrazinamide	1	0.03
R fluoroquinolones	3	0.06-0.12
R isoniazid	7	0.03-0.06
R rifampicin + isoniazid (MDR)	2	0.03

# No cross resistance with other antituberculous antibiotics

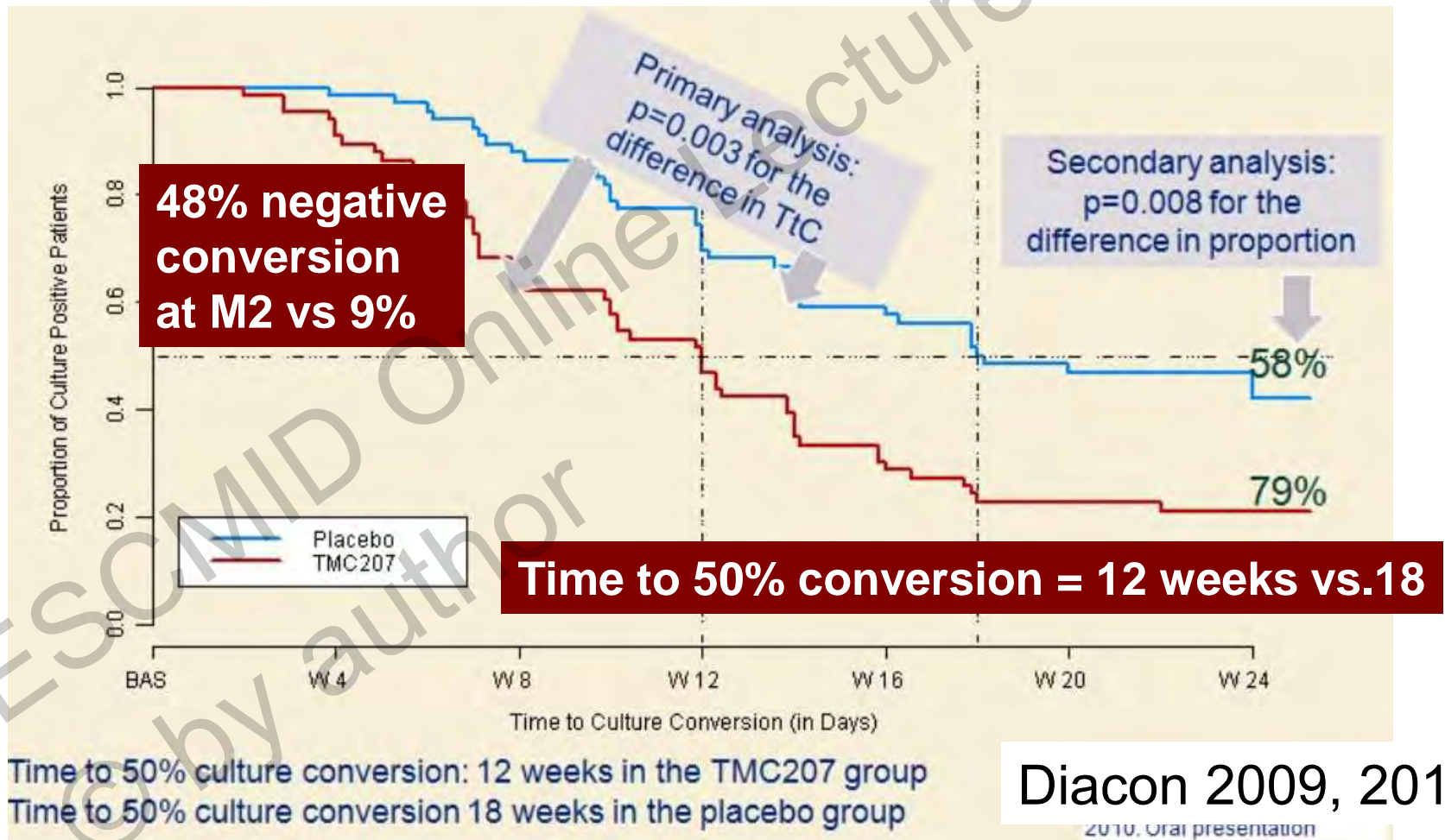
	MIC (mg/L) on 7H11	
	Wild type	Mutant BK12*
<b><u>Bedaquiline</u></b>	<b><u>0.03</u></b>	<b><u>4</u></b>
<b>Isoniazid</b>	<b>0.12</b>	<b>0.12</b>
<b>Rifampicin</b>	<b>0.5</b>	<b>0.12</b>
<b>Ethambutol</b>	<b>2</b>	<b>4</b>
<b>Streptomycin</b>	<b>1</b>	<b>1</b>
<b>Amikacin</b>	<b>1</b>	<b>2</b>
<b>Moxifloxacin</b>	<b>0.25</b>	<b>0.25</b>

\* : mutation A63P in *atpE*

Andries Science 2005

# Efficacy of bedaquiline in MDR-TB

- South Africa, 47 MDR patients
- 5 drugs -combination + bedaquiline or placebo



Diacon 2009, 2012

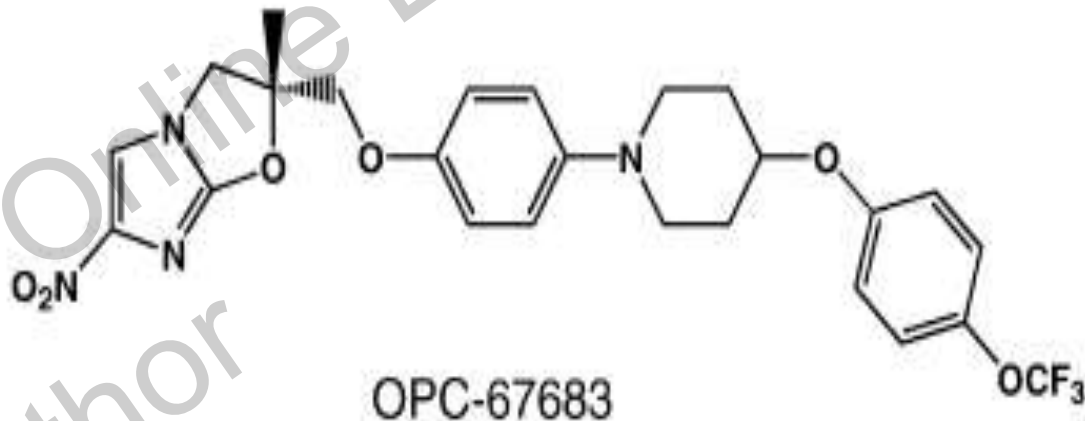


# Delamanid (OPC-67683)

Nitro-imidazole compound

Needs to be activated (reduction) in the bacterial cell

Inhibits mycolic acid synthesis, effect on DNA

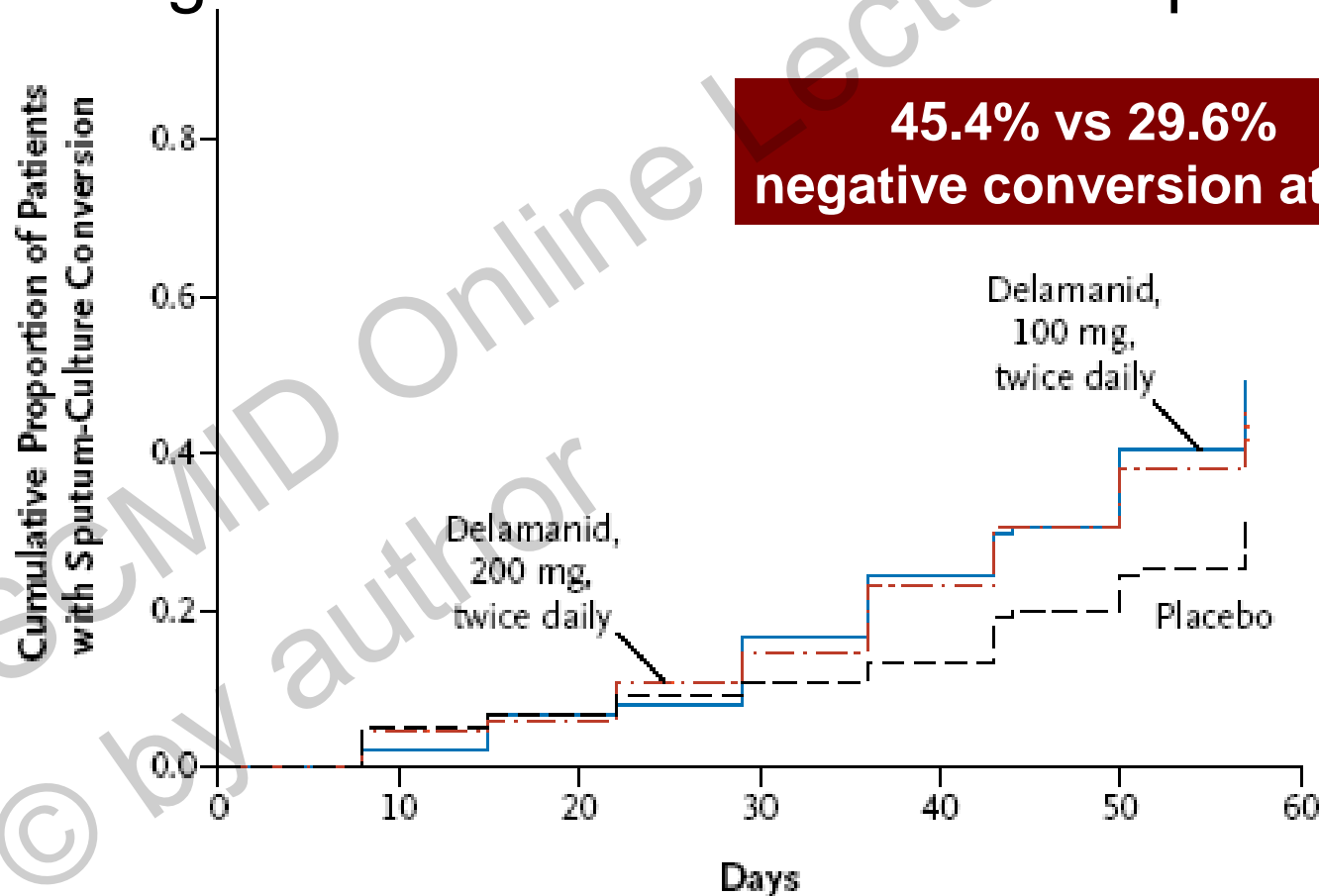


MIC on *M. tuberculosis* : 0.012 mg/L

Matsumoto PLOS Medicine 2006, Rakech PLOS one 2014

# Efficacy of delamanid in MDR TB

- 481 patients HIV- smear+ (Gler NEJM 2011)
- 5 drug-combination + delamanid or placebo



**45.4% vs 29.6%**  
**negative conversion at M2**

# New antituberculosis drugs, regimens, and adjunct therapies: needs, advances, and future prospects

Alimuddin I Zumla, Stephen H Gillespie, Michael Hoelscher, Patrick P J Philips, Stewart T Cole, Ibrahim Abubakar, Timothy D McHugh, Marco Schito, Markus Maeurer, Andrew J Nunn

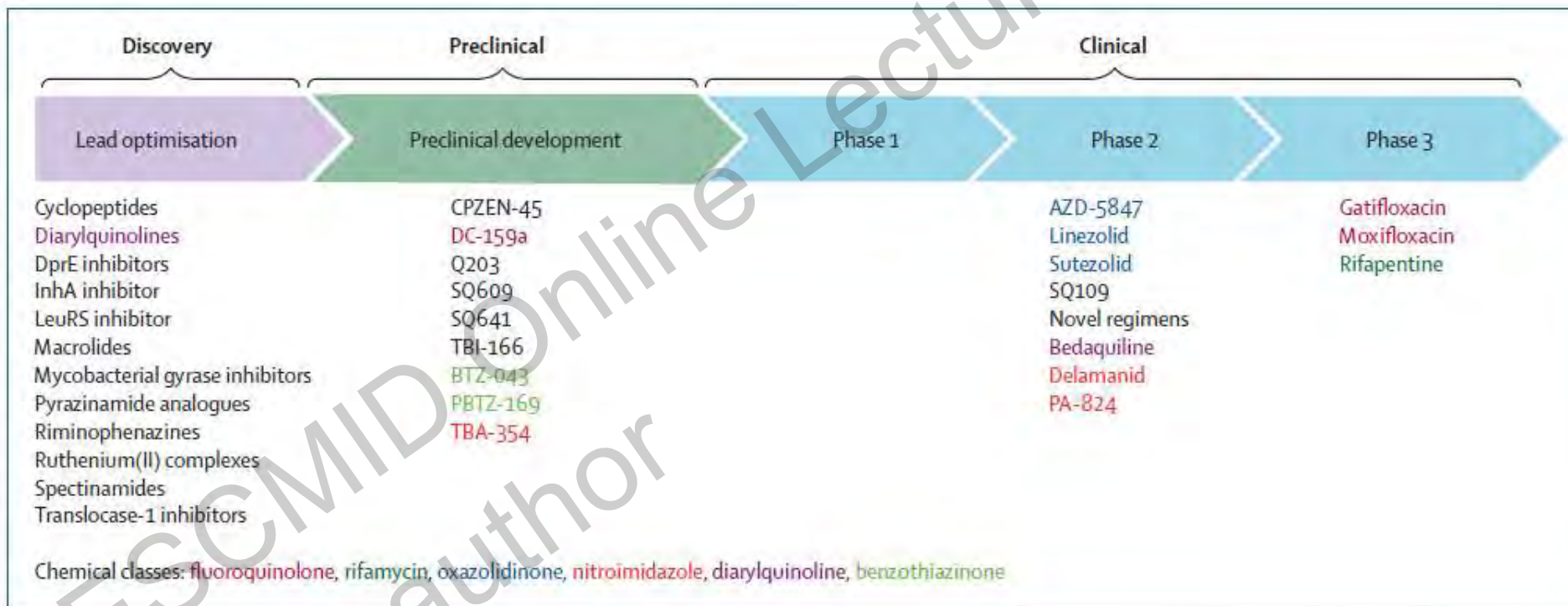


Figure 1: Global pipeline of new tuberculosis drugs<sup>2,22-24,31-48</sup>

Used with permission of the Stop TB Partnership Working Group on New Drugs.

Lancet Infect Dis 2014;  
14: 327-40

# Rapid molecular tests for diagnosis of MDR-TB

## **Xpert® MTB/RIF assay for pulmonary tuberculosis and rifampicin resistance in adults (Review)**

Steingart KR, Sohn H, Schiller I, Kloda LA, Boehme CC, Pai M, Dendukuri N

*Cochrane Database of Systematic Reviews* 2013;

<http://www.thecochranelibrary.com>

## **Advances in tuberculosis diagnostics: the Xpert MTB/RIF assay and future prospects for a point-of-care test**

Stephen D Lawn, Peter Mwaba, Matthew Bates, Amy Platek, Heather Alexander, Ben J Marais, Luis E Cuevas, Timothy D McHugh, Lynn Zijenah, Nathan Kapata, Ibrahim Abubakar, Ruth McNerney, Michael Hoelscher, Ziad A Memish, Giovanni Battista Migliori, Peter Kim, Markus Maeurer, Marco Schito, Alimuddin Zumla

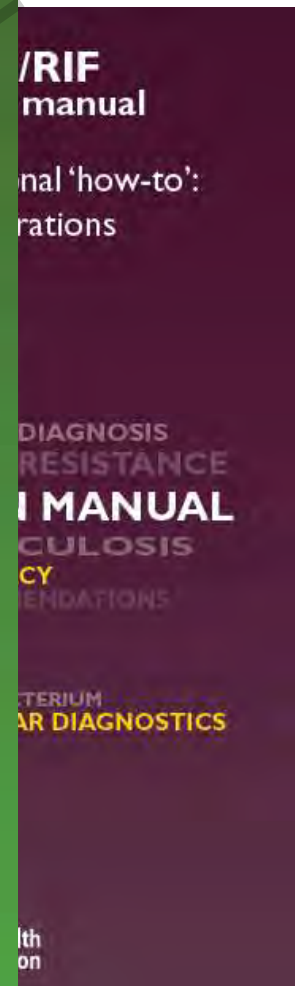
**Lancet Infect Dis** 2013;

**13: 349-61**

[www.thelancet.com/infection](http://www.thelancet.com/infection) Vol 13 April 2013



# WHO publications 2014





# Nucleic acid amplification direct testing (NAAT) for tuberculosis

- PCR started in 1990 using
  - gene (Hermans JCM 1990)
  - RNA (Boddinghaus JCM 1990)
  - IS6110 (Thierry JCM 1990)
- 1996 : recommendations CDC / ATS
- 2008 : recommendations CDC / JAMA
- Pubmed
  - 5000 papers, 350 review, 8 meta-analyses
  - Sarmiento 2003: meta-analysis on smear negative specimens
  - Greco 2009: meta-analysis on smear positive specimens
  - Lawn 2013: review on Xpert®MTB/RIF®

**ATS 1997, Ieven and Goosens 1997, Sarmiento 2003,  
Dinnes 2007, MMWR 2009;58-7-10, Lawn S LID 2013**

# NAAT sensitivity for TB diagnosis

= We would like that all TB patients with Smear-positive specimens are NAAT+

- Cochrane review
  - Pooled sensitivity = **98%** (95% CrI 97- 99%)
- WHO study group review
  - Pooled sensitivity = **98%** (95%CrI 97-99%)

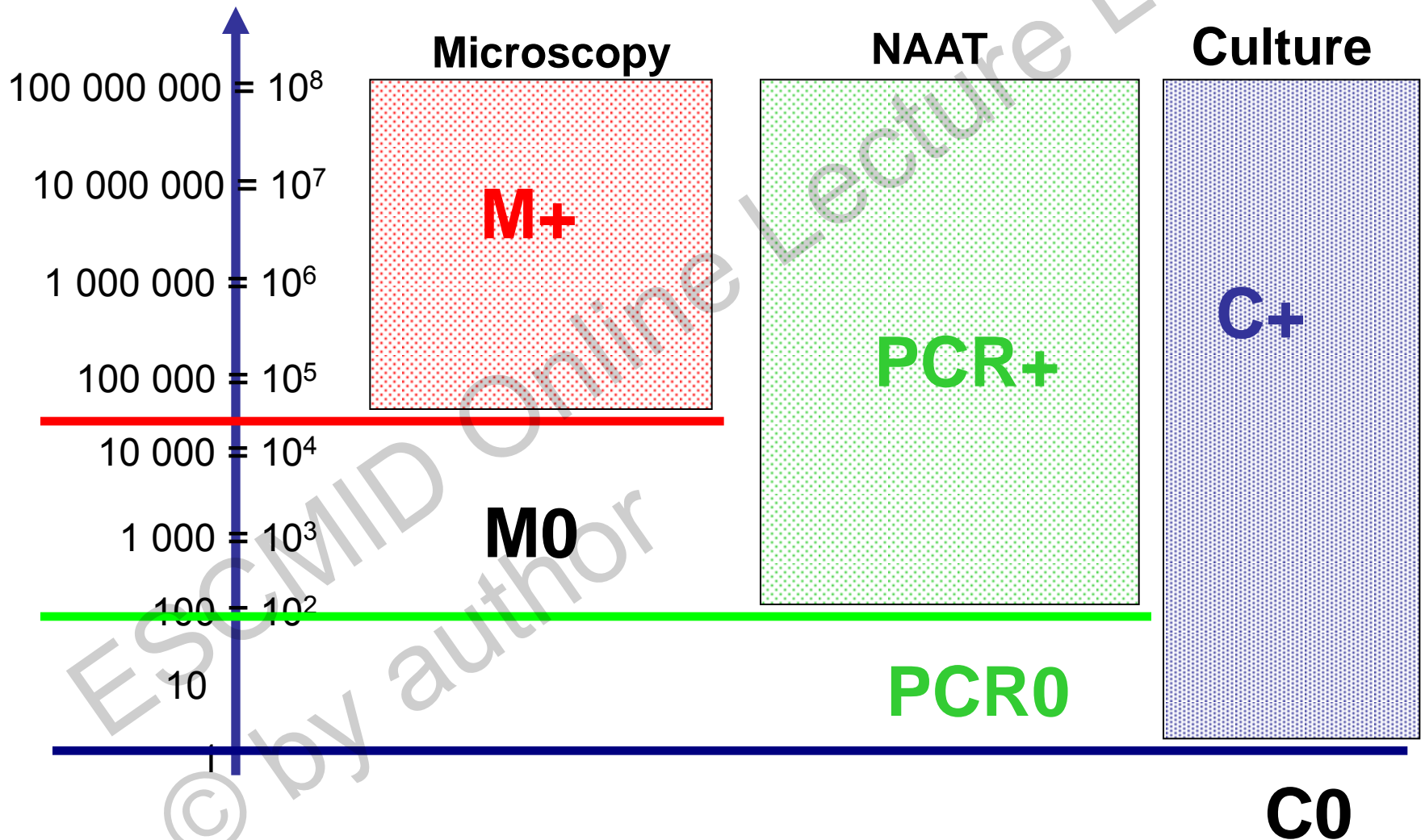
# NAAT sensitivity for TB diagnosis

= We would like that even TB patients with Smear-negative specimens are NAAT+

- **Cochrane review**
  - Sensitivity : **67%** (95% CrI 58% -74%)
- **LID Review (S. Lawn et al. 2013)**
  - Sensitivity : **75%** (range: 47% - 83%)
  - Extrapulmonary TB : **77%** (range 25%-97%)
- **WHO expert group review**
  - Sensitivity : **68%** (95% CrI 61%-74%)

# Sensitivity of NAAT for detecting tuberculosis bacilli

N per ml specimen



# NAAT Performances for TB: specificity

= We would like that patients without TB are NAAT negative

- Cochrane review
  - Specificity : **98%** (95% CrI 97% to 99%)
- LID Review
  - Specificity : **98.6%** (range 88.9% - 100%)
- WHO study group review
  - Specificity : **99%** (95% CrI 98% to 99%)



# Molecular detection of rifampicin resistance

Sensitivity = all Resistant strains are detected  
Specificity = no Susceptible strains are detected

- Cochrane review
  - sensitivity : **94%** (95% CrI 87% to 97%)
  - specificity : **98%** (95% CrI 97% to 99%)
- WHO study group review
  - sensitivity : **95%** (95%CrI 90-97%)
  - specificity : **98%** (95%CrI 97-99%)

# Positive predictive value for detection of rifampicin resistance

For 1000 patients and a specificity of 98%

Prevalence of resistance	30%	2%
Nb of resistant isolates	300	20
Nb of false resistant test	14	20
PPV of detection of rifampicin resistance	96%	50%

# Positive predictive value for detection of rifampicin resistance

For 1000 patients and a specificity of 98%

Prevalence of resistance	30%	2%
Nb of resistant isolates	300	20
Nb	<b>Confirmation with another molecular test and with phenotypic determination is mandatory</b>	
PPV of detection of rifampicin resistance	96%	50%

# The new clinical bacteriology laboratory





Automated susceptibility testing



Identification

Inoculation  
Culture  
incubation



Blood culture  
automate

Whole genome sequencing

Molecular  
detection





# Point-Counterpoint: The Automated Clinical Microbiology Laboratory: Fact or Fantasy?

N.A. Ledebøer, S. Dallas, P.H. Gilligan

JCM Accepts, published online ahead of print on 19 March 2014

J. Clin. Microbiol. doi:10.1128/JCM.00686-14

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## Pros

- **Cost saving for hospital directors (less technicians)**
- **Less skilled workers**
  - **Higher throughput**
  - **Image collection**
- **Can work 24/24**

## Cons

- **High cost for machines**
  - **Large footprints**
- **Less skilled workers**
  - **Not adapted to the emergency needed for infections (batch working)**
- **Not include molecular methods or new techniques**

# Point-Counterpoint: The Automated Clinical Microbiology Laboratory: Fact or Fantasy?

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## Pros

- **Cost saving for hospital**

## Cons

- **High cost for machines**

There are no data thus far that shows that automation of microbiologic culture processes improves patient outcomes

• **Image collection**

- **Can work 24/24**

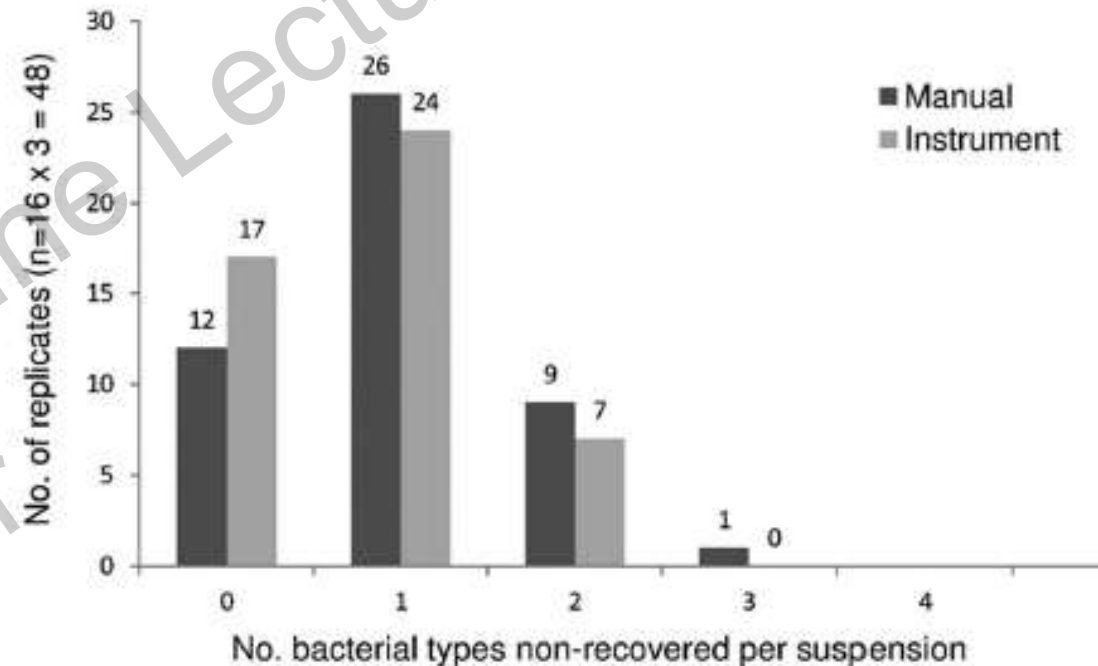
**Infections (batch working)**

- **Not include molecular methods or new techniques**

# Automated versus Manual Sample Inoculations in Routine Clinical Microbiology: a Performance Evaluation of the Fully Automated Inoqula Instrument

P. Froment, H. Marchandin, P. Vande Perre and B. Lamy  
*J. Clin. Microbiol.* 2014, 52(3):796. DOI:  
10.1128/JCM.02341-13.

16 Polymicrobial suspensions  
48 Replicates  
2 to 4 different bacteria



The automate improved the quality and standardization of bacterial isolation from monomicrobial and polymicrobial clinical specimens

## THE WORLD SEPSIS DECLARATION

*Sepsis is one of the most common, least-recognized illnesses in both the developed and developing world. Globally, 20 to 30 million patients are estimated to be afflicted every year, with over 6 million cases of neonatal and early childhood sepsis and over 100,000 cases of maternal sepsis.*

*Worldwide,*

*a person dies from sepsis every few seconds.*

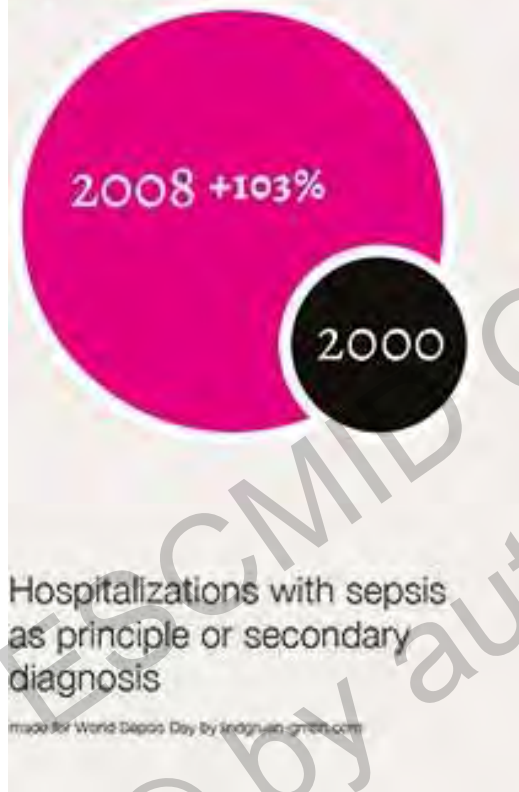
[www.world-sepsis-day.org](http://www.world-sepsis-day.org)

GSA  
WFSICOM  
WFPIOCS  
WFOCN  
ISF  
Sepsis Alliance  
IFEM  
AAP  
GSS  
CSOC

September | World  
13 | Sepsis  
2013 | Day

# Do we have good diagnosis tests for sepsis in CM labs?

The incidence of sepsis is rising dramatically



**Sepsis is a medical emergency**

**Unfortunately, sepsis is still often overlooked and recognized too late.**

Hospital costs of sepsis have doubled



Early sepsis treatment is cost effective. It reduces the number of hospital and critical care bed days for patients.



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## Work Flow Analysis of Around-the-clock Processing of Blood Culture Samples and Integrated MALDI-TOF Mass Spectrometry Analysis for the Diagnosis of Bloodstream Infections

Wilhelm Schneiderhan,<sup>1</sup> Alexander Grundt,<sup>1</sup> Stefan Wörner,<sup>1,2</sup> Peter Findeisen,<sup>1</sup> and Michael Neumaier<sup>1\*</sup>

## Comparison of 2 Blood Culture Media Shows Significant Differences in Bacterial Recovery for Patients on Antimicrobial Therapy

790 • CID 2013:56

Rebecca Zadroga,<sup>1,2</sup> David N. Williams,<sup>2,3</sup> Richard Gottschall,<sup>4</sup> Kevan Hanson,<sup>4</sup> Vickie Nordberg,<sup>4</sup> Marcia Deike,<sup>4</sup> Mike Kuskowski,<sup>5</sup> Lisa Carlson,<sup>6</sup> David P. Nicolau,<sup>7</sup> Christina Sutherland,<sup>7</sup> and Glen T. Hansen<sup>2,4,8</sup>

## Emerging Technologies for Rapid Identification of Bloodstream Pathogens

Atul Kothari<sup>1</sup>, Margie Morgan<sup>2</sup>, David Haake<sup>1,3</sup>

Clinical Infectious Diseases Advance Access published April 24, 2014

# Emerging technologies for rapid identification of bloodstream infections

Kothari A. et al, CID 2014

## List of techniques that can be used

Table 1. Tests for Rapid Identification of Bloodstream Pathogens

Blood Culture Assays	Pathogens Detected	Resistance Markers	Turnaround Time (after blood cultures turn positive)
PNA-FISH	<i>Staphylococcus aureus</i> , CoNS, <i>Enterococcus fecalis</i> , Other Enterococci, <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> , <i>Pseudomonas aeruginosa</i> , <i>Candida albicans</i> , <i>Candida parapsilosis</i> , <i>Candida glabrata</i> , <i>Candida krusei</i> , <i>Candida tropicalis</i>	No	1.5-3 hours
QuickFISH	<i>Staphylococcus aureus</i> , CoNS, <i>Enterococcus fecalis</i> , Other Enterococci, <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> , <i>Pseudomonas aeruginosa</i>	No	<30 minutes
MALDI-ToF*	Gram positive and negative bacteria, yeast, fungi, filamentous fungi, Mycobacteria	In Development	10-30 minutes
Gene Xpert MRSA/SA	<i>Staphylococcus aureus</i>	<i>mec A</i>	<1 hour
Verigene Gram Positive Blood Culture (BC-GP)	<i>Staphylococcus aureus</i> , <i>Staphylococcus epidermidis</i> , <i>Staphylococcus lugdunensis</i> , <i>Streptococcus anginosus</i> group, <i>Streptococcus agalactiae</i> , <i>Streptococcus pneumoniae</i> , <i>Streptococcus pyogenes</i> , <i>Enterococcus fecalis</i> , <i>Enterococcus faecium</i> , <i>Staphylococcus spp.</i> , <i>Streptococcus spp.</i> , <i>Listeria spp.</i>	<i>mec A</i> , van A, van B	2.5 hours
Verigene Gram Negative Blood Culture (BC-GN)**	<i>Escherichia coli</i> , <i>Shigella spp.</i> , <i>Klebsiella pneumoniae</i> , <i>Klebsiella oxytoca</i> , <i>Pseudomonas aeruginosa</i> , <i>Serratia marcescens</i> , <i>Aerobacter spp.</i> , <i>Proteus spp.</i> , <i>Citrobacter spp.</i> , <i>Enterobacter spp.</i>	KPC, NDM, CTX-M, VIM, IMP, OXA	2 hours
Film Array Blood Culture Identification (BC ID)	<i>Staphylococcus aureus</i> , <i>Staphylococcus spp.</i> , <i>Streptococcus agalactiae</i> , <i>Streptococcus pneumoniae</i> , <i>Streptococcus pyogenes</i> , <i>Streptococcus spp.</i> , <i>Enterococcus spp.</i> , <i>Listeria monocytogenes</i> , <i>Hemophilus influenzae</i> , <i>Neisseria meningitidis</i> , <i>Enterobacter cloacae</i> complex, <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> , <i>Klebsiella oxytoca</i> , <i>Pseudomonas aeruginosa</i> , <i>Serratia marcescens</i> , <i>Aerobacter baumannii</i> , <i>Proteus</i> , <i>Candida albicans</i> , <i>Candida glabrata</i> , <i>Candida krusei</i> , <i>Candida parapsilosis</i> , <i>Candida tropicalis</i>	<i>mec A</i> , van A, van B	1 hour
<b>Whole Blood Assays</b>	<b>Pathogens Detected</b>	<b>Resistance Markers</b>	<b>Turnaround Time</b>
Light Cycler*** SeptiFast	<i>Staphylococcus aureus</i> , CoNS, <i>Streptococcus spp.</i> , <i>Streptococcus pneumoniae</i> , <i>Enterococcus fecalis</i> , <i>Enterococcus faecium</i> , <i>Escherichia coli</i> , <i>Klebsiella pneumoniae</i> , <i>Klebsiella oxytoca</i> , <i>Pseudomonas aeruginosa</i> , <i>Enterobacter cloacae / aerogenes</i> , <i>Serratia marcescens</i> , <i>Aerobacter baumannii</i> , <i>Proteus mirabilis</i> , <i>Stenotrophomonas maltophilia</i> , <i>Candida albicans</i> , <i>Candida glabrata</i> , <i>Candida krusei</i> , <i>Candida parapsilosis</i> , <i>Candida tropicalis</i> , <i>Aspergillus fumigatus</i>	No	6 hours
SeptiTest***	>300 different pathogens	No	8-12 hours
T2 Candida****	<i>Candida albicans</i> , <i>Candida glabrata</i> , <i>Candida krusei</i> , <i>Candida parapsilosis</i> , <i>Candida tropicalis</i>	No	3 hours

\*Biomérieux instrument FDA approved, Bruker pending approval

\*\*pending FDA approval

\*\*\*not available in US

# Emerging technologies for rapid identification of bloodstream infections

Kothari A. et al, CID 2014

## List of studies examining impacts on outcomes

Table 2. Studies Examining Impacts on Clinical Outcomes and Healthcare Costs

Intervention	Year of Publication	Antimicrobial Stewardship Intervention	Mortality Benefit	Length of Stay	Cost Saving per patient	Reference
PNA-FISH	2006	Yes	Not studied	-2 days	\$4005	Forrest, 2006 [25]
PNA-FISH	2011	No	Not studied	+2.2 days	Not studied	Holtzman 2011 [27]
PNA-FISH	2006	Yes	Not studies	Not studied	\$1729*	Forrest 2006 [28]
PNA-FISH	2008	No	Yes (16.8% vs 7.9%)	-2 days*	\$19441*	Ly 2008 [26]
Gene Xpert MRSA/SA	2010	Yes	Yes (18% vs 26%)*	-6.2 days	\$21387	Bauer 2010 [30]
MALDI-ToF	2012	Yes	Yes (5.6% vs 10.7%)*	-1.8 days	\$19547*	Perez 2012 [31]
MALDI-ToF	2013	Yes	Yes (12.7% vs 20.3%)	-2.8 days*	Not studied	Huang 2013 [33]
Verigene BC-GP	2013	Yes	No	-21.7 days	\$60729	Sango 2013 [35]

\*not statistically significant



# Emerging technologies for rapid identification of bloodstream infections

Kothari A. et al, CID 2014

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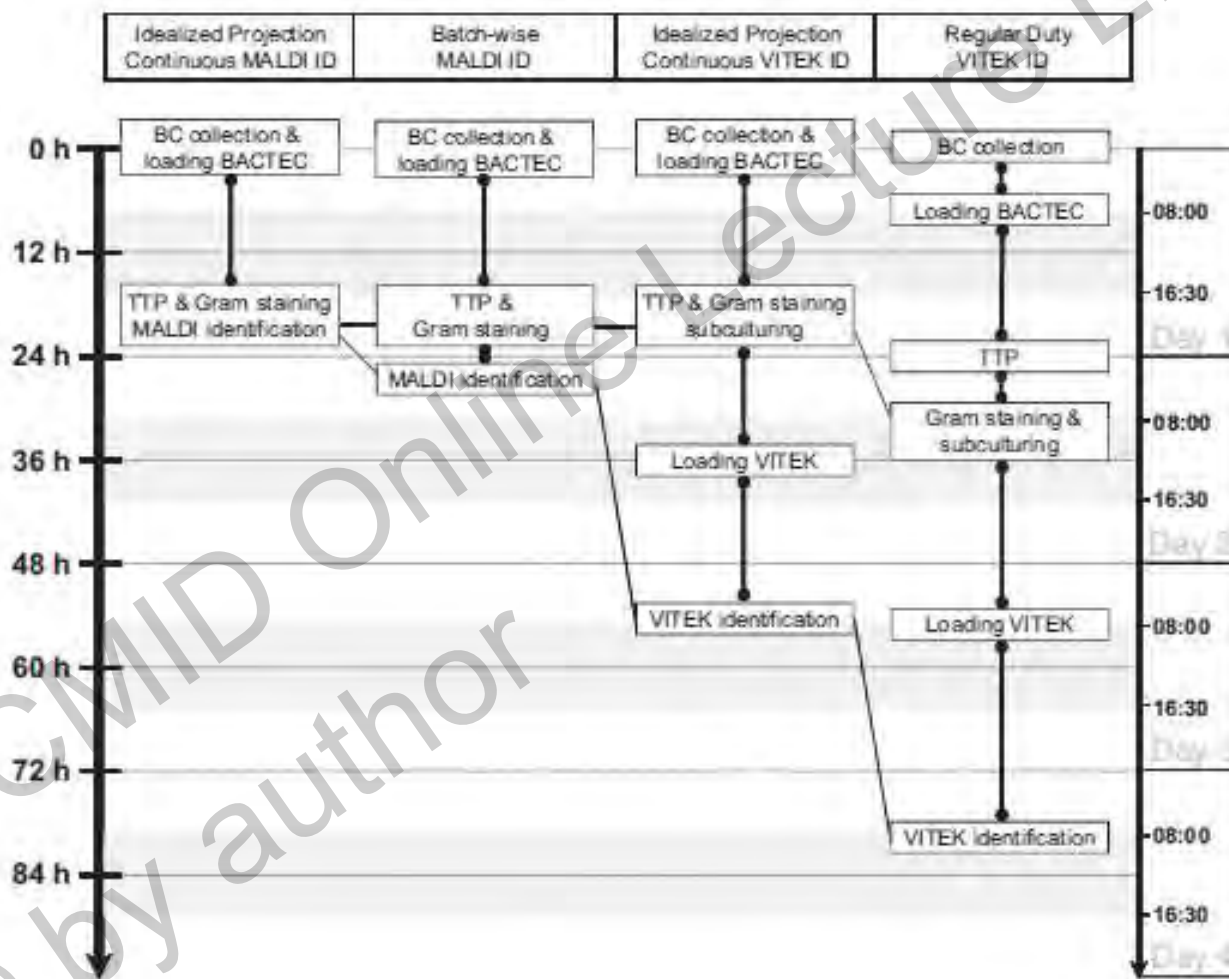
### Impact of rapid technologies on

- Mortality
- Cost
- length of stay in hospital

Needs to be more often examined

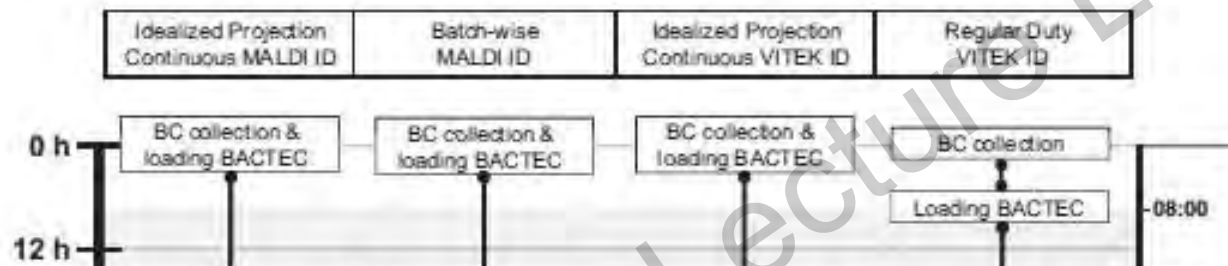
\*not statistically significant

# Workflow for detection and identification of pathogens in Blood cultures



Schneiderhan et al. 2013

# Workflow for detection and identification of pathogens in Blood cultures



**Continuous workflow for culture and identification can shorten time-to-results from 4h to 59 hours with regard to the discontinuous setting**

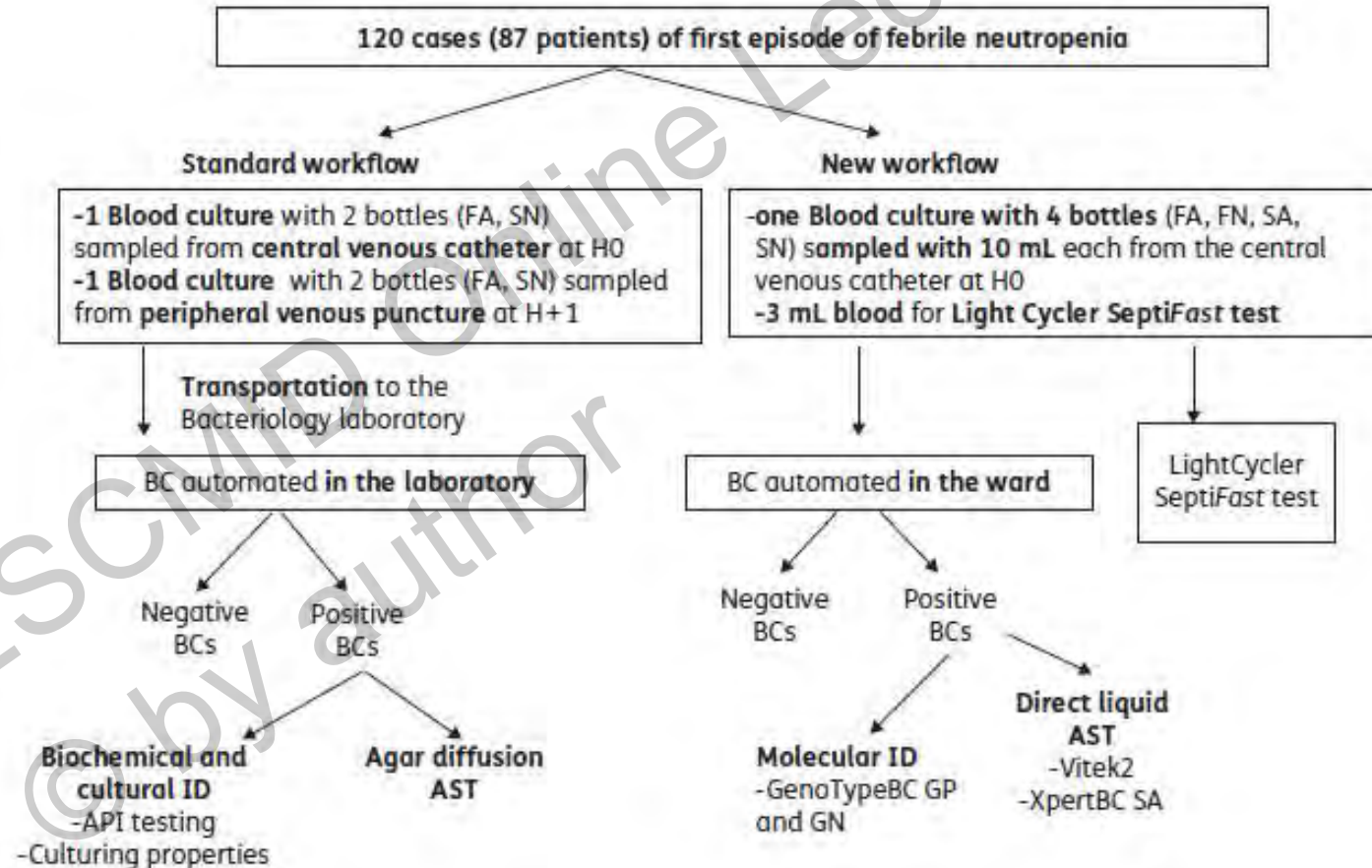




# A new workflow for the microbiological diagnosis of febrile neutropenia in patients with a central venous catheter

Cecile Pautas<sup>1</sup>, Emilie Sbidian<sup>2†</sup>, Yosr Hicheri<sup>1,3†</sup>, Sylvie Bastuji-Garin<sup>2,4</sup>, Stéphane Bretagne<sup>2,5</sup>, Celine Corbel<sup>6</sup>, Laetitia Gregoire<sup>7</sup>, Sébastien Maury<sup>1,2</sup>, Lilia Merabet<sup>6</sup>, Catherine Cordonnier<sup>1,2</sup> and Emmanuelle Cambau<sup>6,8,9\*†</sup>

Journal of Antimicrobial Chemotherapy 2013



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Journal of Antimicrobial Chemotherapy 2013

120 cases (87 patients) of first episode of febrile neutropenia

Standard  
Process

Ideal  
process  
with  
continuous  
workflow

# Continuous workflow decreased time-to-results to less than 24h

**Table 1.** Turnaround times for blood culture and identification and susceptibility testing results for 29 episodes positive with both wor

Time to microbiological results	Median time (range)	
	standard workflow	study workflow
Time to BC-positive signal <sup>a</sup>		
all episodes (n=29)	13 h 01 (9 h 31–43 h 33)	12 h 25 (7 h 55–25 h 37)
episodes with streptococci (n=13)	10 h 39 (9 h 31–13 h 37)	9 h 55 (7 h 55–17 h 55)
episodes with CoNS (n=7)	20 h 13 (17 h–43 h 27)	16 h 24 (14 h 34–25 h 18)
episodes with other bacteria (n=9)	14 h 10 (11 h 18–21 h 55)	16 h 54 (11 h 18–25 h 37)
Time to results of identification <sup>b</sup>		
all episodes (n=26)	69 h (22 h–405 h)	18 h 16 (11 h 74–55 h 37)
episodes with streptococci (n=11)	156 h (82 h 48–405 h)	15 h 55 (12 h 57–47 h 55)
episodes with CoNS (n=6)	66 h (50 h–75 h)	21 h 58 (18 h 53–31 h 18)
episodes with other bacteria (n=9)	36 h 12 (22 h–69 h)	21 h 55 (16 h 20–37 h 23)
Time to results of antibiotic susceptibility testing (AST)		
methicillin-resistance in staphylococci (n=9) <sup>c</sup>	70 h 36 (50 h–130 h)	17 h 30 (15 h 51–31 h 18)
all episodes with valid AST results (n=18) <sup>d</sup>	56 h (22 h–245 h)	22 h 10 (13 h 10–43 h 37)

\* Pautas et al. JAC 2013

# Continuous workflow decreased time-to -results to less than 24h

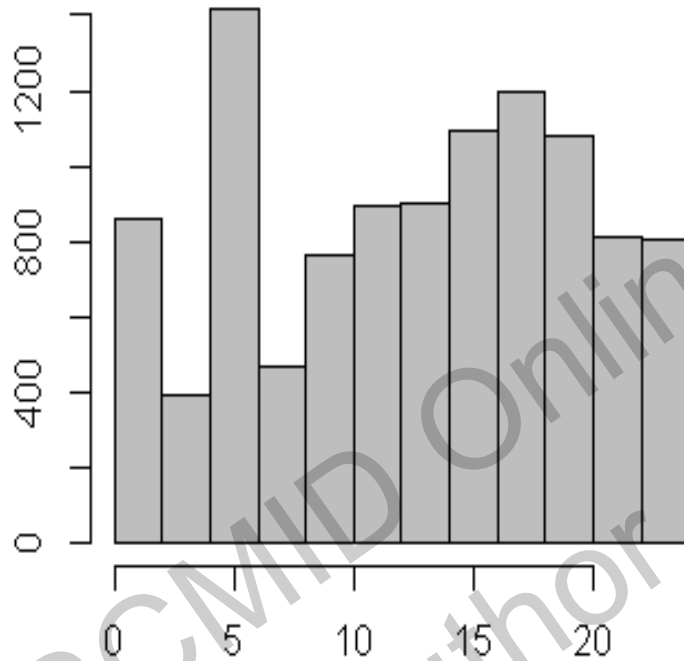
**Table 1.** Turnaround times for blood culture and identification and susceptibility testing results for 29 episodes positive with both wor

Time to microbiological results	Median time (range)	
	standard workflow	study workflow
Time to BC-positive signal <sup>a</sup>		
all episodes		
episodes with <i>C. albicans</i> (n=8)	68 h (30 h-73 h)	21 h 36 (18 h 33-31 h 18)
episodes with other bacteria (n=9)	36 h 12 (22 h-69 h)	21 h 55 (16 h 20-37 h 23)
Time to results of antibiotic susceptibility testing (AST)		
methicillin-resistance in staphylococci (n=9) <sup>c</sup>	70 h 36 (50 h-130 h)	17 h 30 (15 h 51-31 h 18)
all episodes with valid AST results (n=18) <sup>d</sup>	56 h (22 h-245 h)	22 h 10 (13 h 10-43 h 37)

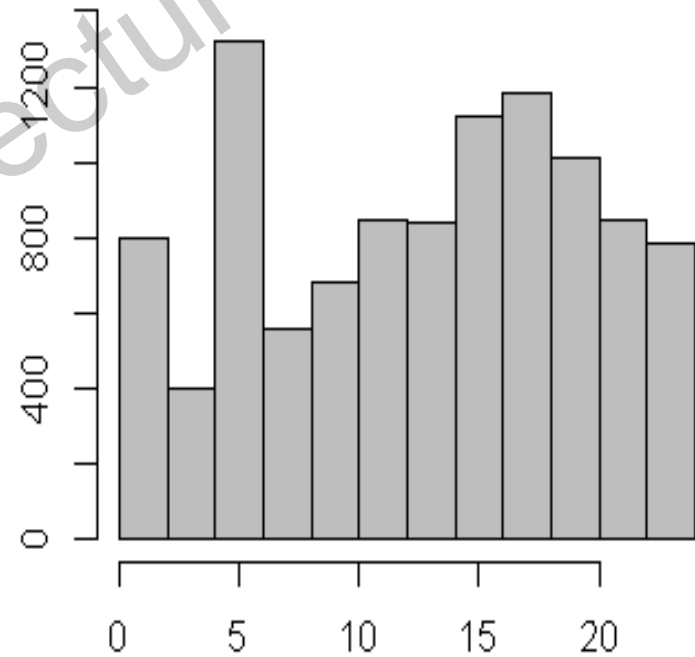
**Median time to positivity of BC = 12h25**  
**Median time to identification = 18h16**  
**Median time to AST results = 22h10**

\* Pautas et al. JAC 2013

Have you looked at what time the blood cultures were collected?



2011

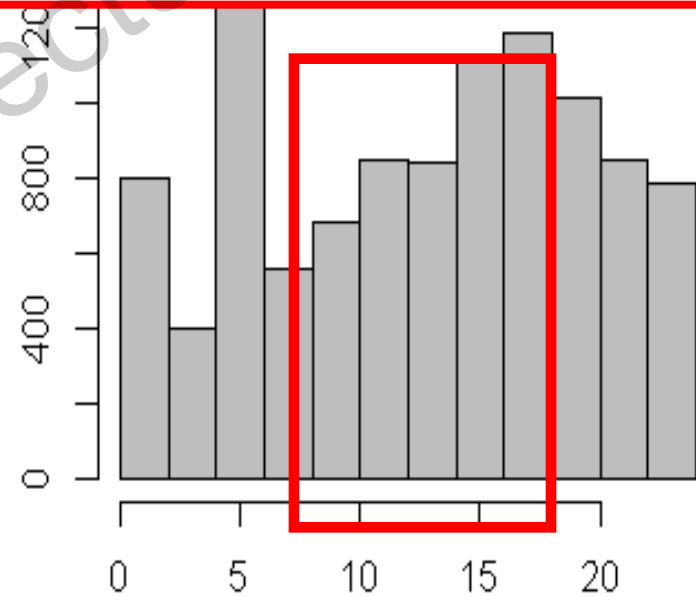
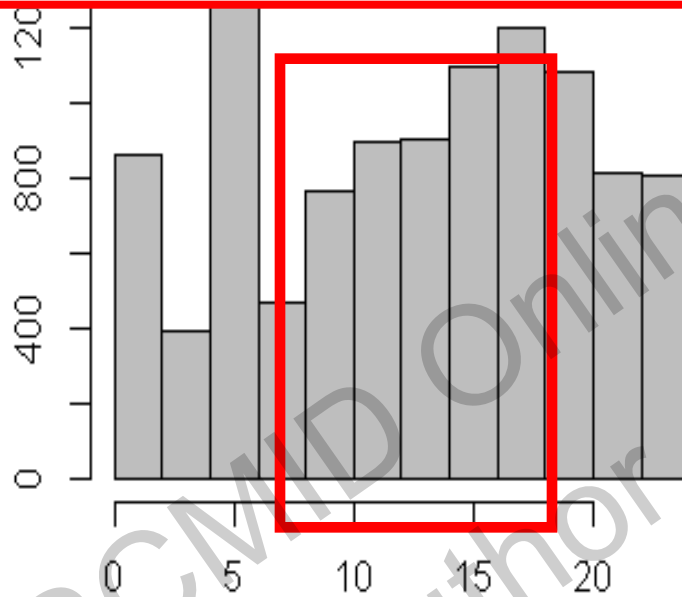


2012

Data for Lariboisière-Fernand Widal, 1000-bed University Hospital

# Have a look at the time when the blood cultures are collected!

**50% of BC are sampled during the day**



**50% of BC are sampled during the night**



# MDR bacteria are still evolving crazy

[Proc Natl Acad Sci U S A](#). 2014 Apr 1;111(13):4988-93. doi: 10.1073/pnas.1321364111. Epub 2014 Mar 17.

## **Molecular dissection of the evolution of carbapenem-resistant multilocus sequence type 258 *Klebsiella pneumoniae*.**

[Deleo FR](#)<sup>1</sup>, [Chen L](#), [Porcella SF](#), [Martens CA](#), [Kobayashi SD](#), [Porter AR](#), [Chavda KD](#), [Jacobs MR](#), [Mathema B](#), [Olsen RJ](#), [Bonomo RA](#), [Musser JM](#), [Kreiswirth BN](#).  
[Author information](#)

[N Engl J Med](#). 2014 Apr 17;370(16):1524-31. doi: 10.1056/NEJMoa1303359.

## **Transferable vancomycin resistance in a community-associated MRSA lineage.**

[Rossi F](#)<sup>1</sup>, [Diaz L](#), [Wollam A](#), [Panesso D](#), [Zhou Y](#), [Rincon S](#), [Narechania A](#), [Xing G](#), [Di Gioia TS](#), [Doi A](#), [Tran TT](#), [Reyes J](#), [Munita JM](#), [Carvajal LP](#), [Hernandez-Roldan A](#), [Brandão D](#), [van der Heijden IM](#), [Murray BE](#), [Planet PJ](#), [Weinstock GM](#), [Arias CA](#).

# Threat of antibiotic resistance

Confronting the threat of multidrug-resistant Gram-negative bacteria  
in critically ill patients

Jonathan Cohen\*

*J Antimicrob Chemother* 2013; 68: 490–491

[Ann N Y Acad Sci](#). 2014 Apr 16. doi: 10.1111/nyas.12399. [Epub ahead of print]

**The risk/benefit of predicting a post-antibiotic era: Is the alarm working?**

[Fowler T](#)<sup>1</sup>, [Walker D](#), [Davies SC](#).

[Soc Sci Med](#). 2014 Jun;110:81-8. doi: 10.1016/j.socscimed.2014.03.030. Epub 2014 Mar 28.

**Cultures of resistance? A Bourdieusian analysis of doctors' antibiotic prescribing.**

[Broom A](#)<sup>1</sup>, [Broom J](#)<sup>2</sup>, [Kirby E](#)<sup>3</sup>.

# World Alliance Against Antibiotic Resistance (WAAAR)

1. Awareness of antibiotic resistance (ATB-R)
2. Financed national plans for fighting against ATB-R
3. Antibiotics of assured quality in every part of the world
4. Integrated surveillance of ATB-R and ATB use
5. Development of new diagnostic tests
6. Revisit the use of ATB and its business
7. Educational efforts to change
8. Prevention of ATB-R bacteria transmission including environment
9. Increase research and drug development
10. “lives saved by antibiotics” in the list of the world immaterial heritage (UNESCO)

Thank you for your attention

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