



Clinical Microbiology in *action*

Cases from the Brigham & Women's Hospital

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Topics within this talk

- The vital role of the technologist in patient care
- The integration of the clinical microbiology laboratory with multiple levels of healthcare decision makers
- The challenges of decision-making in the face of ever-increasing diagnostic complexity

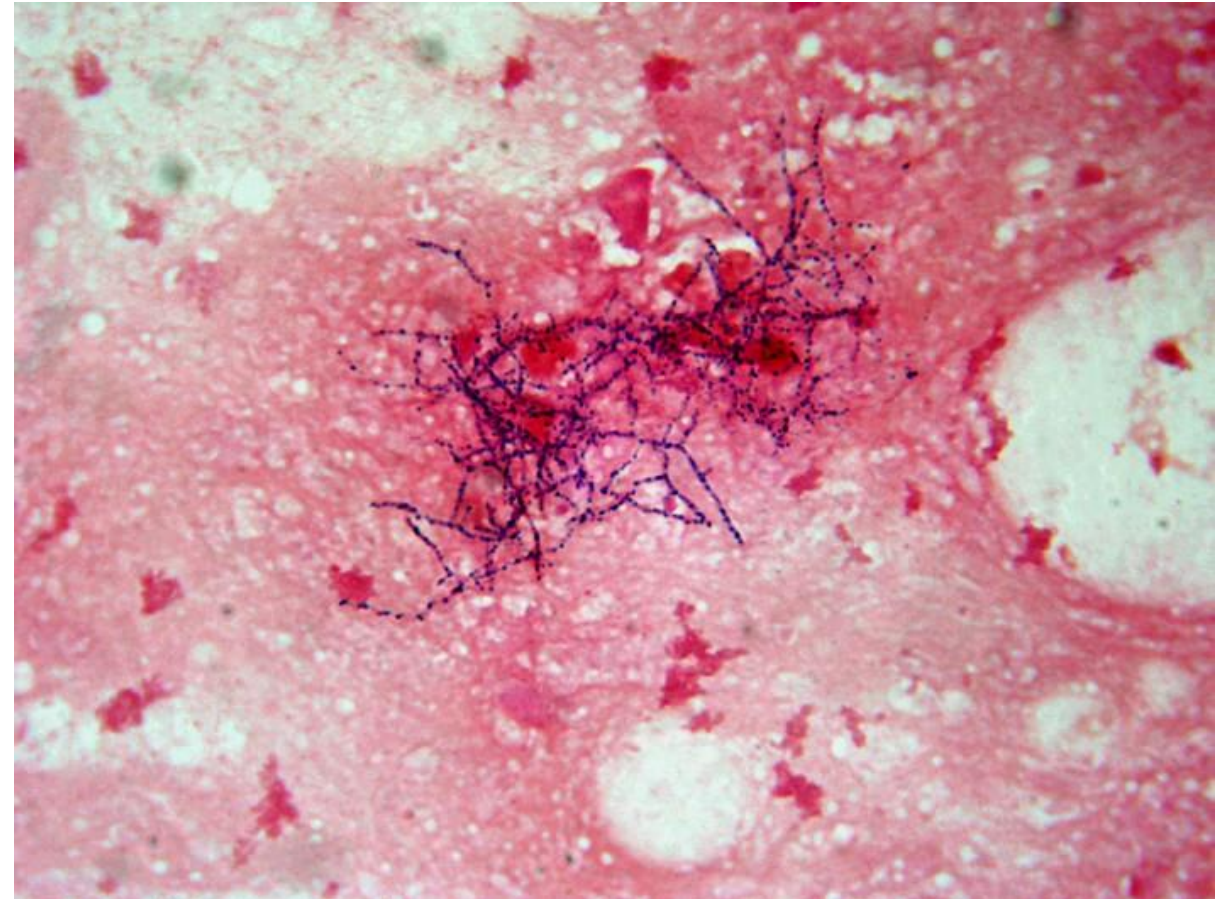


A case of the coughs

- A 30 year old healthy female presents to her pulmonologist in June of 2016 with a history of bronchiectasis and recurrent pneumonia
- She had 3 episodes of pneumonia between January and April 2016 during which she coughed up sputum that started out as yellowish-green but eventually became tinged with blood
- Each episode seemed to respond to antibiotics

A case of the coughs

- A sputum culture is obtained at the time and is negative for mycobacteria, fungi and bacteria
- Gram stain shows...





A case of the coughs

- Because she appears well, no treatment is initiated at the time
- The following month (August 2016) has another episode of pneumonia for which she is treated with cefuroxime
- In November 2017 she delivers a healthy full term baby with no complications
- In March of 2018 she again presents to her pulmonologist with 'rust-colored' sputum, fever to 101F, and night sweats
- She receives another course of cefuroxime but her symptoms persist, therefore she is switched to a longer course of levofloxacin



A case of the coughs

- Another sputum is obtained

Specimen Source/ Description

SPECIAL REQUESTS

GRAM STAIN

GRAM STAIN

GRAM STAIN

CULTURE / TEST

SPUTUM

None

4+ POLYS

1+ EPITHELIAL CELLS

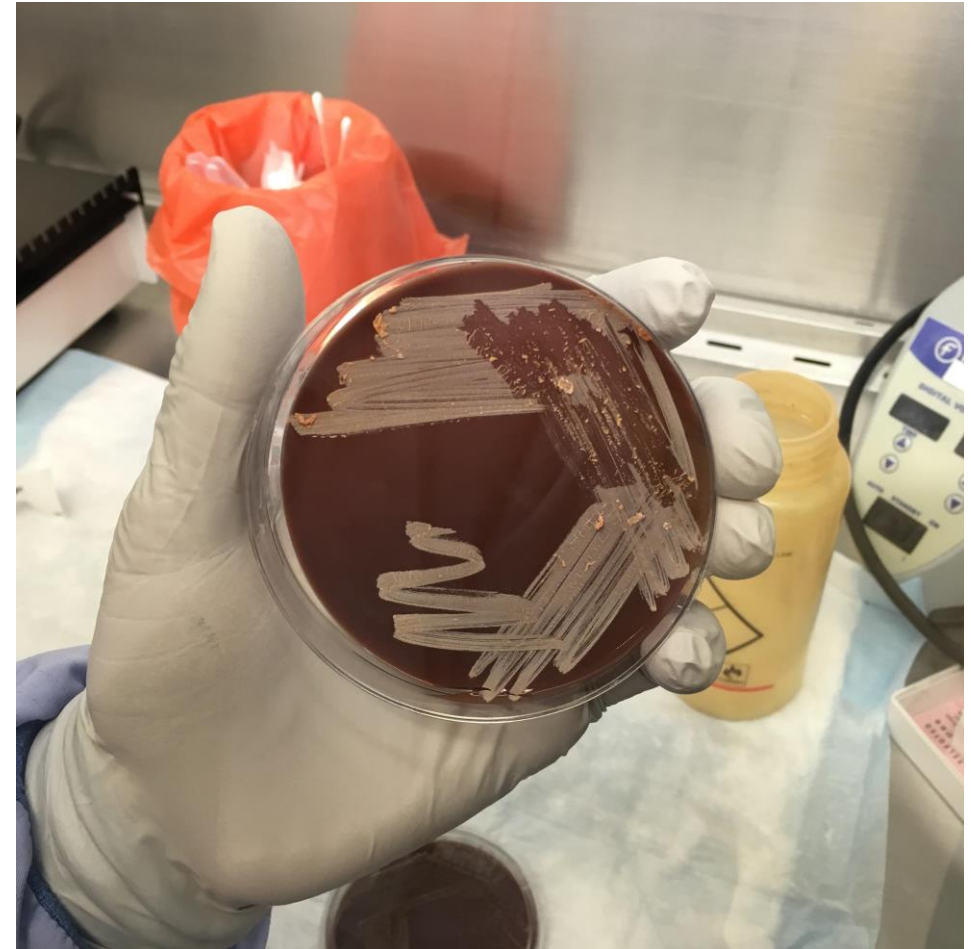
3+ BRANCHING GRAM POSITIVE RODS

2+ ORAL FLORA

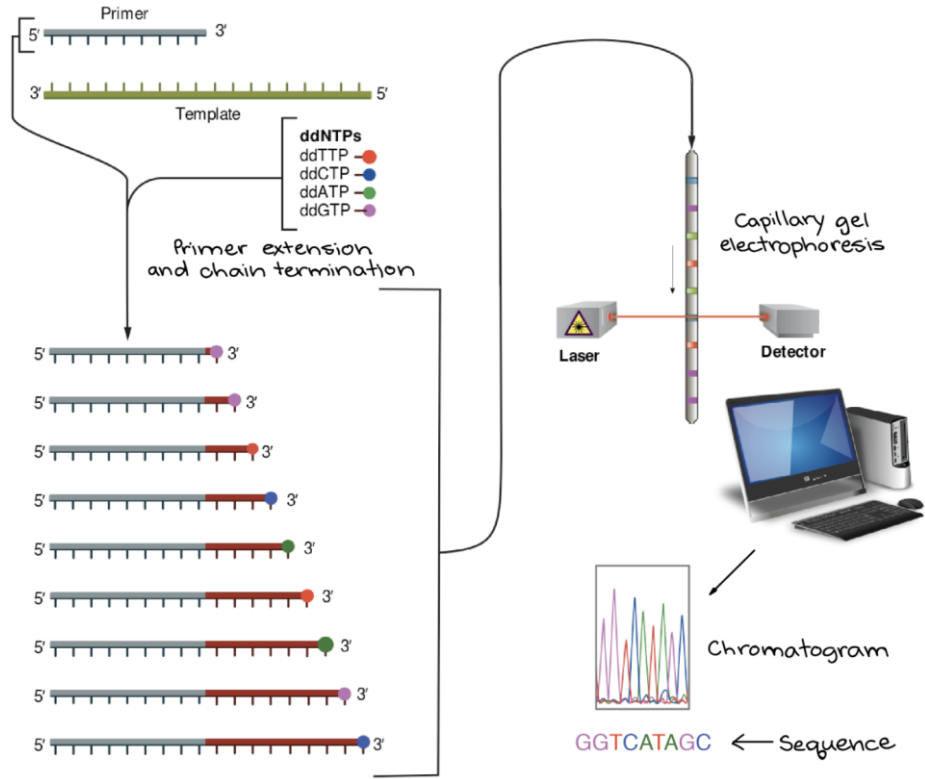
- What's going on here?

One in a million

- An astute technologist picked out a single small colony in the 2nd quadrant that appeared different than the surrounding oral flora
- The isolate failed to grow on Lowenstein-Jensen agar but did grow on chocolate and BCYE
- Gram stain matched sputum from 2016 and from 2018



When in doubt, sequence

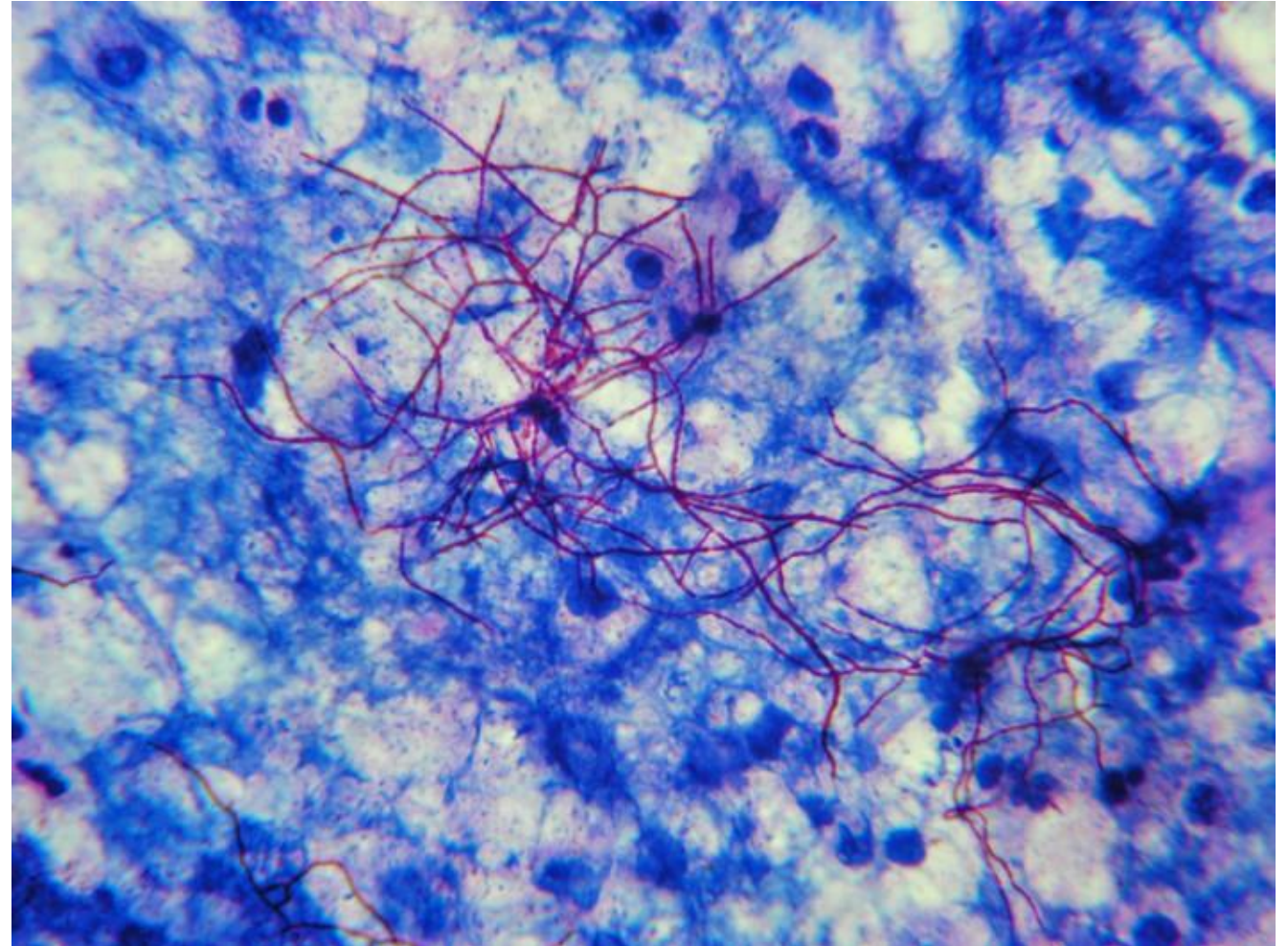


- 16S ribosomal gene sequencing was performed using a colony of pure growth
- ‘Broad-based’ bacterial primers successfully amplified the gene target
- Subsequent Sanger sequencing yielded a contig of ~1500 base pairs
- Using local sequence alignment software and a curated database of bacterial genomes, the genetic sequence had a 99.8% match to

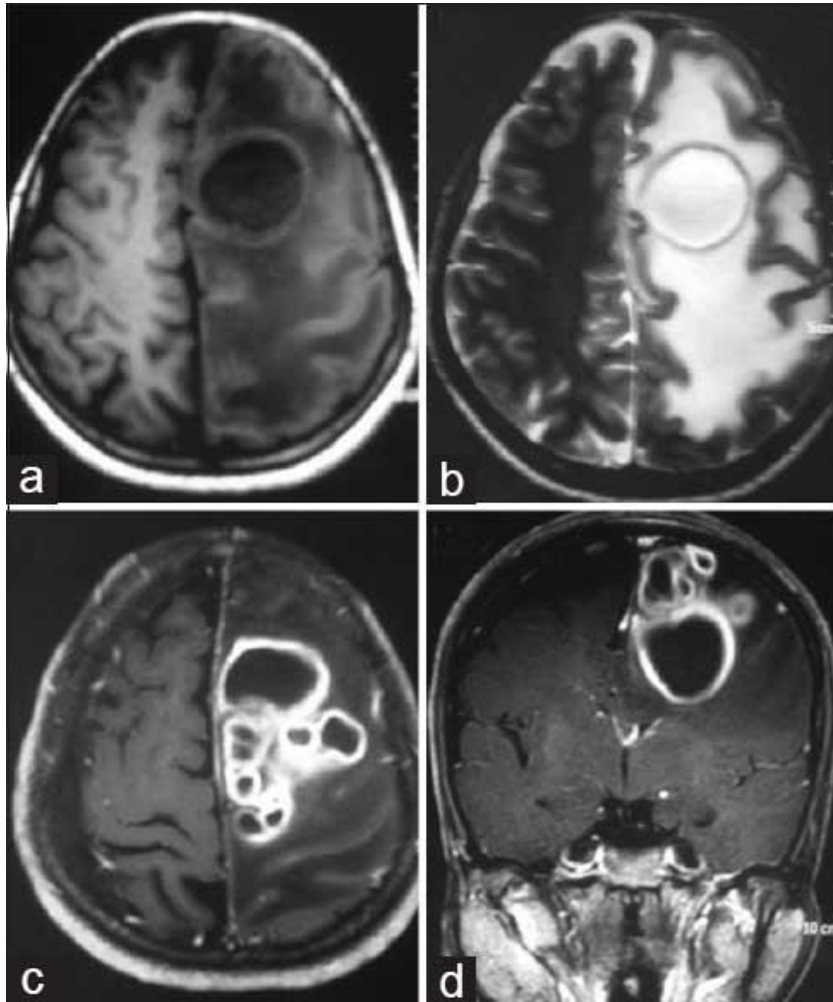
Nocardia asiatica

Nocardia : Everywhere you want to be

- Gram positive
- Filamentous
- Branching
- Modified acid-fast
- Soil saprophytes



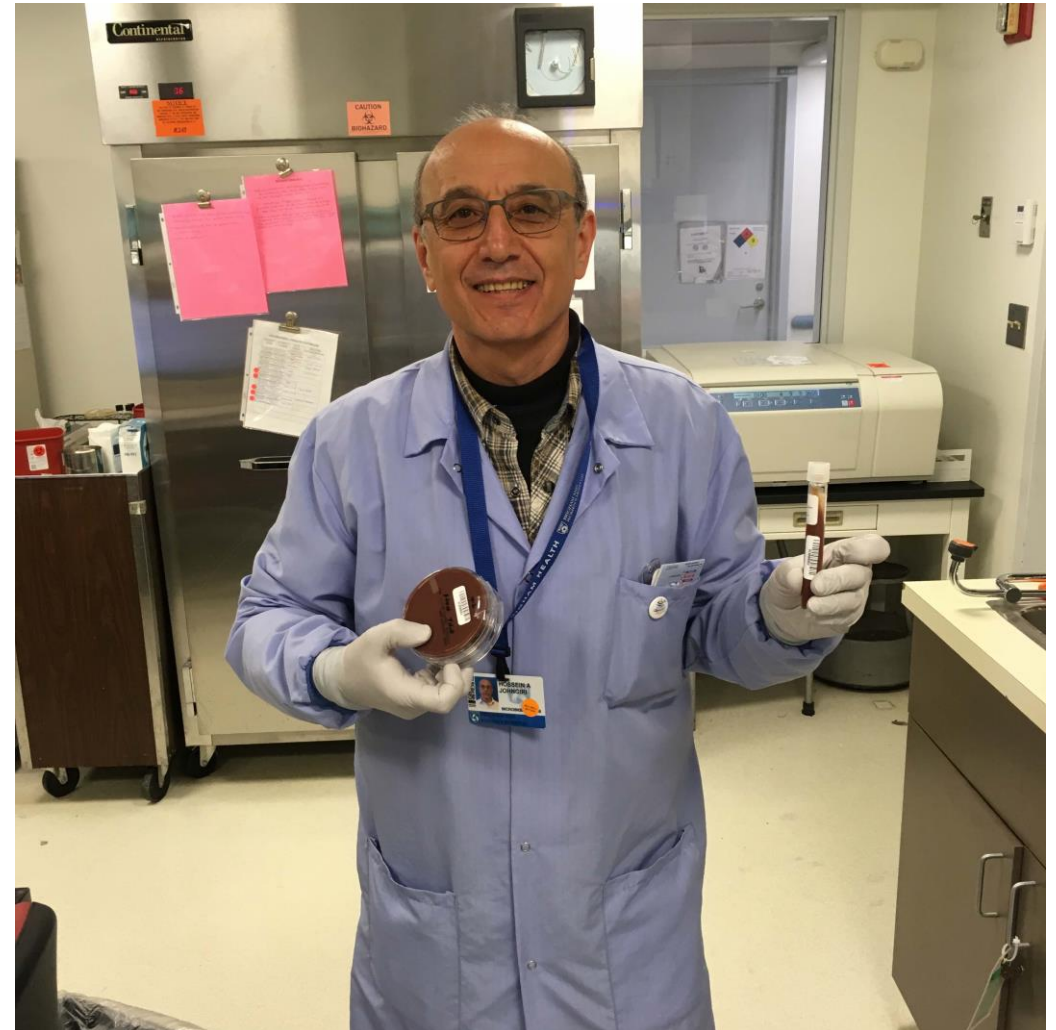
Nocardia : Everywhere you want to be



- Can cause disease in healthy and immunocompromised hosts
- 3 major clinical manifestations:
 - Cutaneous
 - Pulmonary
 - Disseminated (CNS in particular)

Micro saves the day (again)

- The patient was switched off levofloxacin and started on trimethoprim-sulfamethoxazole
- Her symptoms appear to have resolved and she is awaiting infectious diseases consultation for further management





Resistance is not futile

- 50 year old female with a history of uterine fibroids who presented to her general practitioner in India in August of 2017 with a complaint of R leg swelling for 2 weeks
- An ultrasound showed a 16x13cm mass attached to her uterus that was compressing her ureters
- She underwent a surgery to remove the tumor, which was identified as a leiomyosarcoma, and ureteral stents placed to maintain patency of her urinary tract
- She spent several days in the surgical intensive care unit in India



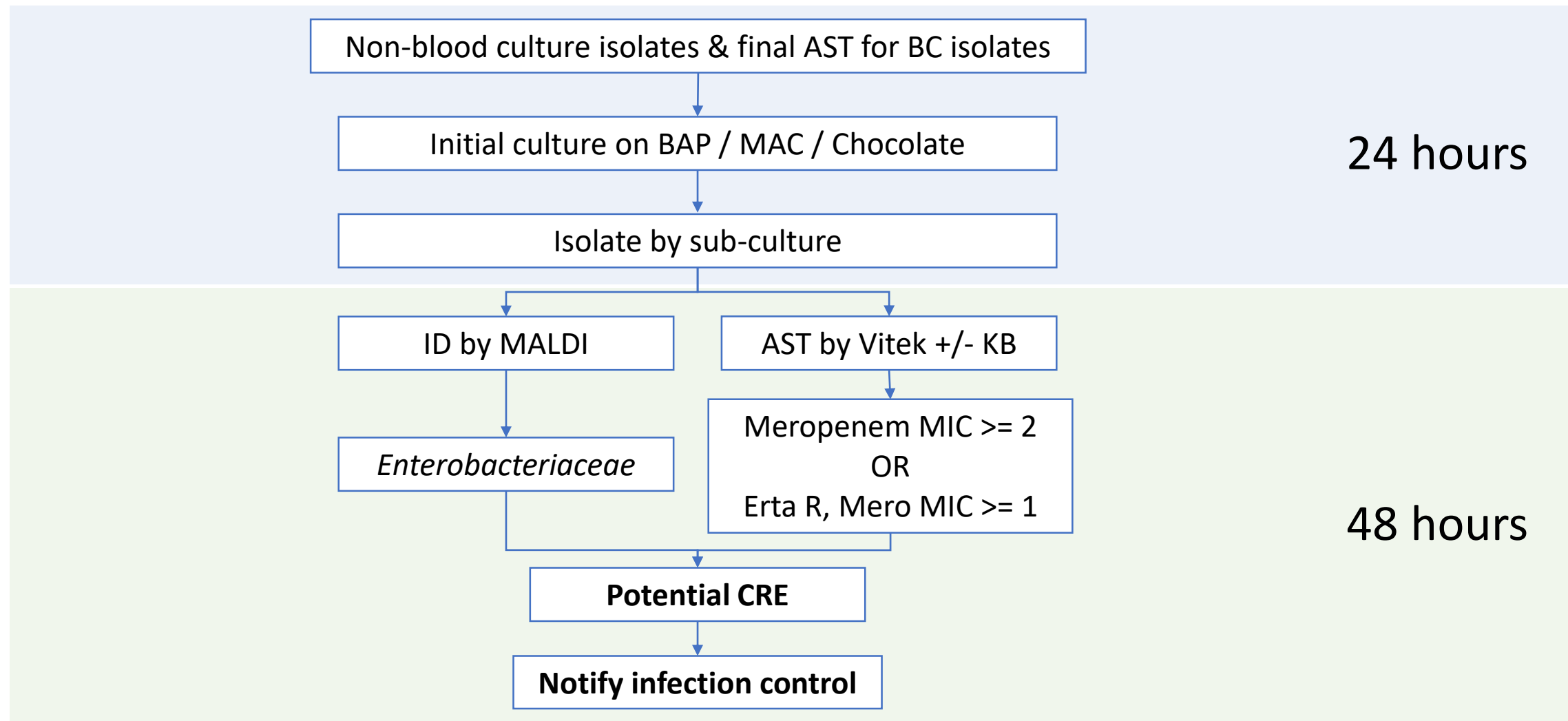
Resistance is not futile

- She comes to the Dana Farber Cancer Institute for further management of her cancer
- She is asymptomatic
- On routine pre-chemotherapy surveillance, she is noted on her urinalysis to have 50 WBCs
- A urine culture is obtained

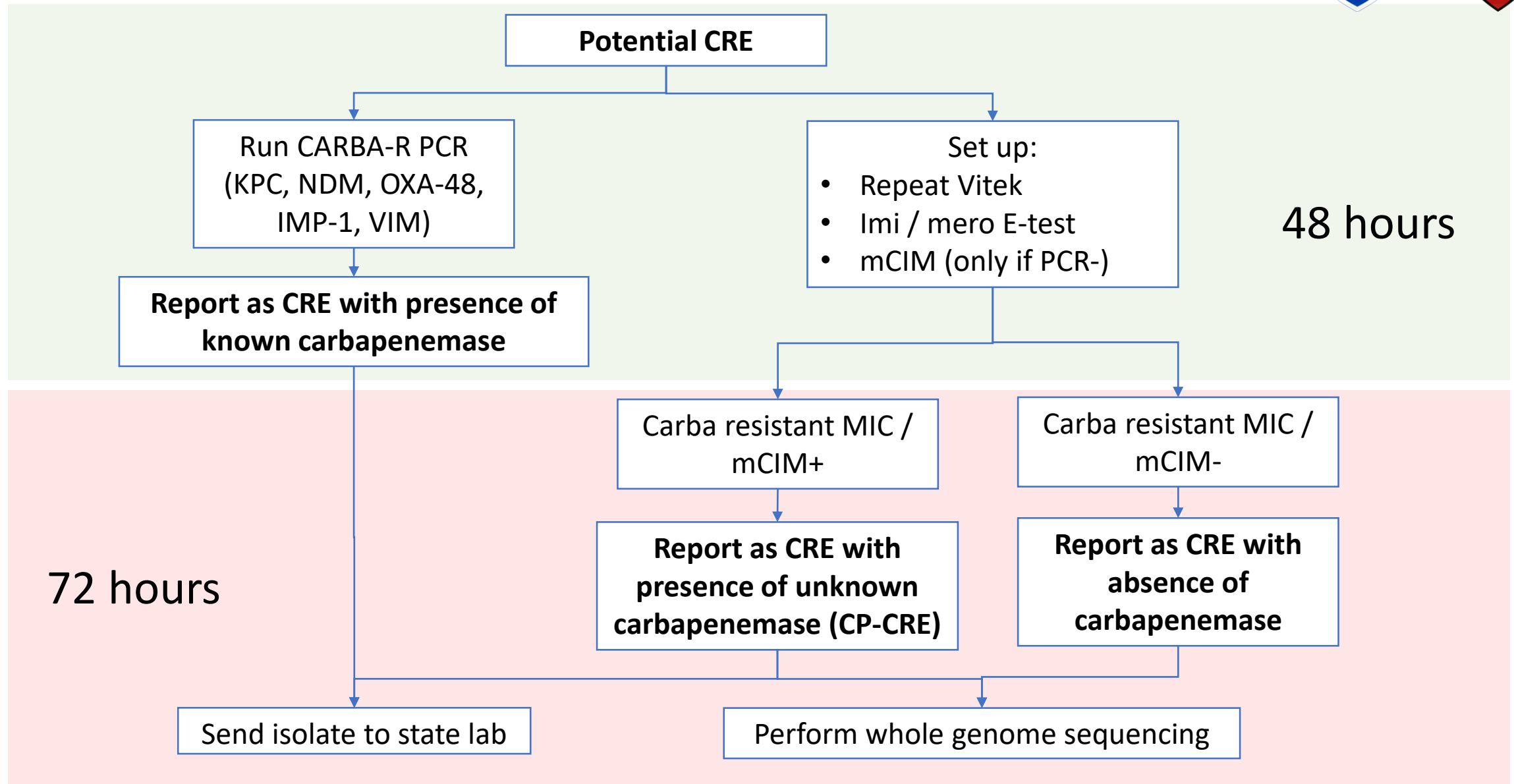
KLEBSIELLA PNEUMONIAE

| Antibiotic | | Interpretation | Value |
|-------------------------------|---|----------------|-------|
| Amikacin | | Resistant | ≥64 |
| Amoxicillin-clavulanate | | Resistant | ≥32 |
| Ampicillin | | Resistant | ≥32 |
| Cefazolin | | Resistant | ≥64 |
| Cefepime | | Resistant | ≥64 |
| Cefoxitin | | Resistant | ≥64 |
| Ceftazidime | | Resistant | ≥64 |
| Ceftriaxone | | Resistant | ≥64 |
| Ciprofloxacin | | Resistant | ≥4 |
| Gentamicin | | Resistant | ≥16 |
| Levofloxacin | | Resistant | ≥8 |
| Meropenem | | Resistant | ≥16 |
| Nitrofurantoin | | Resistant | ≥512 |
| Piperacillin-tazobactam | | Resistant | ≥128 |
| Tetracycline | | Resistant | ≥16 |
| Tobramycin | | Resistant | ≥16 |
| Trimethoprim/sulfamethoxazole | | Resistant | 80 |
| Comments | KLEBSIELLA PNEUMONIAE | | |
| | 100,000 colony forming units per ml KLEBSIELLA PNEUMONIAE | | |

CRE protocol activated!



CRE protocol activated!





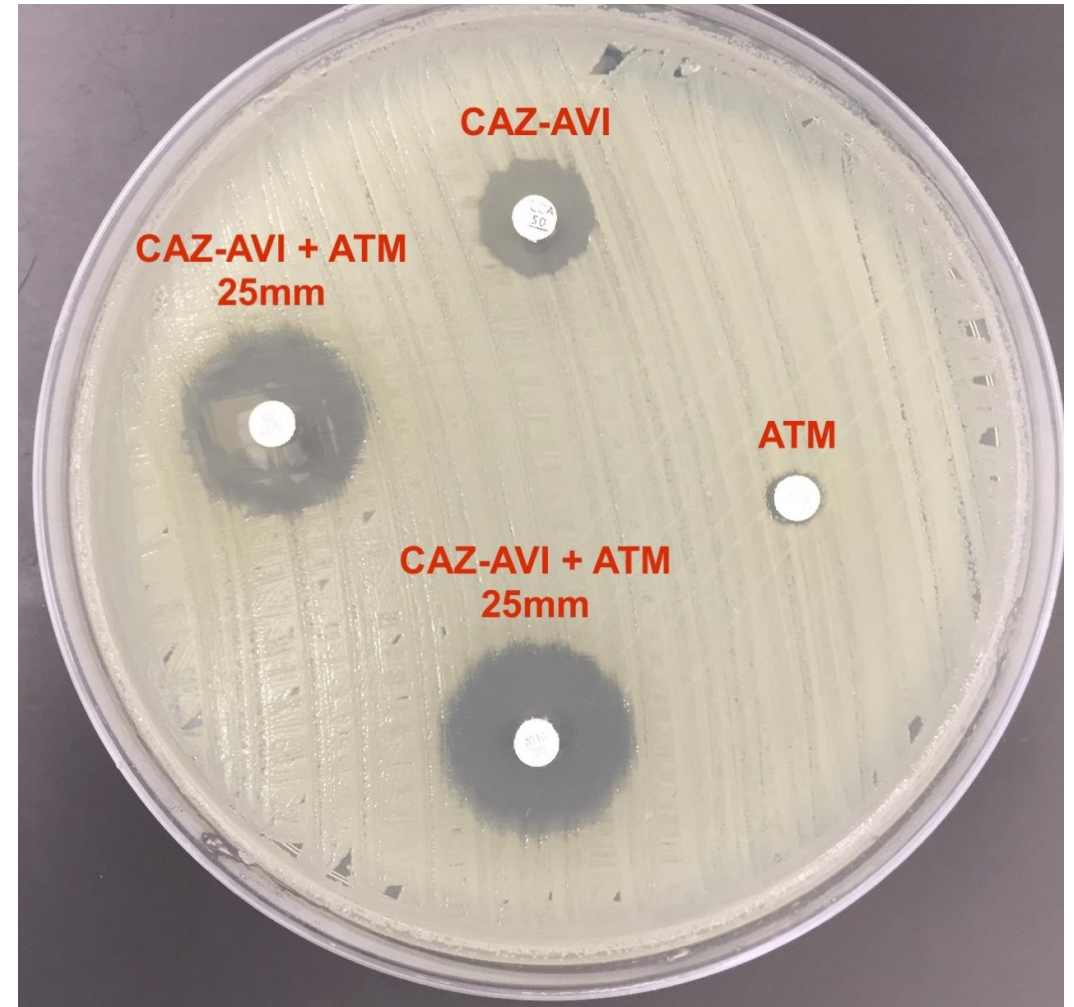
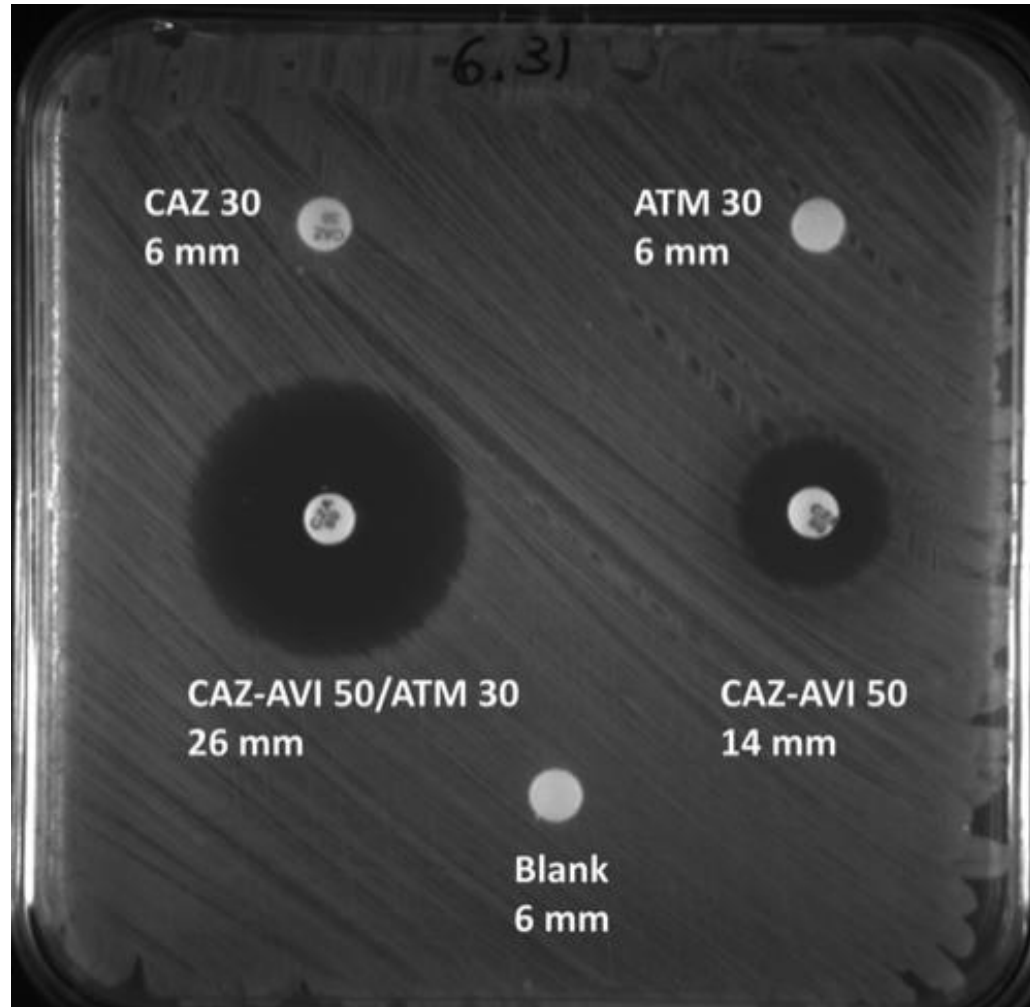
The Wild Wild East

- An NDM-1 metalloproteinase was identified
- Special unvalidated antibiotic synergy testing was initiated

Can Ceftazidime-Avibactam and Aztreonam Overcome β -Lactam Resistance Conferred by Metallo- β -Lactamases in *Enterobacteriaceae*?

Steven Marshall,^a Andrea M. Hujer,^{a,b} Laura J. Rojas,^{a,b,c}
Krisztina M. Papp-Wallace,^a Romney M. Humphries,^d Brad Spellberg,^e
Kristine M. Hujer,^{a,b} Emma K. Marshall,^a Susan D. Rudin,^{a,b} Federico Perez,^{a,b}
Brigid M. Wilson,^a Ronald B. Wasserman,^f Linda Chikowski,^g David L. Paterson,^h
Alejandro J. Vila,ⁱ David van Duin,^j Barry N. Kreiswirth,^k Henry F. Chambers,^l
Vance G. Fowler, Jr.,^m Michael R. Jacobs,ⁿ Mark E. Pulse,^o William J. Weiss,^o
Robert A. Bonomo^{a,b,c,p}

The Wild Wild East



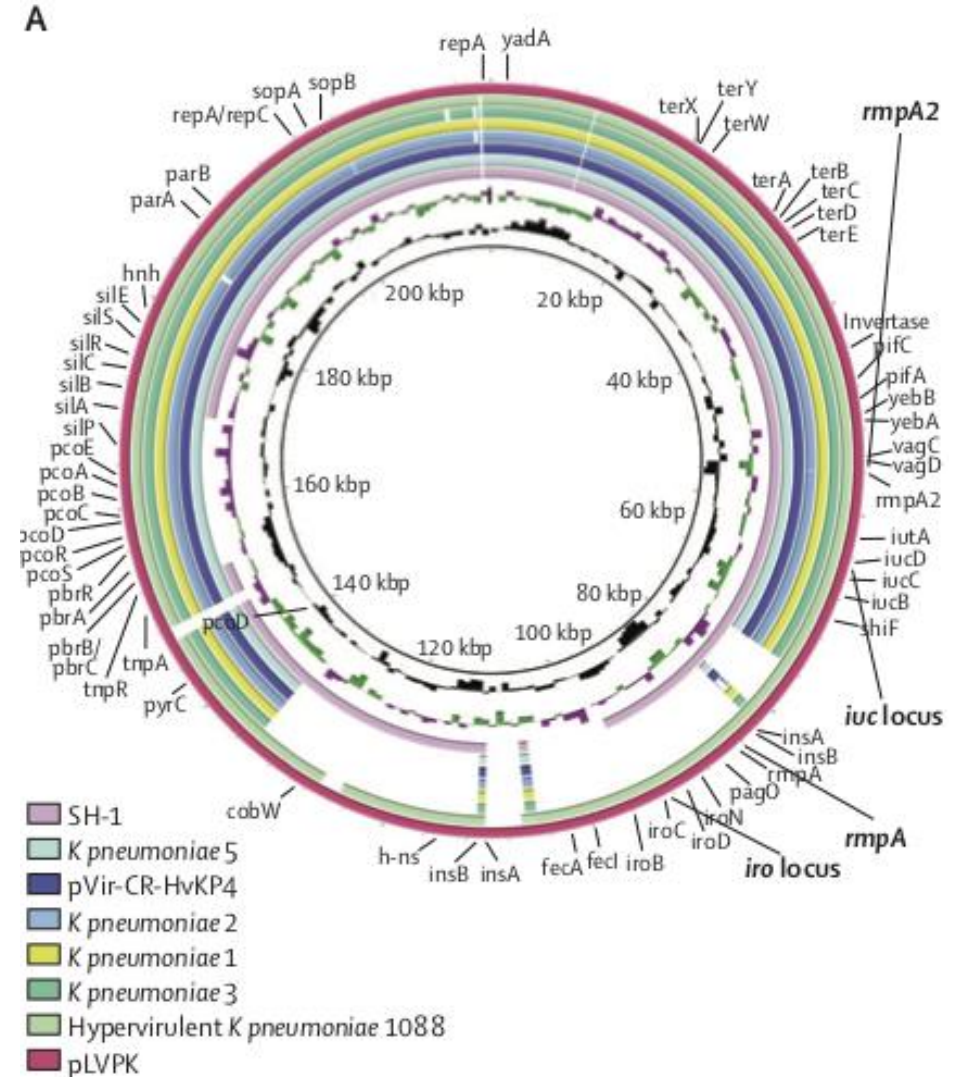


Micro saves the day (again)

- Patient had stents exchanged and was empirically started on meropenem and tigecycline but could not tolerate therapy
- She was switched to ceftazidime-avibactam + aztreonam and all subsequent urine cultures were negative for organisms
- She was able to start her chemotherapy

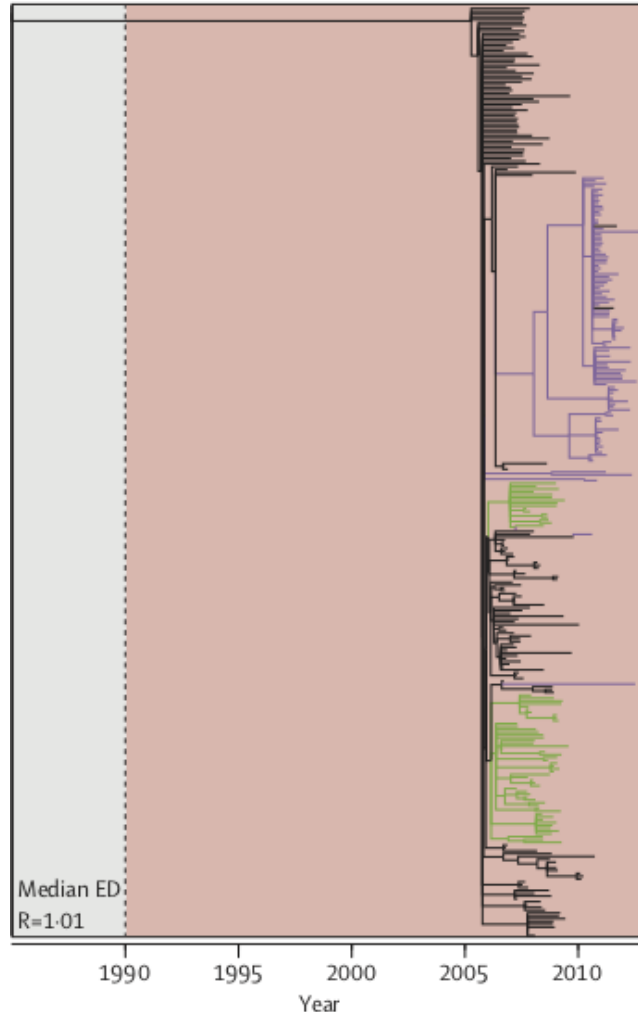
From one test, many answers (and even more questions)

- From a single sequence
 - Genetic basis of antibiotic susceptibility
 - Genetic basis of virulence
 - Presence of mobile genetic elements
 - Identification of new mechanisms of resistance



From one test, many answers (and even more questions)

D ST1(027)



E ST3(001)



- From comparison of multiple sequences
 - Identification of high-risk clones for infection control, state / national labs
 - Identification of nosocomial transmission events
 - Prediction of treatment response
 - Evolutionary dynamics of antibiotic resistance

Back to our patient

| Class | Subclass | Gene(s) |
|-----------------|--------------------------|----------------------------------|
| Beta-lactams | Penicillins | <i>SHV-11</i> |
| | Beta-lactam combinations | <i>CTX-M-15</i> |
| | Cephalosporins | <i>OXA-11</i> |
| | Carbapenems | <i>TEM-199</i> |
| Macrolides | | <i>msr(E)</i> |
| Aminoglycosides | | <i>APH(3')</i> <i>AAC(6')</i> |
| Rifamycins | | <i>arr-6</i> |
| DHFR | | <i>sul1</i> |
| Fosfomycin | | <i>fosA5</i> |
| Multidrug | | <i>emrD</i> |

- A second *K. pneumoniae* isolate obtained at the same time was identical except it lacked the NDM-1 gene
 - Dynamic acquisition and loss?
- 2 subsequent NDM-1+ isolates have been identified since October
 - No travel history for both

We serve multiple masters

Direct
patient care

Inpatient / outpatient
populations

Financial
imperatives

Policy makers

Providers

Antimicrobial
stewardship

Infection
control

State / national public
health agencies

Academic
researchers

Clinical microbiology
lab



Our deliverables are deceptively simple

- Organism identification and quantification
- Susceptibilities to anti-infectives
- Assaying indirect biomarkers of infection
 - β -D-glucan
 - Antigen / antibody tests

Our true deliverables span a much wider range



- For the provider
 - Communication of critical results
 - Interpretation of test results
 - Offline / unvalidated testing for challenging cases
 - Prompt send out to reference laboratories or initiation of additional testing
- For ASPs
 - Calculation of cumulative antimicrobial susceptibility reports
- For infection control
 - Surveillance of high-risk organisms
 - Identification of new resistant phenotypes
 - Identification of newly emerging or highly dangerous pathogens
- For all
 - Diagnostic stewardship



A major headache

- 27 year old female presents in February 2018 with a 1 day history of severe headache and vomiting.
- She denies fever, chills and has no sick contacts
- The week before her presentation she had travelled to Mexico for vacation and stayed at a resort
 - While there, she ate some soft cheeses

A major headache

- A lumbar puncture was performed

| CSF CHEMISTRIES | |
|--------------------|----------|
| Lactate, CSF | 2.6 * |
| Glucose, CSF | 63 * |
| Total Protein, CSF | 70.5 * ▲ |

- She is started on broad-spectrum antibiotics and antivirals
- A specimen was run on the Biofire meningitis/encephalitis platform

| CSF COUNTS AND DIFF | | |
|-----------------------|-------------|-------------|
| Color, CSF | COLORLESS | COLORLESS |
| Turbidity/Appearan... | SL HAZY ! | SL HAZY ! |
| RBC, CSF | 183 ! | 8 ! |
| Nucleated cells, CSF | 263 * !!▲ | 463 * !!▲ |
| Blasts, CSF (%) | 0 | 0 |
| Bands, CSF (%) | 0 | 0 |
| Neutrophils, CSF (%) | 74 | 58 |
| Lymphs, CSF (%) | 12 | 21 |
| Monos, CSF (%) | 13 | 19 |
| Eos, CSF (%) | 0 | 0 |
| Basos, CSF (%) | 0 | 1 |
| Plasma cells, CSF | | 1 |
| NRBC#, CSF | 0 | 0 |
| Xanthochromia, CSF | NOT PRESENT | NOT PRESENT |

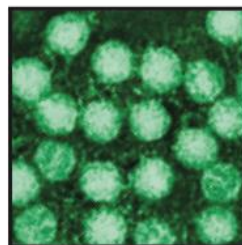
What is the **BIO** **FIRE**?

- The first FDA-approved syndromic panel for diagnosis of meningitis / encephalitis



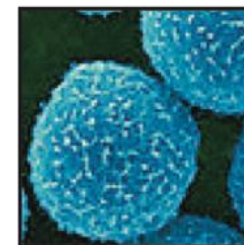
Bacteria

Escherichia coli K1
Haemophilus influenzae
Listeria monocytogenes
Neisseria meningitidis
Streptococcus agalactiae
Streptococcus pneumoniae



Viruses

Cytomegalovirus (CMV)
 Enterovirus
 Herpes simplex virus 1 (HSV-1)
 Herpes simplex virus 2 (HSV-2)
 Human herpesvirus 6 (HHV-6)
 Human parechovirus
 Varicella zoster virus (VZV)



Fungi

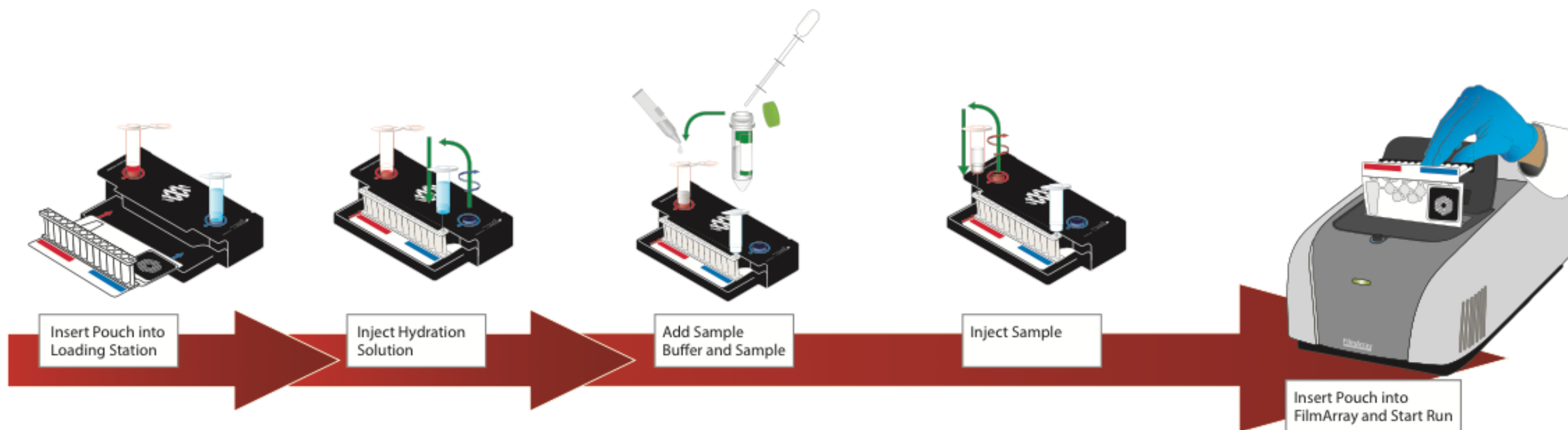
Cryptococcus neoformans/gattii



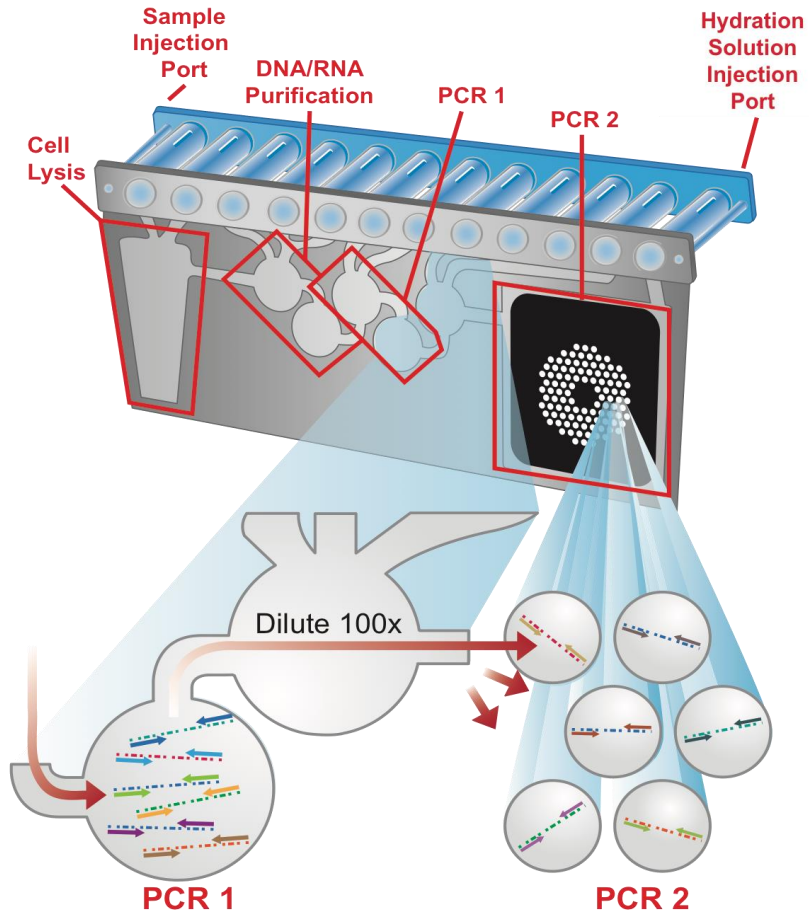
Why use the BIO FIRE?

- Diagnosis of meningitis and encephalitis is really hard
 - Non-specific symptoms
 - Deadly for some but not all organisms
 - Sensitivity of traditional diagnostics is poor
- Minimal hands-on (5') and rapid turnaround (60') times
- Detects most common pathogens
- High sensitivity

The FilmArray

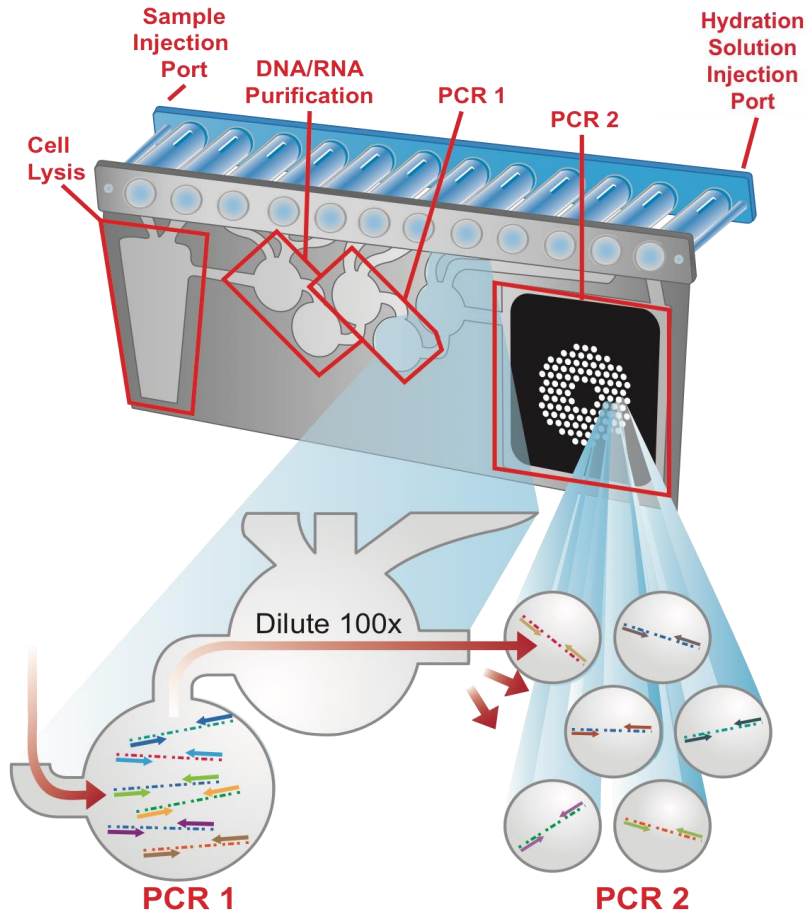


The FilmArray



- Step 1: Cell lysis
 - Ceramic beads break up human and bacterial cells and virions
 - Free RNA/DNA bound by magnetic beads and moved to purification chamber

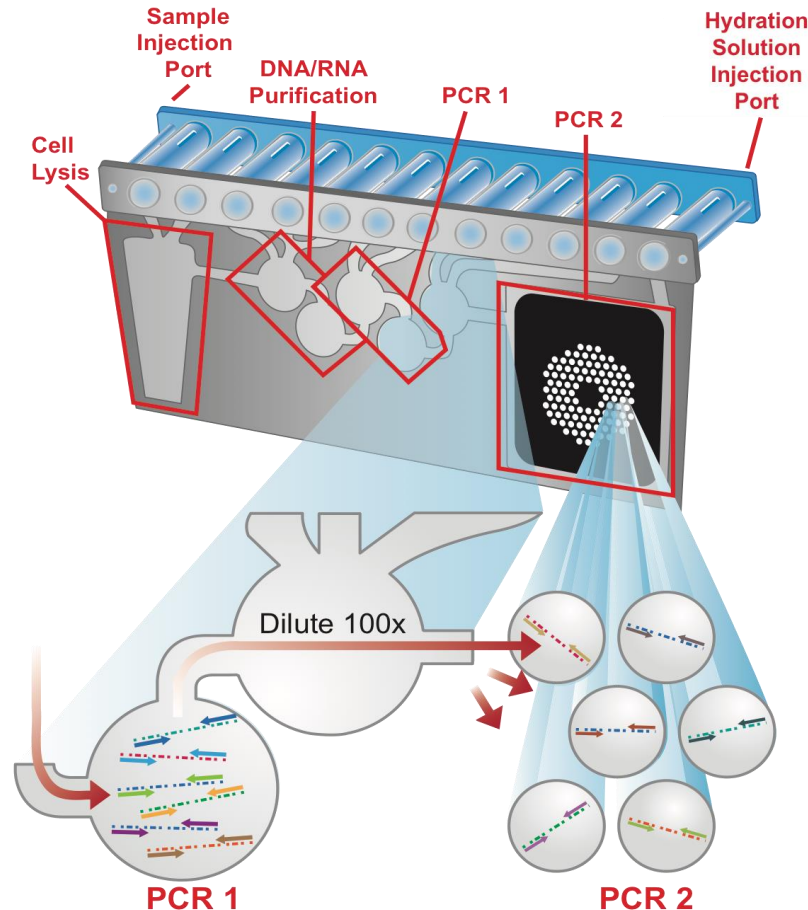
The FilmArray



• Step 2: Purification

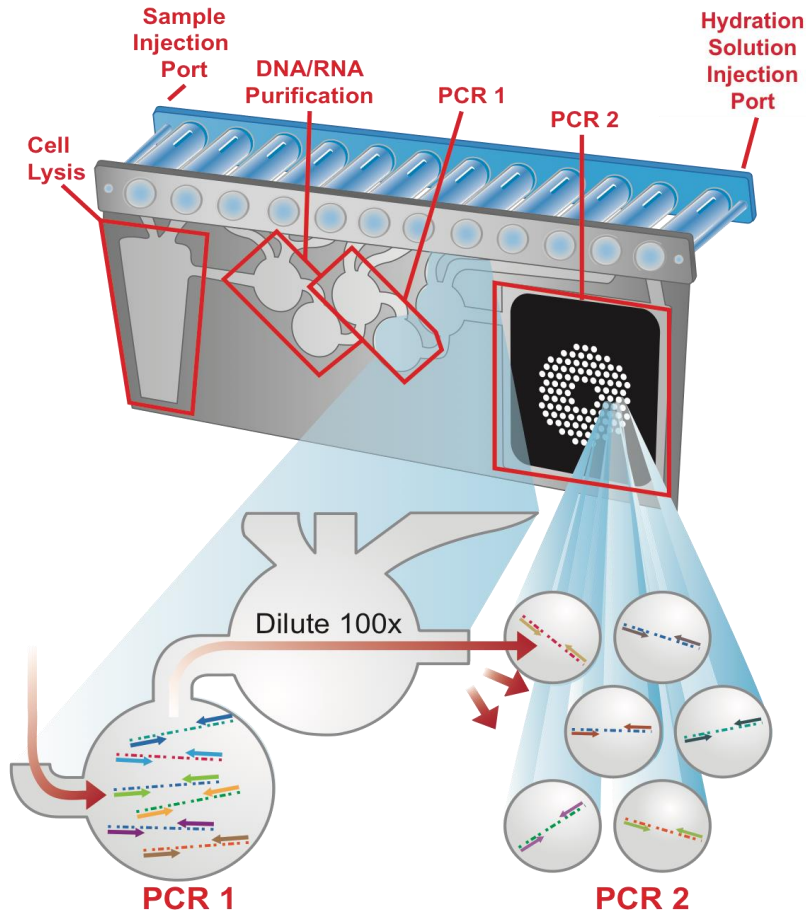
- Beads held stationary by a magnet outside of the pouch while remaining cellular debris is washed away
- Beads released and elution buffer washes the nucleic acid off of them
- Beads are again held by magnets as eluted nucleic acids move to next chamber

The FilmArray



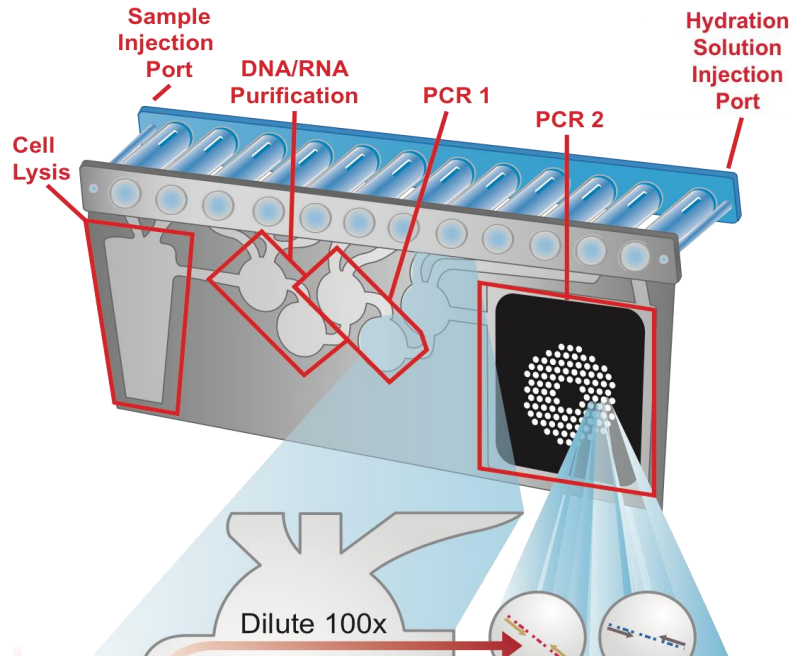
- Step 3: PCR stage 1
 - RNA converted to DNA
 - Multiplex PCR amplification
 - Buffer added to dilute remaining unbound PCR primers
 - Fresh master mix added and solution added to each cell of microarray

The FilmArray



- Step 4: PCR stage 2
 - Wells spotted with 2nd stage primers highly specific to amplicons made in the first stage
 - Detected through addition of a fluorescent dye that binds ds-DNA
- Step 5: DNA melting
 - Compares the measured T_m to known T_m of ds-DNA increases specificity

The FilmArray



- Advantages
 - Closed system
 - Required volume: 200ul
 - Hands on time: ~5 minutes
 - Analysis time: 60 minutes
- Disadvantages
 - Closed system

Clin Chem Lab Med. 2017 Aug 1. pii: /j/cclm.ahead-of-print/cclm-2017-0518/cclm-2017-0518.xml. doi: 10.1515/cclm-2017-0518. [Epub ahead of print]

False negative results caused by erroneous automated result interpretation algorithm on the FilmArray 2.0 instrument.

Lee CK¹, Chiu L¹, Yan G¹, Chew KL¹, Yan B¹, Jureen R¹, Loh TP¹.

Back to our patient

- Antibiotics were discontinued and the patient's symptoms resolved in 48 hours
- Infection was presumed acquired in Mexico

| | |
|-----------------------|--------------|
| CRYPTOCOCCUS NEO/GAT | Not Detected |
| CYTOMEGALOVIRUS | Not Detected |
| ENTEROVIRUS | DETECTION ! |
| ESCHERICHIA COLI K1 | Not Detected |
| H. INFLUENZAE | Not Detected |
| HSV 1 | Not Detected |
| HSV 2 | Not Detected |
| HUMAN HERPESVIRUS 6 | Not Detected |
| HUMAN PARECHOVIRUS | Not Detected |
| L. MONOCYTOGENES | Not Detected |
| N. MENINGITIDIS | Not Detected |
| S. AGALACTIAE (GRP B) | Not Detected |
| STREP. PNEUMO | Not Detected |
| VZV | Not Detected |



Challenges

BE THOROUGH

High penalty for incorrect identification or AST results

BE FAST

Intense pressure to report results as fast as possible

- The availability of effective empiric antibiotic therapy provides the lab a safety window within which they can perform a workup with relatively low harm to the patient



A false sense of security

BE THOROUGH

High penalty for incorrect identification or AST results

BE FAST

Intense pressure to report results as fast as possible

- Empiric antibiotic therapy is not without significant problems
 - Inadequate therapy associated with poor treatment outcomes
 - Overly broad therapy associated with selection of drug resistant organisms



A false sense of security

BE THOROUGH

High penalty for incorrect identification or AST results

BE FAST

Intense pressure to report results as fast as possible

- The philosophy of clinical microbiology labs has been to favor patient-level outcomes over population-level outcomes
- Delays in reporting make de-escalation difficult



The future?

BE THOROUGH

High penalty for incorrect identification or AST results

BE FAST

Intense pressure to report results as fast as possible

- Advances in the sensitivity, specificity and throughput of molecular platforms has engendered intense academic and commercial interest
- The hope is for these technologies to resolve the tension between accuracy and turnaround time

Syndromic panels are IN

| Syndrome | Manufacturer | Model | Market entry |
|---------------------------|--------------|------------------------------------|--------------|
| Bloodstream infection | Biofire | FilmArray | 2013 |
| | Luminex | Verigene GP | 2012 |
| | | Verigene GN | 2014 |
| | Accelerate | Pheno | 2017 |
| Respiratory infection | Biofire | FilmArray | 2011 |
| | GenMark | eSensor RVP | 2013 |
| | | ePlex | 2017 |
| | Luminex | xTAG RVP v1 | 2008 |
| | | RVP Fast | 2011 |
| | | Verigene Respiratory Pathogen Flex | 2015 |
| | | NxTAG | 2015 |
| Meningitis / encephalitis | Biofire | FilmArray | 2015 |



Why newer platforms will not replace traditional diagnostics (yet)

- Do not perform comprehensive AST*
- Finite range of targets
- Poor performance in polymicrobial infections
- Slightly decreased sensitivity



Why newer platforms will not replace traditional diagnostics (yet)

- Results from such platforms will arrive within a few hours and should be viewed as 'preliminary'
 - The understanding that additional confirmation and/or information will arrive in the following 24 – 48 hours through older algorithms
 - Would not change initial empiric management
- Cost and hands-on time prohibits running on every patient sample

With more data comes more complexity

| | | TRUTH | |
|----|----------|----------|------------|
| | | Infected | Uninfected |
| T1 | Positive | TP | FP |
| | Negative | FN | TN |

With more data comes more complexity

| | | TRUTH | |
|----|----------|--|--|
| | | Infected | Uninfected |
| T1 | Positive | <div> <div>T2</div> <div> <div>+</div> <div>T1TP T2TP</div> </div> <div> <div>-</div> <div>T1TP T2FP</div> </div> </div> | <div> <div>T2</div> <div> <div>+</div> <div>T1FP T2FP</div> </div> <div> <div>-</div> <div>T1FP T2TN</div> </div> </div> |
| | Negative | <div> <div>T2</div> <div> <div>+</div> <div>T1FN T2TP</div> </div> <div> <div>-</div> <div>T1FN T2FN</div> </div> </div> | <div> <div>T2</div> <div> <div>+</div> <div>T1TN T2FP</div> </div> <div> <div>-</div> <div>T1TN T2TN</div> </div> </div> |



Take home points

- The clinical micro lab is a dynamic and exciting environment
- Much of what we do depends heavily on the skills of the lab staff
- Our work connects not only to direct patient care but to systems of public health
- The field is undergoing a sea change in terms of our approaches and requires a new set of critical thinking skills

Thank you!

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