

# What is the Future of Microbiologic Diagnosis?



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## Déclaration d'intérêts

- **Contracted Research: ContraFect, TenNor Therapeutics Limited, and BioFire**
- **Consultant: Curetis, PathoQuest, Selux Diagnostics, 1928 Diagnostics, PhAST, Torus Biosystems, Day Zero Diagnostics, Mammoth Biosciences, CARB-X, Qvella, Netflix**
- **Mayo Clinic and I have a relationship with Adaptive Phage Therapeutics and Pathogenomix**
- **Patents: *Bordetella pertussis/parapertussis* PCR; device/method for sonication; anti-biofilm substance**

# Outline

- **The challenge**
- **Technology revolution**
  - Proteomics
  - NAATs & sequencing-based diagnostics
  - Other diagnostic types (imaging, metabolomic)
  - Point of care diagnostics
  - Advanced host response assessment for infectious diseases
- **How to move to the future**
  - Demonstration of clinical utility
- **The deliverables**
  - Improved health
  - Changes to healthcare delivery
  - Better understanding of infectious diseases and their mimics

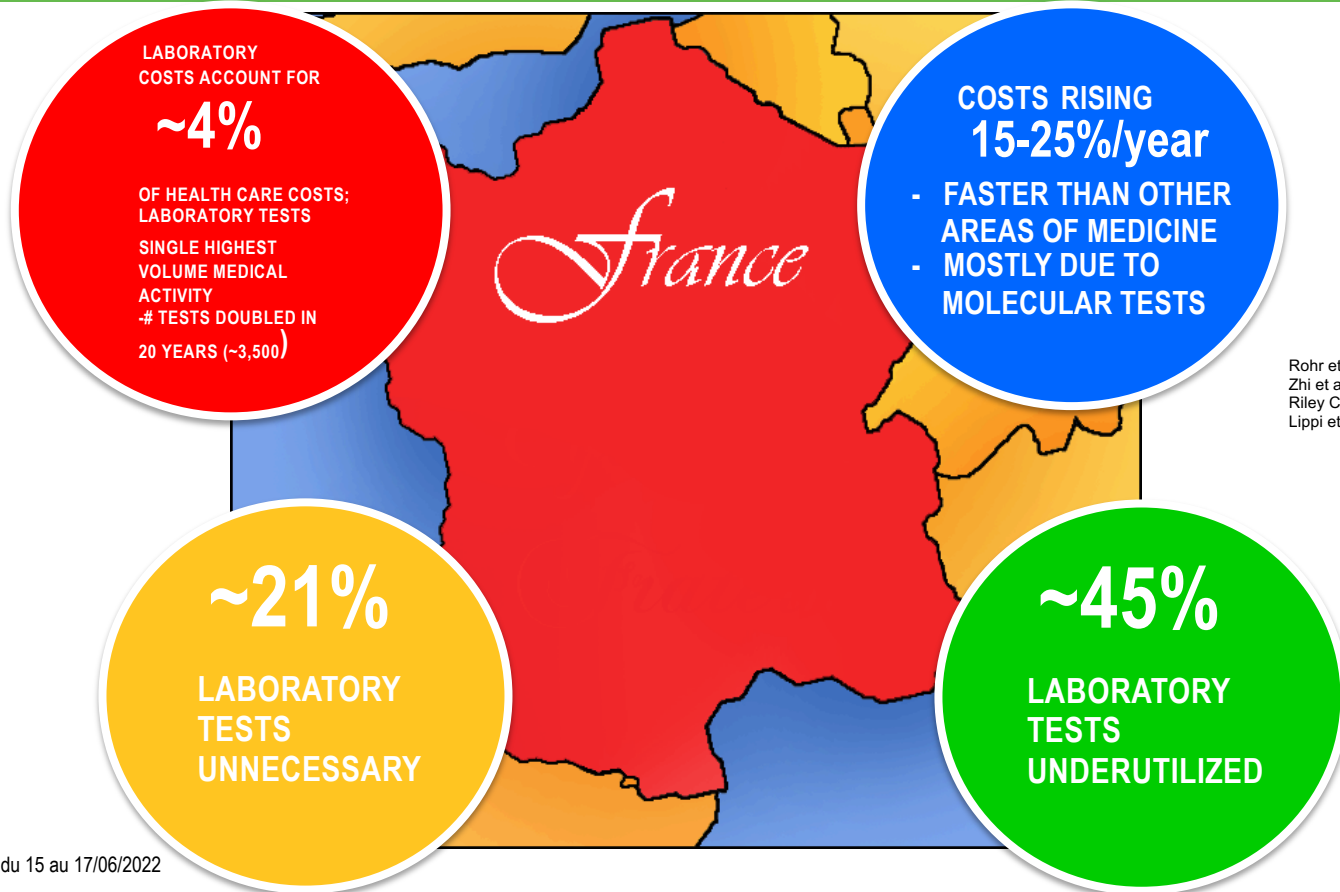
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# The Challenge

- **The need:**
  - Antimicrobial resistance
  - Emerging infectious diseases
  - Availability of better therapeutics
- **The barriers:**
  - Infectious diseases management predicated on “culture of empiricism”
  - Inappropriate test utilization
  - Lack of understanding of value of advanced diagnostics
  - Cost of advanced diagnostics

# 70% Clinical Decisions Substantially Based on Results of Diagnostic Tests

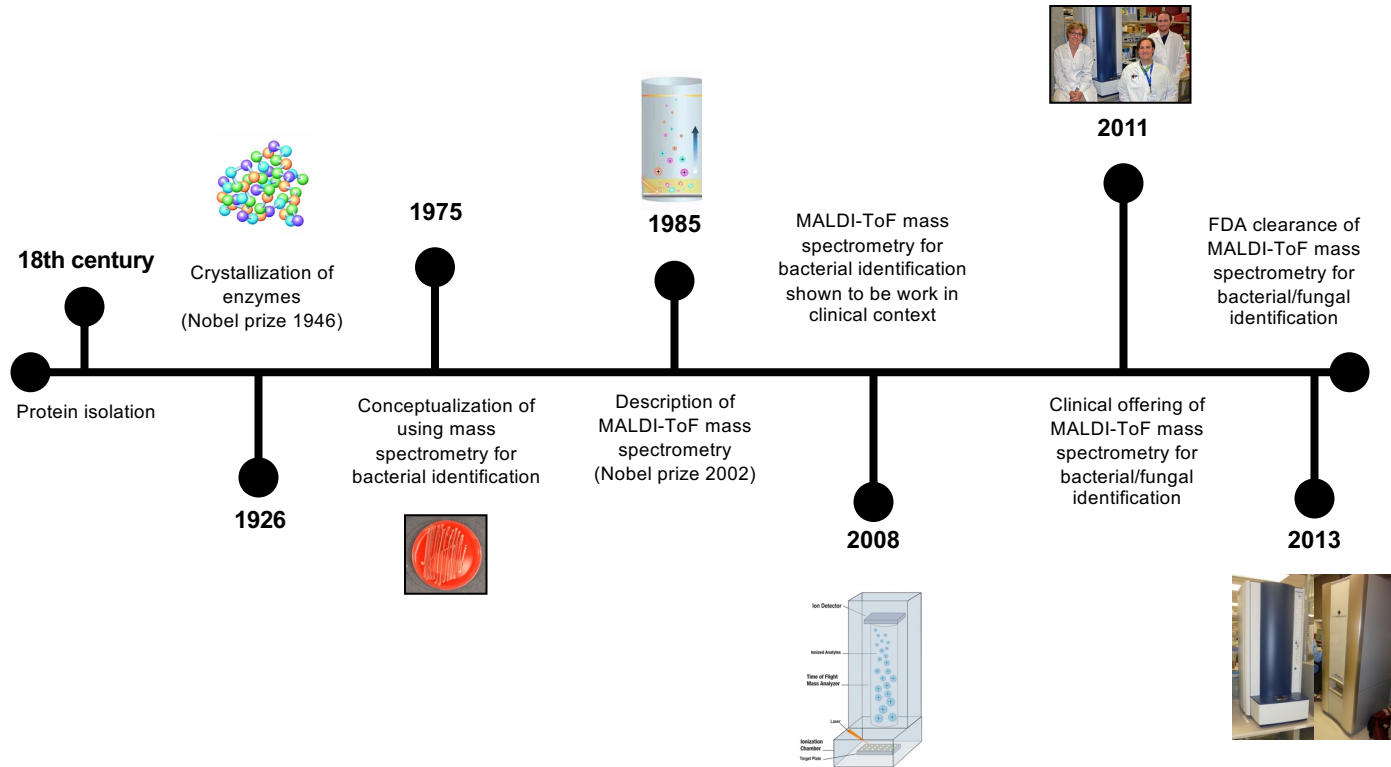


Rohr et al. PLoS One 2016;11:e0149856  
Zhi et al. PLoS One. 2013;8(11):e78962  
Riley Clin Microbiol Newslett 2017;39:69-73  
Lippi et al. Ann Transl Med 2017;5(4):82

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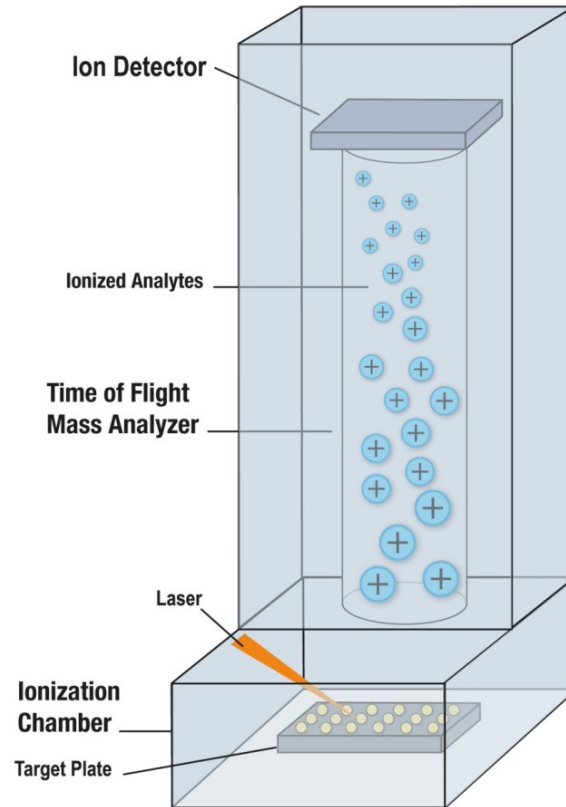
# Technology Revolution Proteomics





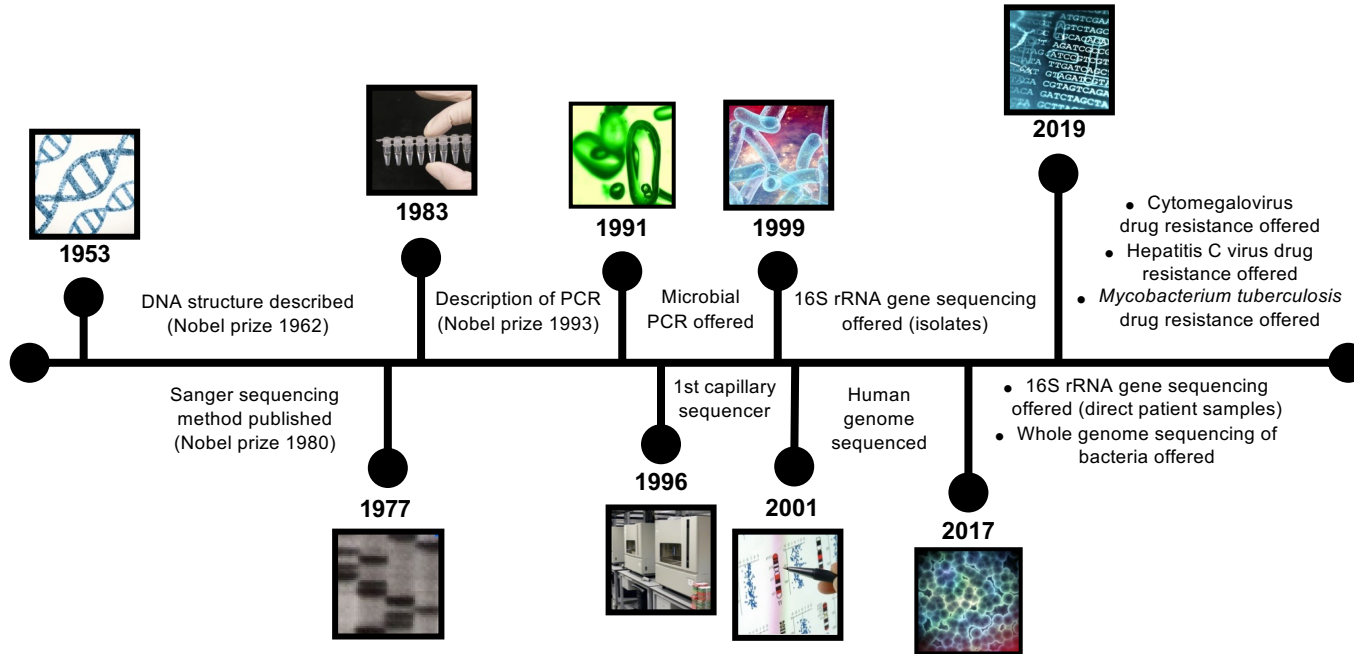
# Technology Revolution

## MALDI ToF Mass Spectrometry



# Technology Revolution

## NAAT and Sequencing-Based Diagnostics



# Technology Revolution: Microbial Real-Time PCR Assays Developed @ Mayo Clinic, 2000–2020

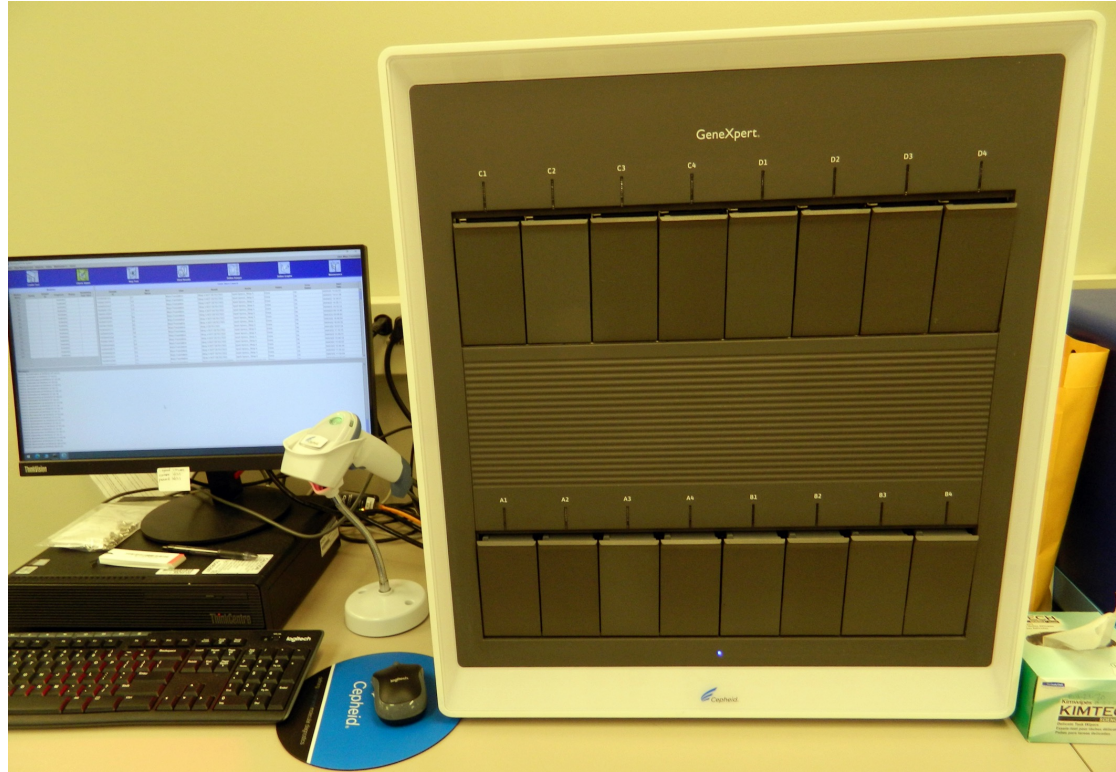
2000	Herpes simplex virus*
2001	<i>Bacillus anthracis</i>
	Cytomegalovirus*
	<i>Toxoplasma gondii</i> *
2002	<i>Bordetella pertussis/parapertussis</i> *
	<i>Streptococcus pyogenes</i>
	Enterovirus
	Varicella zoster virus*
2003	<i>Tropheryma whipplei</i> *
	<i>Babesia microti</i> *
	BK virus*
	<i>Ehrlichia/Anaplasma</i> species*
	Epstein–Barr virus (qualitative)*
	JC virus*
	<i>Borrelia burgdorferi</i> *
2004	<i>Bartonella</i> species*
	<i>vanA/vanB</i> *
	Parvovirus B19*
2005	Influenza A/B
	<i>Pneumocystis jirovecii</i>
	BK virus (quantitative)
	Epstein–Barr virus (quantitative)
	Human herpesvirus-6
	<i>Plasmodium</i> species
	West Nile virus
2007	<i>Clostridioides difficile</i>
	<i>Coccidioides immitis/posadasii</i>
	<i>Staphylococcus aureus</i>
	<i>Mycobacterium chelonae/abscessus</i>
2008	Adenovirus

2009	<i>Legionella</i> species
	<i>bla</i> <sup>KPC</sup>
	<i>Mycobacterium tuberculosis</i> complex
2010	<i>Salmonella</i> species
	<i>Shigella</i> species
	<i>Campylobacter</i> species
	<i>Yersinia</i> species
	Shiga toxin
2011	<i>Mycoplasma genitalium</i>
	<i>Mycoplasma hominis</i>
	<i>Ureaplasma urealyticum/parvum</i>
	<i>Histoplasma/Blastomyces</i> species
2012	<i>Mycobacterium tuberculosis</i> complex species
2013	<i>Coxiella burnetii</i>
	<i>bla</i> <sup>NDM</sup>
	<i>Mycoplasma pneumoniae</i>
2014	<i>Babesia</i> species
	<i>Borrelia mayonii</i>
2015	Microsporidia
2016	<i>Borrelia miyamotoi</i>
2017	<i>bla</i> <sup>OXA48-like</sup>
	<i>bla</i> <sup>VIM</sup>
	<i>Kingella kingae</i>
	<i>Acanthamoeba</i> species
	Free-living amoeba
	Norovirus
2020	<i>mecA</i>
	<i>Helicobacter pylori</i>
	<i>Candida auris</i>
	SARS coronavirus-2

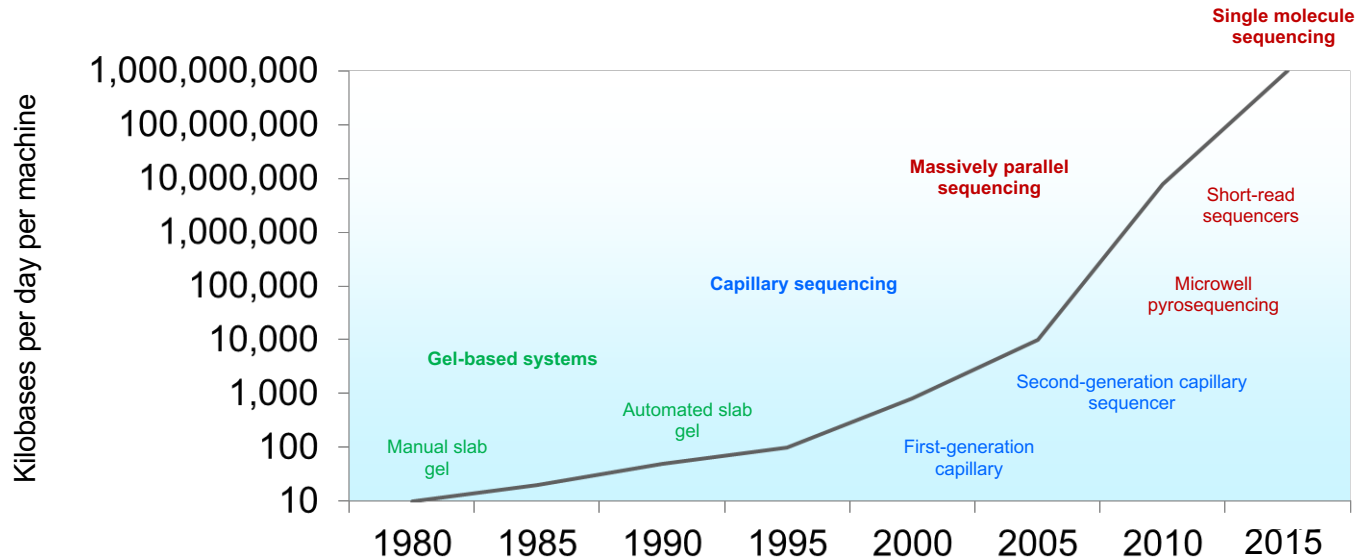
Microorganisms marked by an asterisk (\*) were detected with conventional PCR assays with Southern blot detection developed and deployed in the 1990s prior to conversion to real-time PCR assays

Patel. Clin Chem. 2022;68:10-15

# Technology Revolution: Multiplex and Rapid Automated NAATS

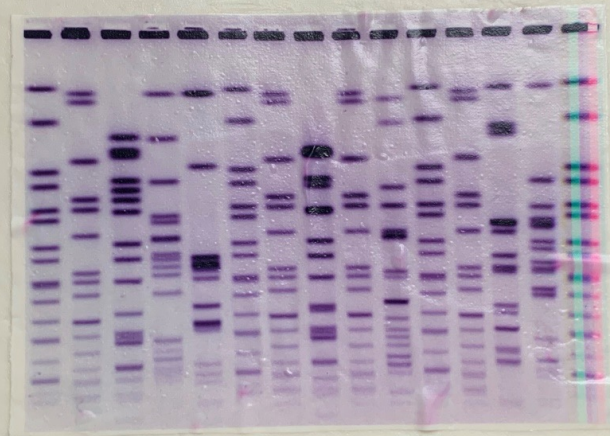


# Technology Revolution: Improvements in DNA Sequencing Rates



Based on Stratton et al. Nature 2009;458:719

Farewell PFGE!

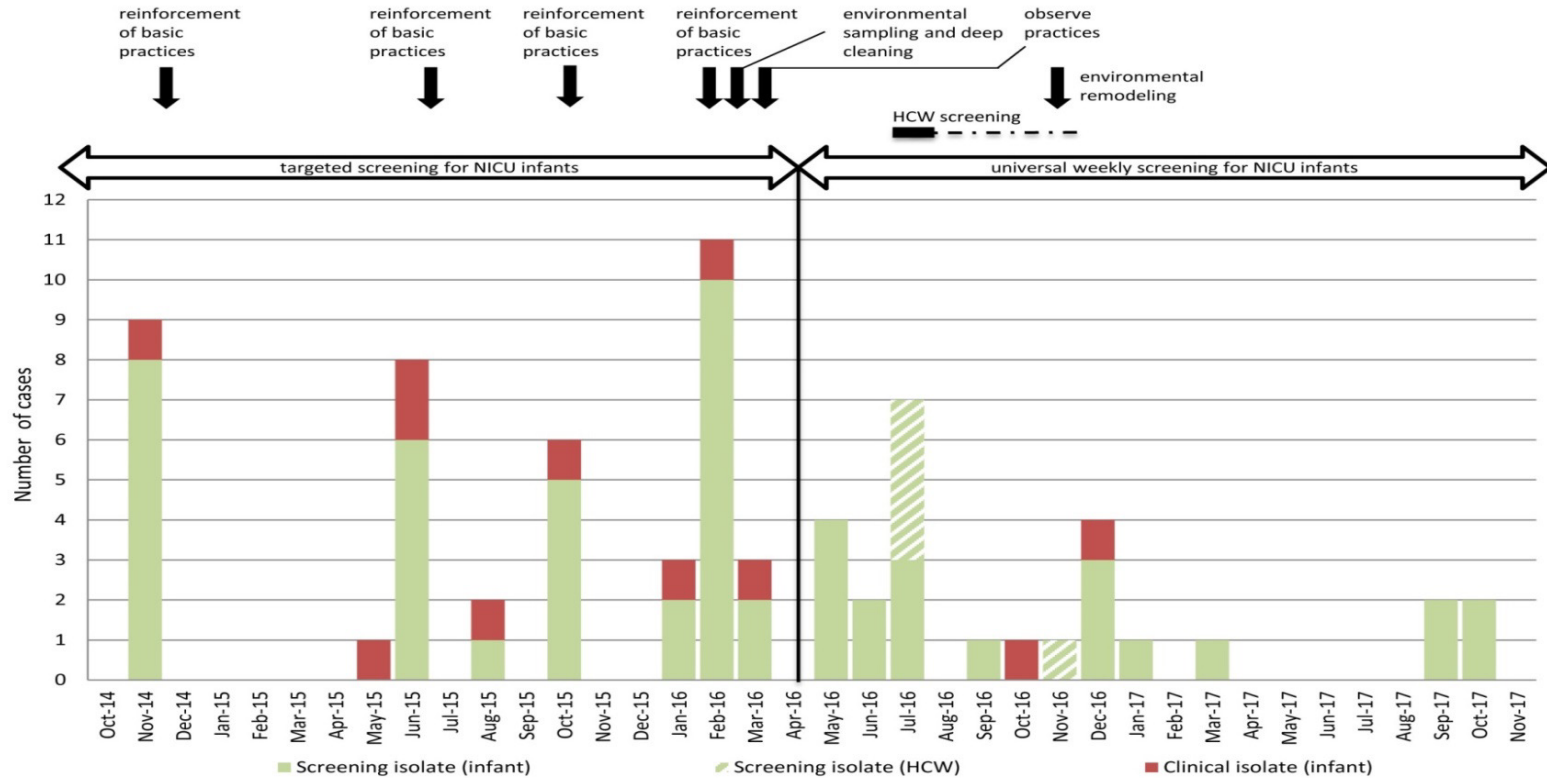


1982-2020



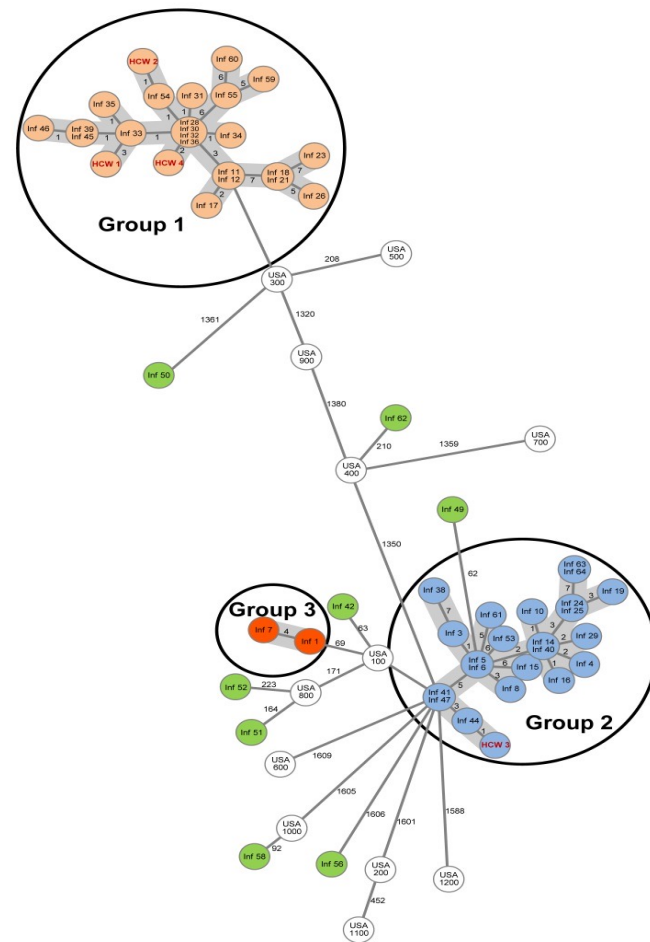
# Technology Revolution: Whole Genome Sequencing

## Neonatal Intensive Care Unit MRSA Cases by Initial Source & Interventions Taken



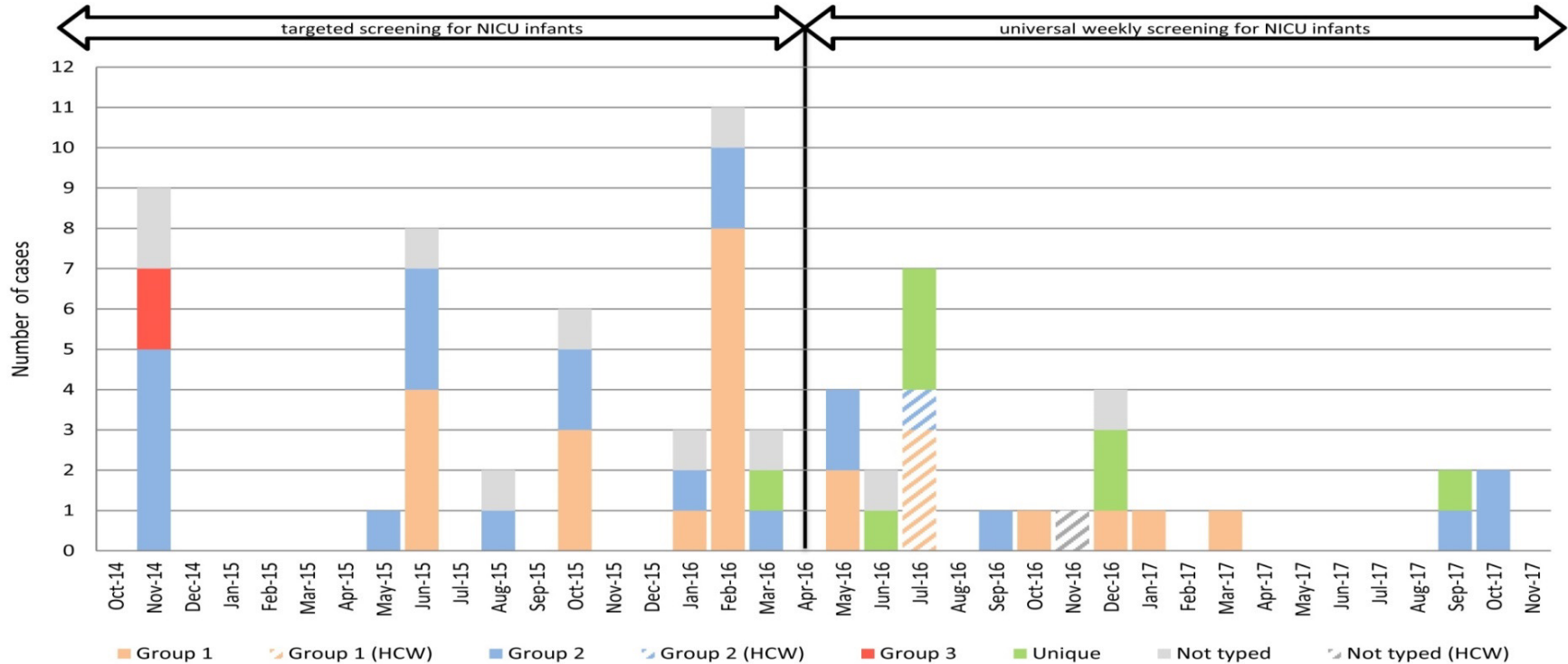


# Neonatal Intensive Care Unit Isolates (Clinical & Surveillance) Neonates & Healthcare Workers



# NICU Outbreaks (!) Timeline

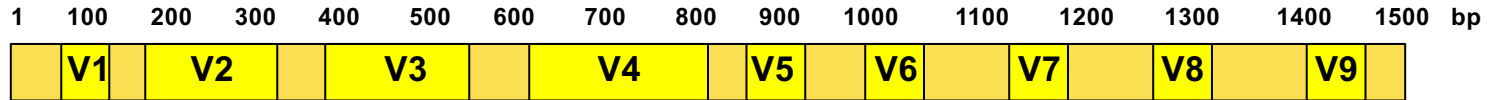
## MRSA Cases Shown by WGS Group



# Species Validated for WGS (cgMLST)

- *Acinetobacter baumannii*
- *Campylobacter jejuni/coli*
- *Clostridioides difficile*
- *Cutibacterium acnes*
- *Enterobacter cloacae*
- *Enterococcus faecalis*
- *Enterococcus faecium*
- *Escherichia coli*
- *Klebsiella pneumoniae*
- *Legionella pneumophila*
- *Pseudomonas aeruginosa*
- *Serratia marcescens*
- *Staphylococcus aureus*
- *Staphylococcus epidermidis*
- *Staphylococcus lugdunensis*
- *Streptococcus agalactiae*
- *Streptococcus pyogenes*

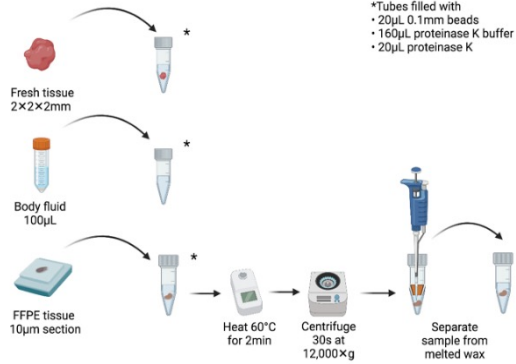
# Technology Revolution: Targeted Metagenomic Sequencing (16S Ribosomal RNA Gene)



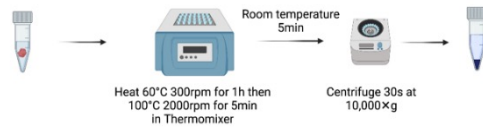
# 16S rRNA Gene PCR/Sanger Sequencing in Clinical Practice

- Retrospective analysis specimens clinically analyzed by 16S rRNA PCR/Sanger sequencing (04-2017→ 03-2019)
- 566 specimens, 460 patients
- 17% (97/566) 16S rRNA gene PCR/Sanger sequencing positive
  - 90% (88/97) positive results → clinical infections
- 22% positive specimens (21/97), 16S rRNA gene detected but bacterium not identified by Sanger sequencing (mixtures)

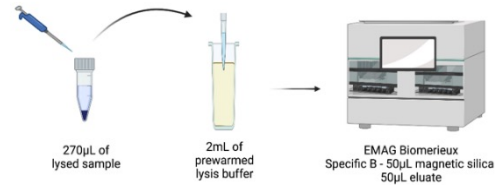
### 1 Sample processing



### 2 Digestion and lysis



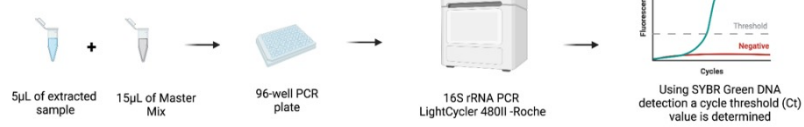
### 3 Extraction



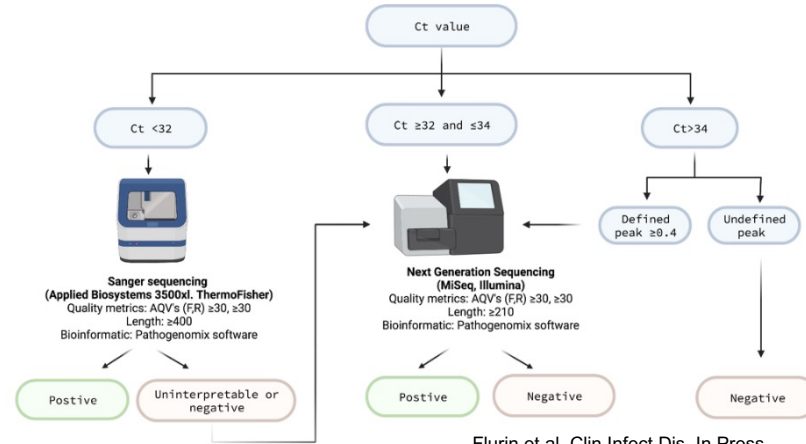
### 4a 16S rRNA gene and primers



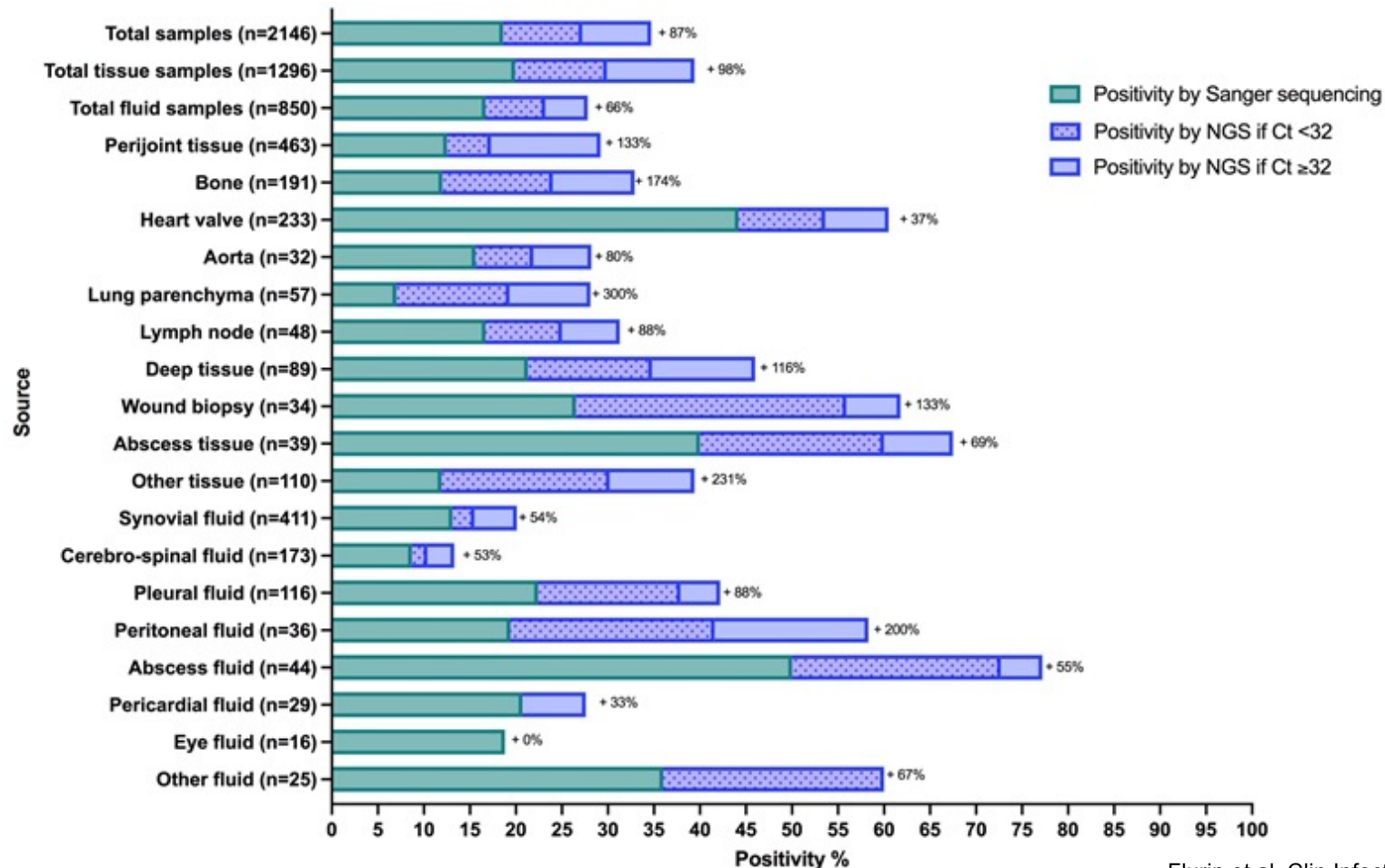
### 4b 16S rRNA PCR



### 5 Sequencing algorithm



## Yield of next-generation sequencing (NGS)







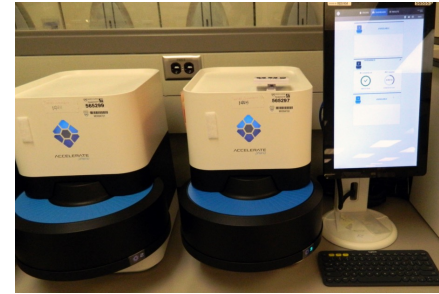
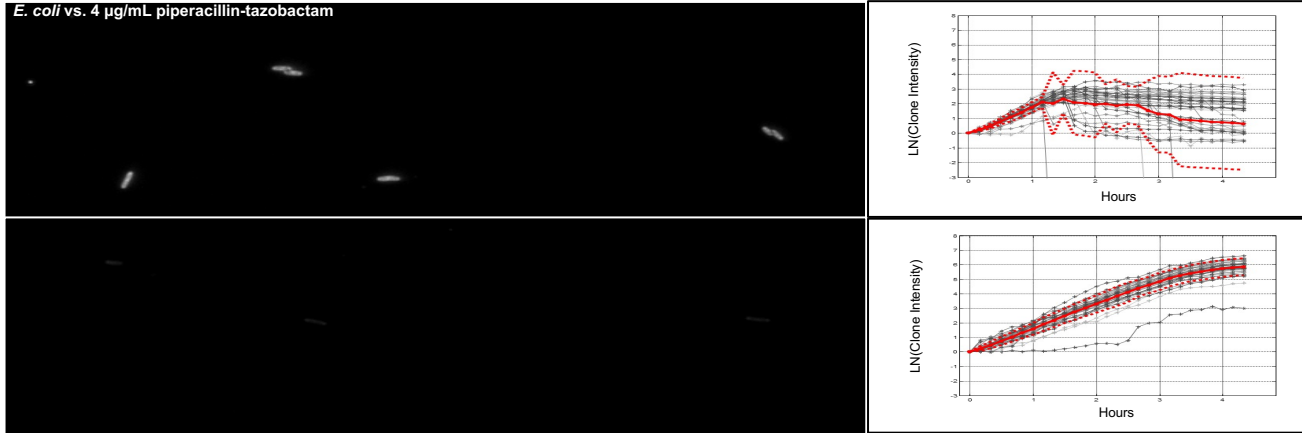


# Technology Revolution: Single Cell Imaging

## Application: Antibacterial Susceptibility Testing

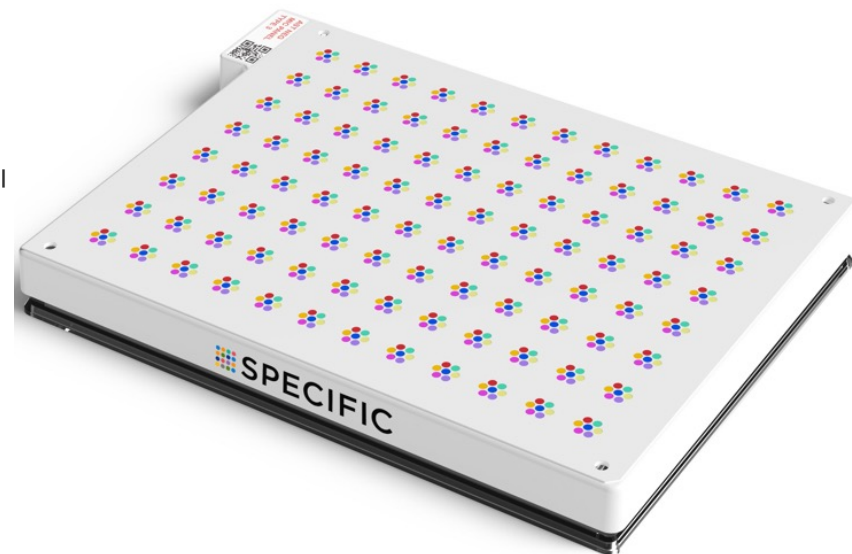
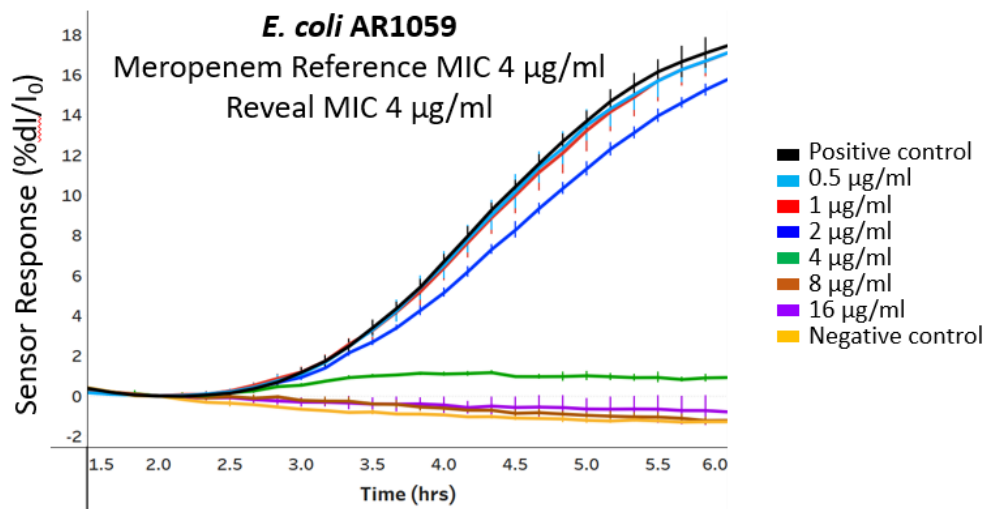
- Time-lapse darkfield imaging, analysis of bacterial growth
  - Individual bacterial response to single antibiotic concentration over time

*E. coli* vs. 4 µg/mL piperacillin-tazobactam



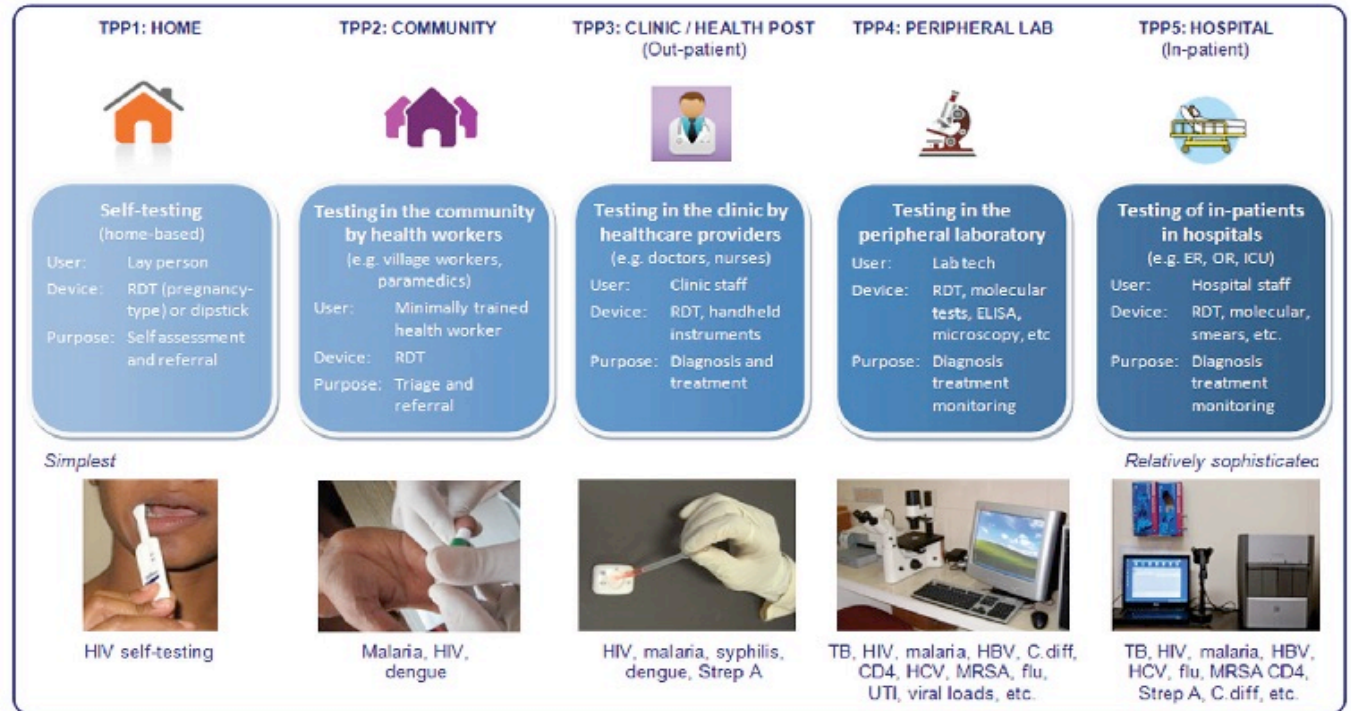
# Technology Revolution: Metabolomics

## Application: Antibacterial Susceptibility Testing



# Technology Revolution

## Point of Care Diagnostics



[www.asm.org/index.php/colloquium-reports/item/6421-changing-diagnostic-paradigms-for-microbiology](http://www.asm.org/index.php/colloquium-reports/item/6421-changing-diagnostic-paradigms-for-microbiology)

# Technology Revolution: Advanced Host Response Assessment for Infectious Diseases

- **Advanced human transcriptomic and/or proteomic analyses may be able to**
  - Point to underlying viral, bacterial, fungal, parasitic infections and non-infectious processes
  - Possibly, point to specific microbial etiologies
  - Blood, body fluids, tissues?

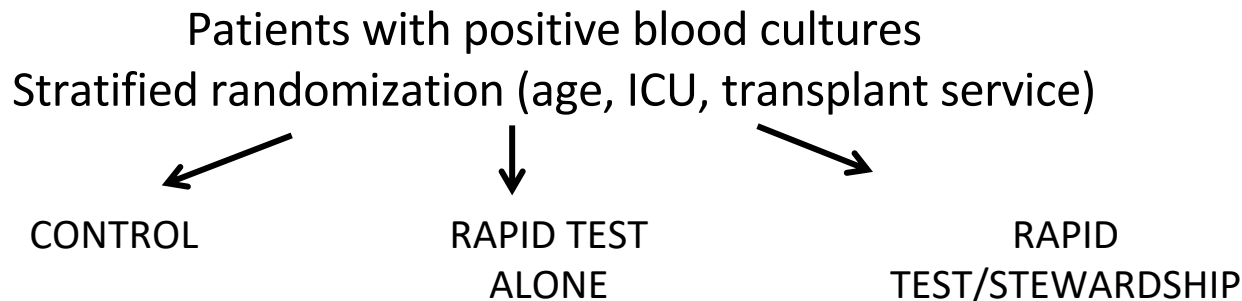
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# How to Move to the Future?

- **MALDI-TOF mass spectrometry example**
- **Demonstration of clinical utility**
  - COVID-19 example
  - Advanced diagnostics - sometimes not so clear...

# Randomized Controlled Clinical Trial BioFire BCID, Mayo Clinic 2013-2014



- Gram stain called to service
- Standard subculture and susceptibility (1-3 d)

- Gram stain called to service
- Standard subculture and susceptibility (1-3 d)
- Rapid test plus lab call with comments (1 h)

- Gram stain called to service
- Standard subculture and susceptibility (1-3 d)
- Rapid test plus lab call with comments (1 h)
- ID MD/pharmacist call with specific treatment recommendations

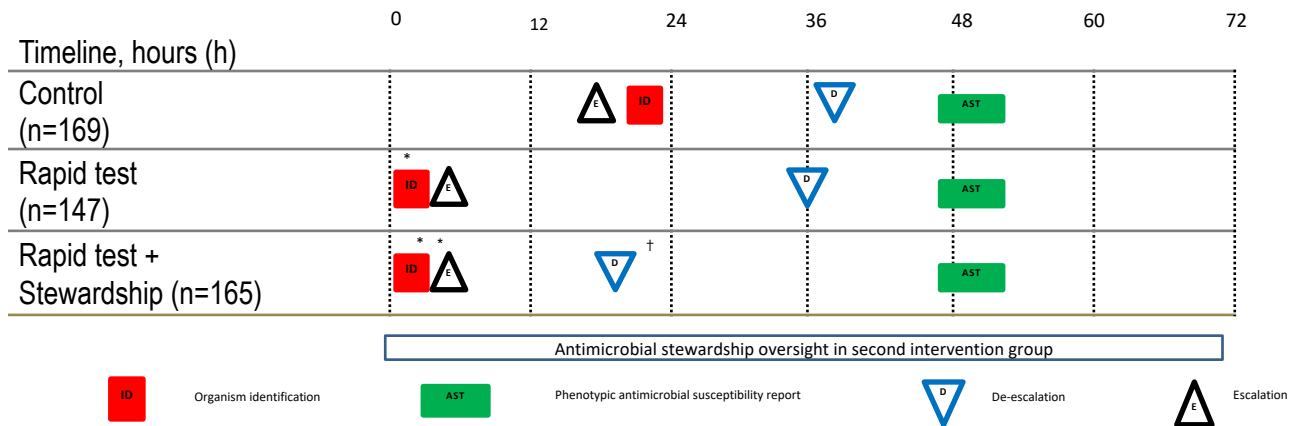
Banerjee et al. Clin Infect Dis 2015;61:1071

Supported by the National Institute of Allergy And Infectious Diseases of the National Institutes of Health under Award Number UM1AI104681 (Antibacterial Resistance Leadership Group)





# Comparison of Time To Identification, Susceptibility Results, and Antibiotic Modifications



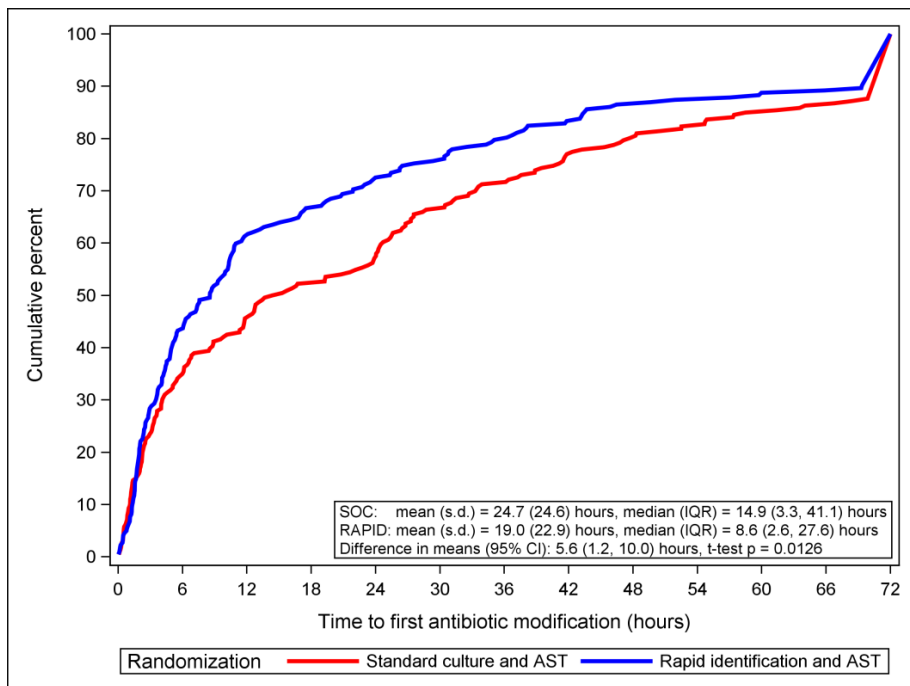
\*Significant vs. control; †Significant vs. control and rapid multiplex PCR alone

# Thoughtful, Clear and Informative Test Reporting

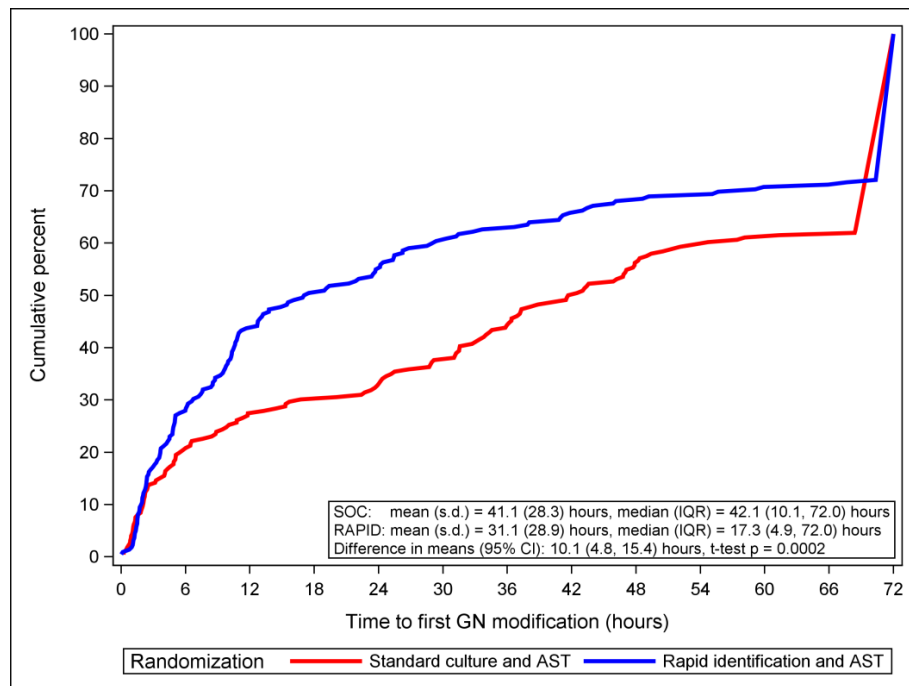
FilmArray BCID2 Result	Report	Isolate Comments
<i>Staphylococcus aureus</i> + <i>mecA/C</i> and MREJ (MRSA) +	<i>Staphylococcus aureus</i> complex	<i>mecA</i> detected Methicillin (oxacillin)-resistant <i>Staphylococcus aureus</i> complex (MRSA). MRSA is predictably resistant to beta-lactam antibiotics (except ceftaroline). Vancomycin or other anti-MRSA treatment recommended for initial therapy pending susceptibility results. Semi-Urgent Result Consult Infectious Diseases
<i>Staphylococcus aureus</i> + <i>mecA/C</i> and MREJ (MRSA) -	<i>Staphylococcus aureus</i> complex	<i>mecA</i> not detected Methicillin (oxacillin)-susceptible <i>Staphylococcus aureus</i> complex. An anti-staphylococcal beta-lactam, such as cefazolin, nafcillin or oxacillin, is recommended for empiric therapy, unless clinically contraindicated. Consult Infectious Diseases
<i>Staphylococcus epidermidis</i> + <i>mecA/C</i> +	<i>Staphylococcus epidermidis</i>	<i>mecA</i> detected Methicillin (oxacillin)-resistant <i>Staphylococcus epidermidis</i> . Possible blood culture contaminant (unless isolated from more than one blood culture draw or clinical case suggests pathogenicity). No antibiotic treatment is indicated for blood culture contaminants.

# Randomized Trial Evaluating Clinical Impact of RAPid Identification and Susceptibility Testing for Gram-negative Bacteremia: RAPIDS-GN

## Time to any antibiotic change

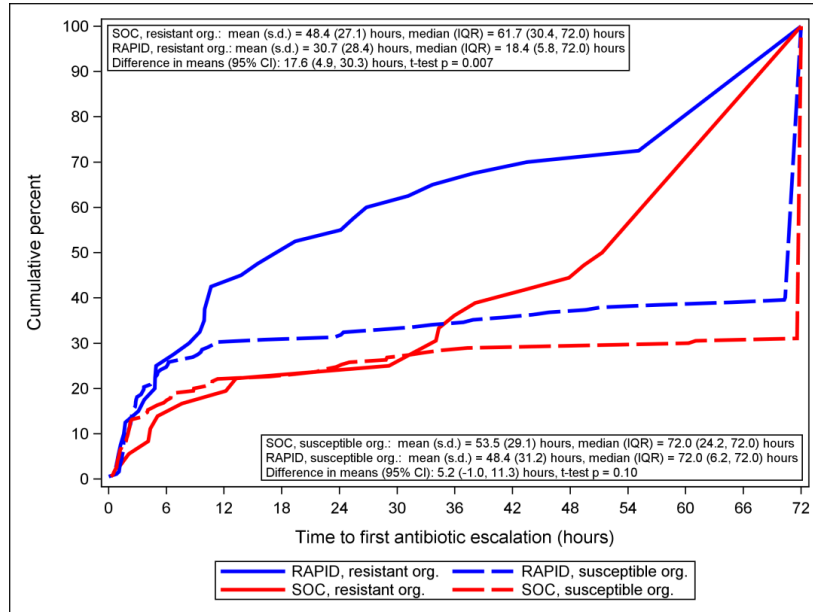


## Time to GN antibiotic change

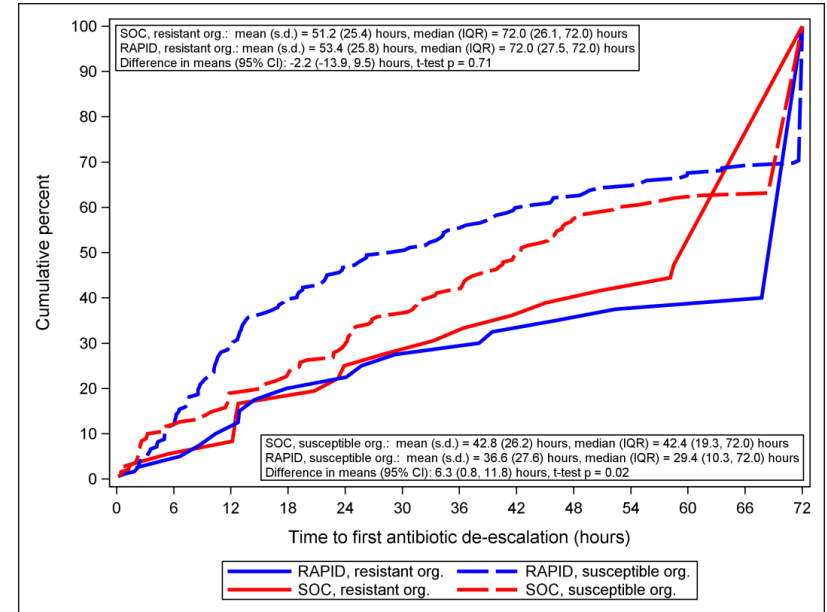


# Randomized Trial Evaluating Clinical Impact of RAPid Identification and Susceptibility Testing for Gram-negative Bacteremia: RAPIDS-GN

## Time to antibiotic escalation

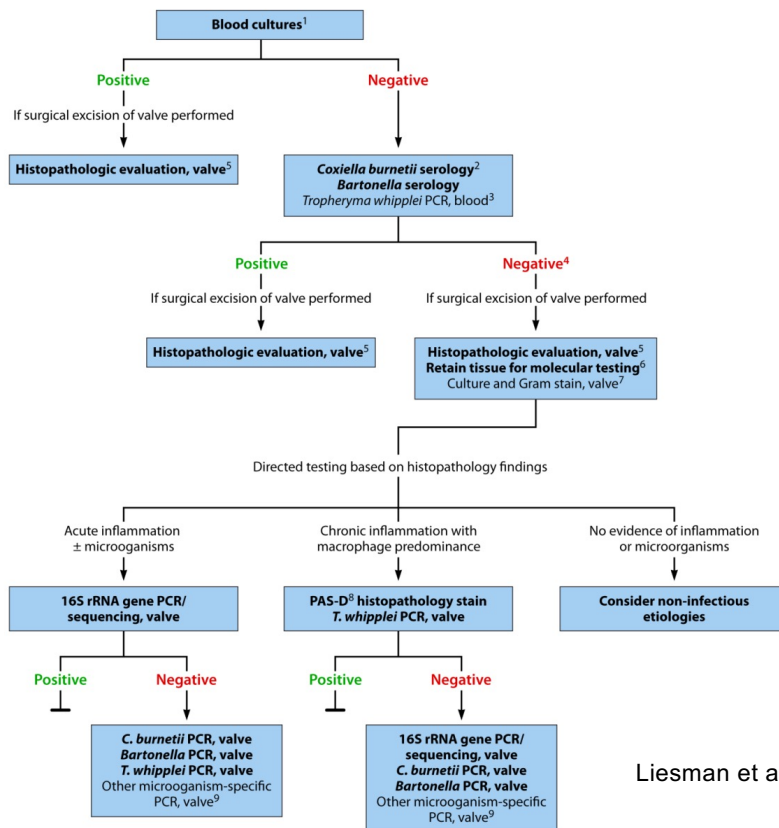


## Time to antibiotic de-escalation



# Importance of Guidelines, Diagnostic Algorithms, Diagnostic Stewardship

## Microbiologic and Pathological Diagnostic Algorithm For Endocarditis



Liesman et al J Clin Microbiol. 201;55(9):2599-608

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# The Deliverables: Improved Patient Health

- **Improved patient health and hopefully...**
  - Avoid unneeded testing and treatment (patient benefit)
  - Decrease transmission of infectious diseases (societal benefit)
  - Curb emergence of antimicrobial resistance (patient and societal benefit)

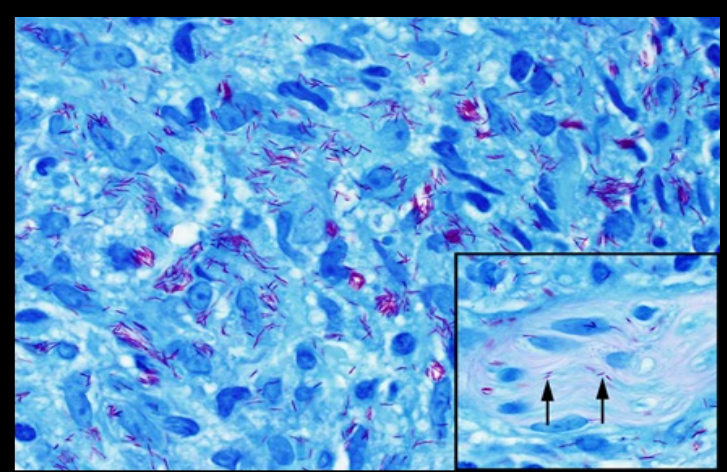
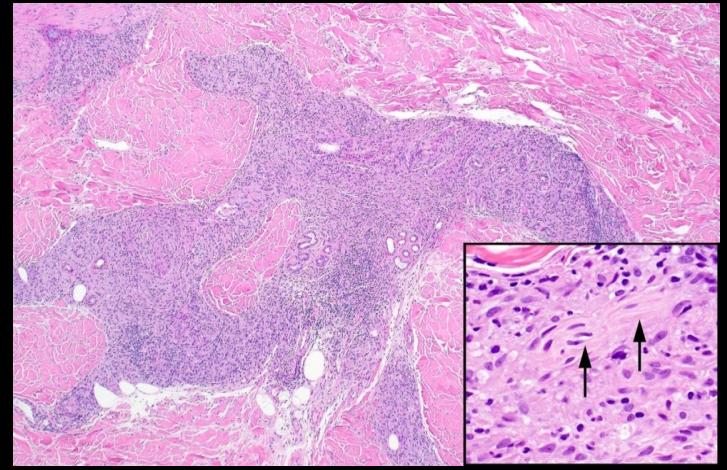
# The Deliverables: Changes in Healthcare Delivery

- **Home-based testing**
- **Patient self-testing**
- **Testing @ non-traditional sites**
- **Test prioritization approaches in traditional healthcare sites (e.g., emergency departments, urgent care settings)**
- **etc.**

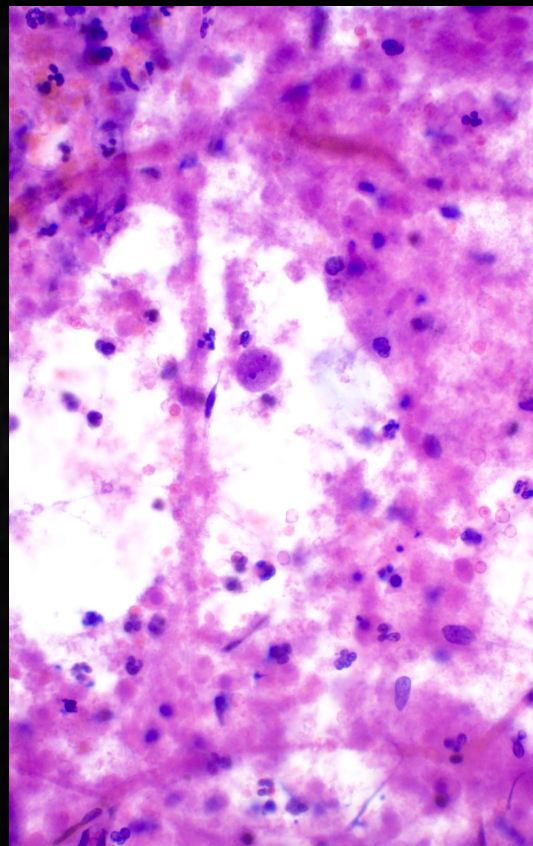
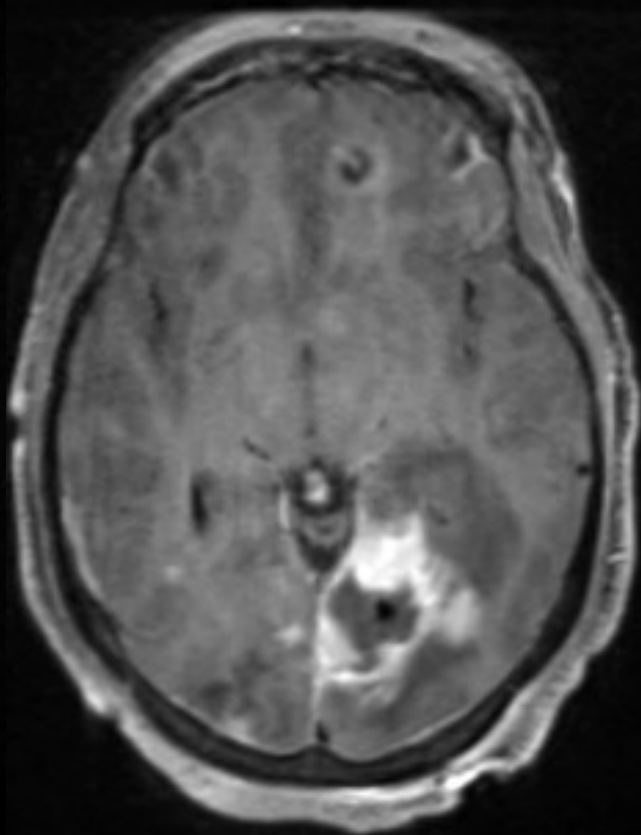
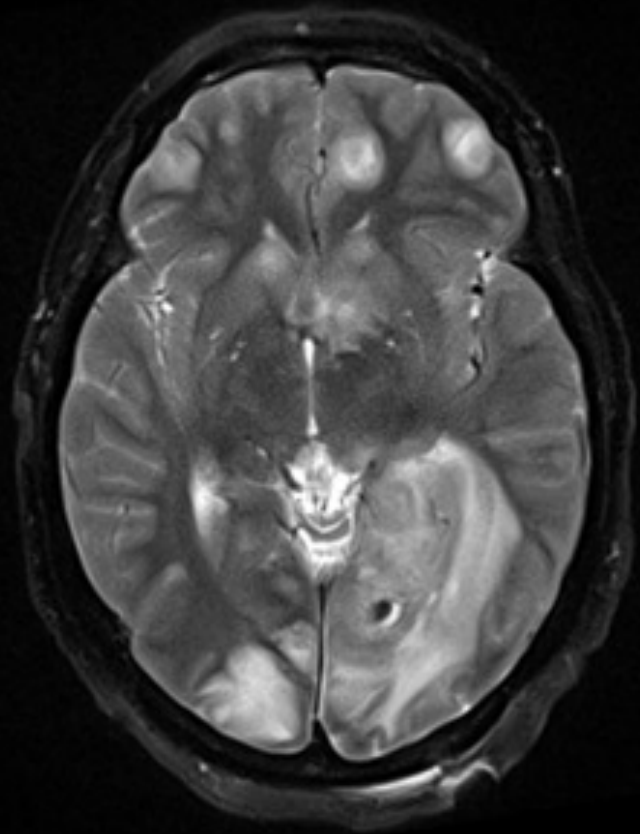


- **Discovery through testing**

- *Borrelia mayonii*, *E. muris* subspecies *eauclairensis*, *Yersinia rochesterensis*
  - Many more novel species remain to be described
- Understanding antimicrobial resistance
  - Characterization of vancomycin resistance in *Paenibacillus popilliae*
  - Recognition of the frequency of resistance in *Helicobacter pylori*
- New diseases – hyperammonemia associated with *Ureaplasma* species



Source:	BRAIN, RIGHT- Tissue. Rlght frontal lobe		
Fresh vs FFPE:	Fresh		
Cp value:	31.86		
FORWARD			
Abundance	Read length	Result	% ID
	22606	222Amoebophilus	96.8
	405	225Cutibacterium acnes **	100
	274	224Methylothera	95.5
	222	222Amoebophilus	96.4
		Chryseobacterium lactis / Chryseobacterium aquaticum / Chryseobacterium aurantiacum /	
	211	223Chryseobacterium vietnamense	99.1
	148	224Amoebophilus	96.4
	140	222Enterococcus faecium / Enterococcus hirae / Enterococcus pseudoavium	100
	138	222Amoebophilus	96.8
	128	222Amoebophilus	96
	117	222Amoebophilus	96.4
	106	222Amoebophilus	96.4
REVERSE			
	19971	228Amoebophilus	98.2
	754	229Amoebophilus	98.3
	353	226Cutibacterium acnes / Cutibacterium namnetense *	100
	179	227Methylophilus	98.7
	170	228Amoebophilus	97.8
	144	228Amoebophilus	97.8
		Enterococcus durans / Enterococcus faecium / Enterococcus hirae / Enterococcus ratti /	
		Enterococcus thailandicus / Enterococcus villorum / Enterococcus xinjiangensis / Enterococcus	
	139	227lactis / Enterococcus pseudoavium	100
		Chryseobacterium artocarpi / Chryseobacterium antibioticum / Chryseobacterium carnipullorum	
	131	228/ Chryseobacterium shigense / Chryseobacterium ureilyticum / Chryseobacterium vrystaatense	99.6
	118	228Amoebophilus	97.4
	118	228Amoebophilus	97.4
	104	228Amoebophilus	97.8



# Conclusions

- **Many current microbiology diagnostics antiquated, poorly used**
- **Technology revolution**
  - Bringing major advances
- **How to move to the future**
  - Demonstration of value
- **The deliverables**
  - Improved health (patients, society, antimicrobial resistance)
  - Changes to healthcare delivery
  - Better understanding of infectious diseases and their mimics (discovery through testing)



R01 AR056647  
R01 AI91594  
R21 AI125870  
UM1 AI104681