

# Advanced Molecular Detection and Epidemiology

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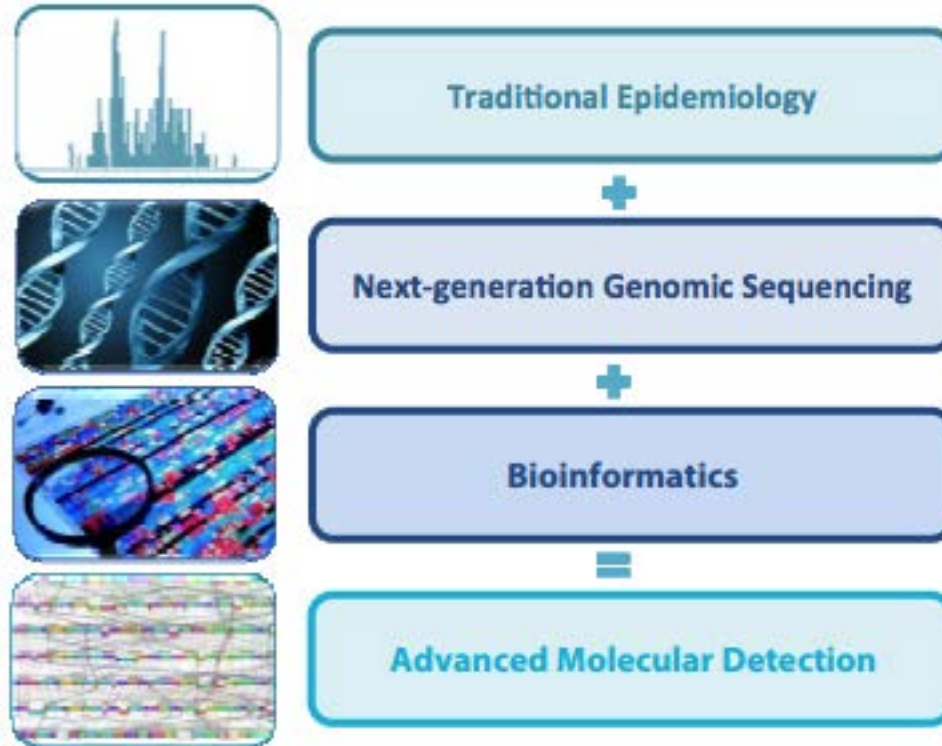
- **No Financial Disclosures**
- **The findings and conclusions in this presentation are those of the author and do not necessarily represent the official position of the Centers for Disease Control and Prevention.**



## Preview

- **Impact of incorporating whole genome sequencing (WGS)**
- **Role of epidemiological data in interpretation of WGS data**
- **Potential use of Advanced Molecular Detection technology in infection control**

# What is Advanced Molecular Detection (AMD)?



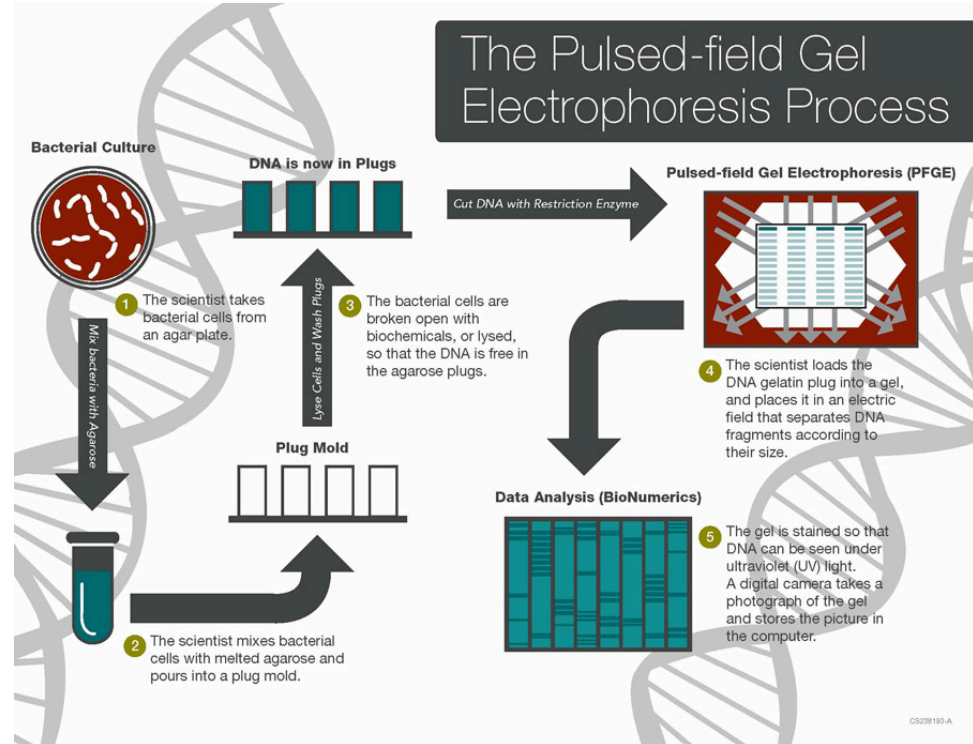


# Advanced Molecular Detection (AMD)

- **Impact public health**
  - Investigate outbreaks
  - Understand antibiotic resistance
  - Diagnose new pathogens
  - Identify control points
- **New technologies**
  - Next generation sequencing platforms
  - Supercomputers

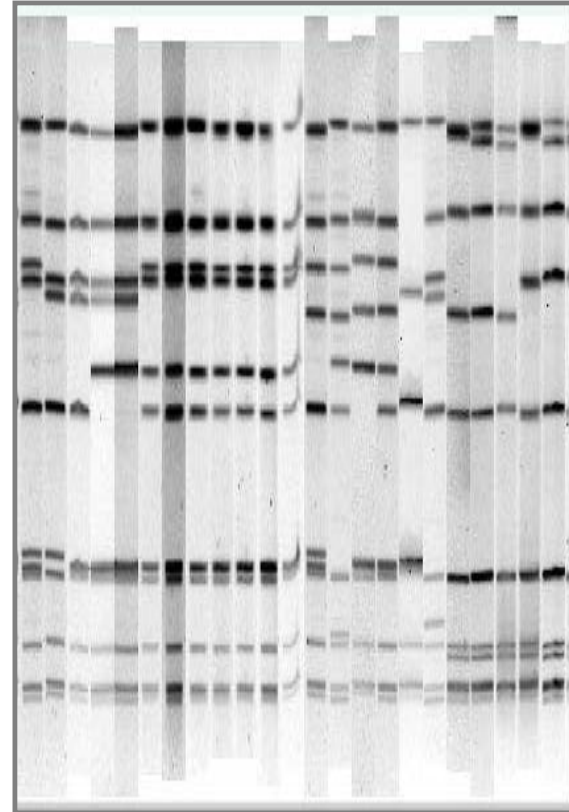
# Pulsed-Field Gel Electrophoresis (PFGE): Strengths

- **Universal subtyping method**
- **Reproducible**
- **Outbreaks**
  - Flag disseminated foodborne outbreaks (space, time)
  - Identify isolates that are likely to have come from a common source



# Pulsed-Field Gel Electrophoresis (PFGE): Limitations

- **Relatedness is not a true phylogenetic measure**
  - Related isolates may have different PFGE pattern
  - Unrelated isolates may have same PFGE pattern
- **Cannot discriminate between epidemiologically unrelated isolates**
- **Information only at the cut sites**
- **PFGE  $\neq$  WGS**



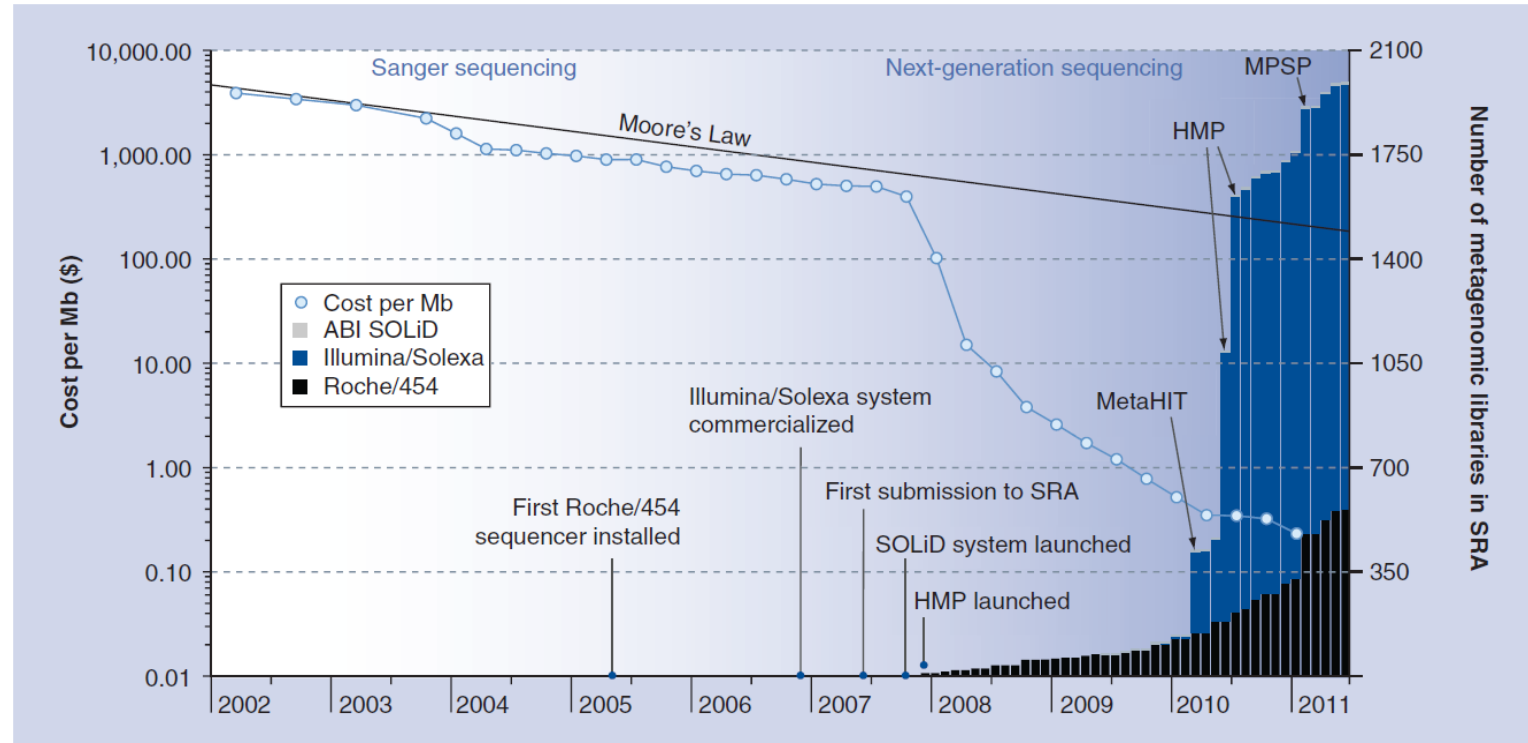
# Whole Genome Sequencing (WGS): Strengths

- High resolution sequence data
- Information at every point in the genome
  - True phylogenetic relatedness
  - Transmission path
  - Identify control points





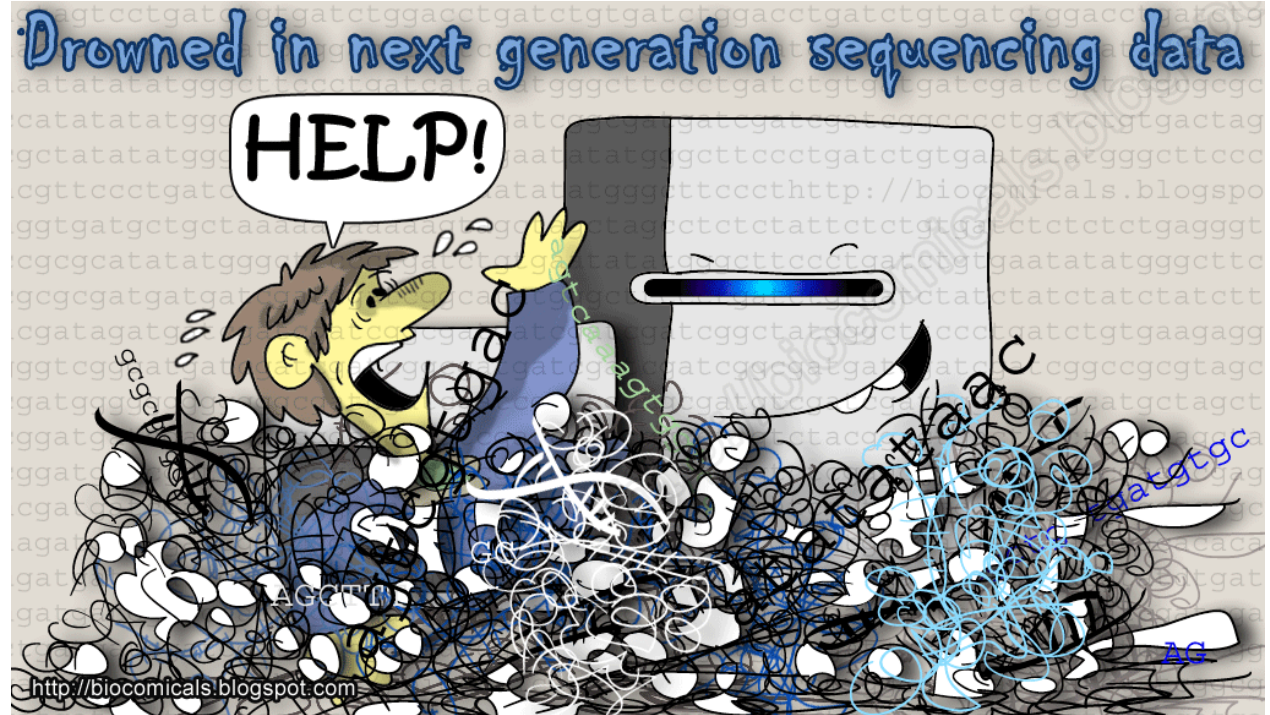
# Impact of Next Generation Sequencing



# Impact of Next Generation Sequencing

## ■ Challenges

- Data overload
- Interpretation

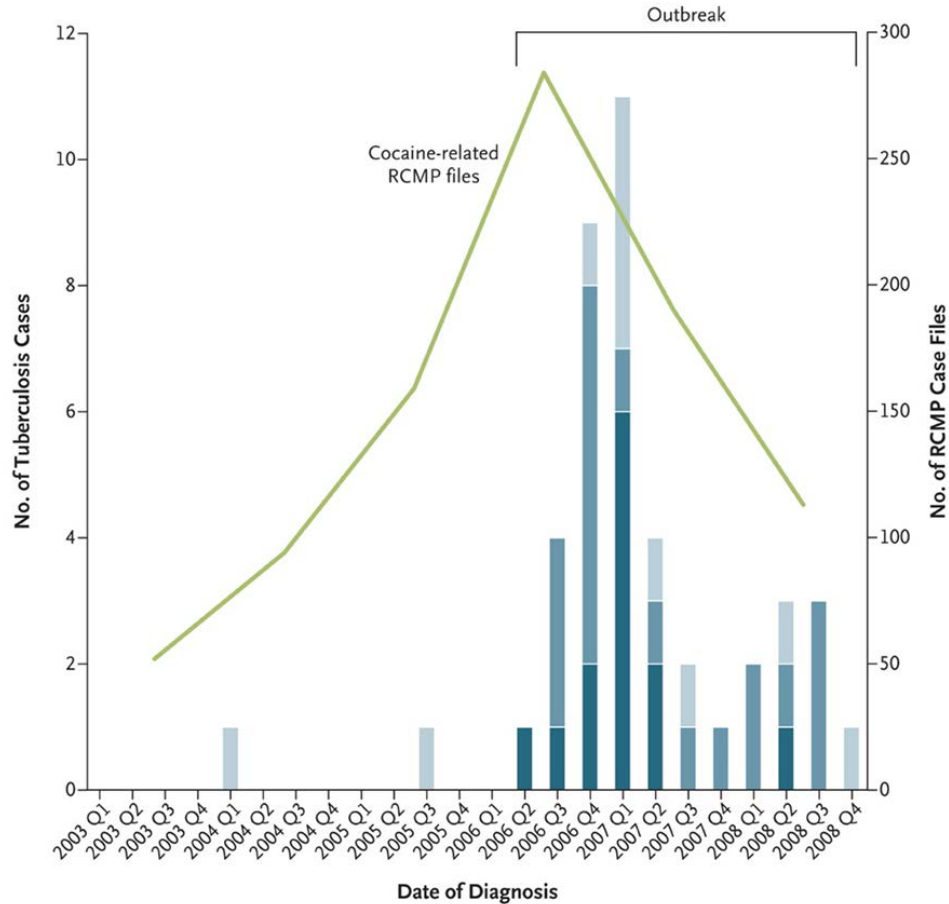




# Integration of Epidemiology and WGS



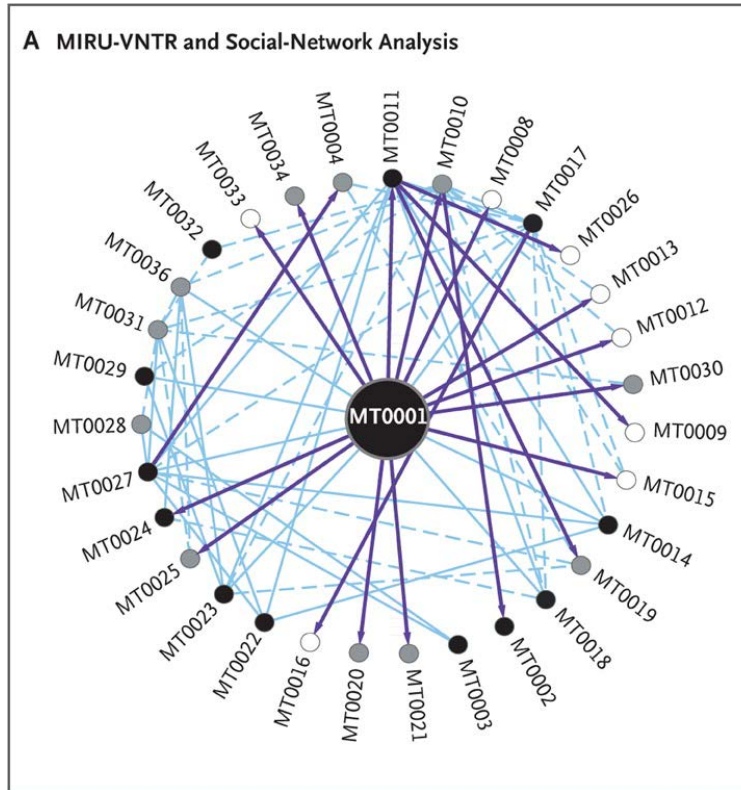
# *Mycobacterium tuberculosis* Outbreak



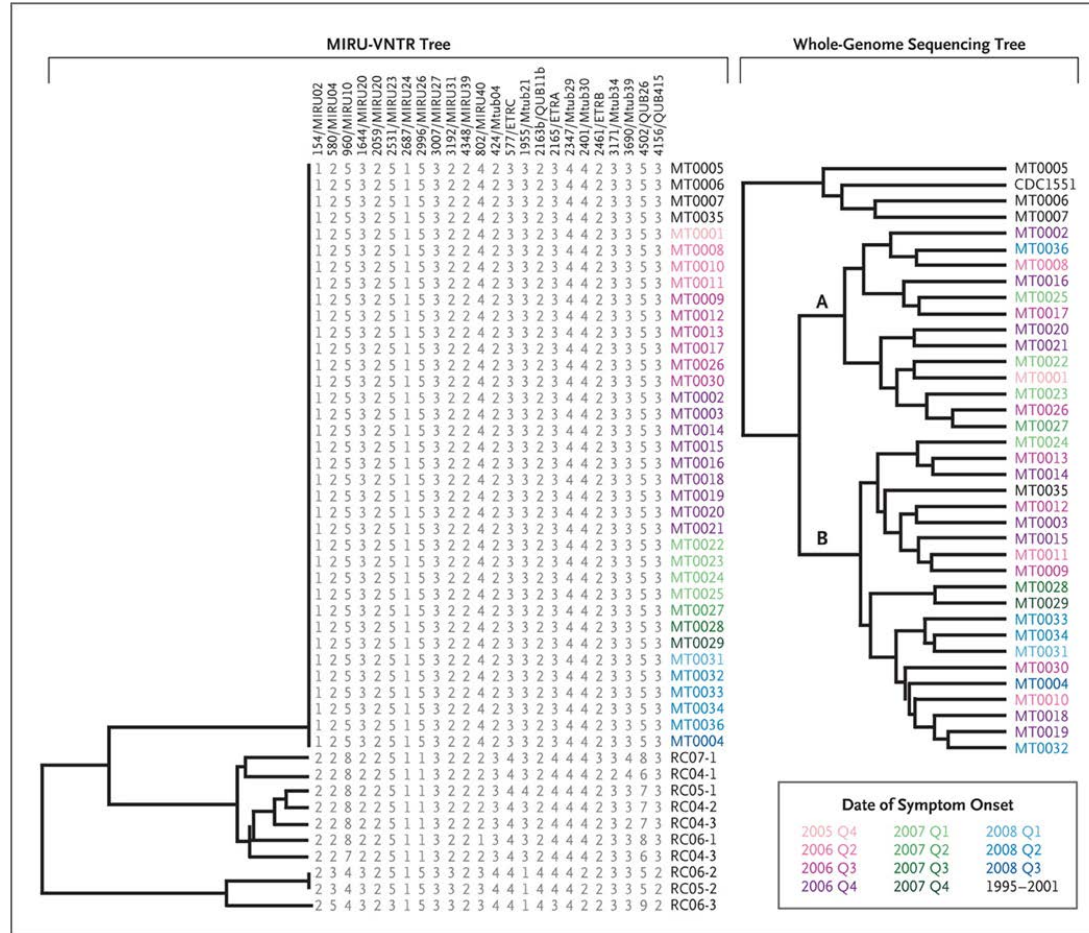




# *Mycobacterium tuberculosis* Outbreak

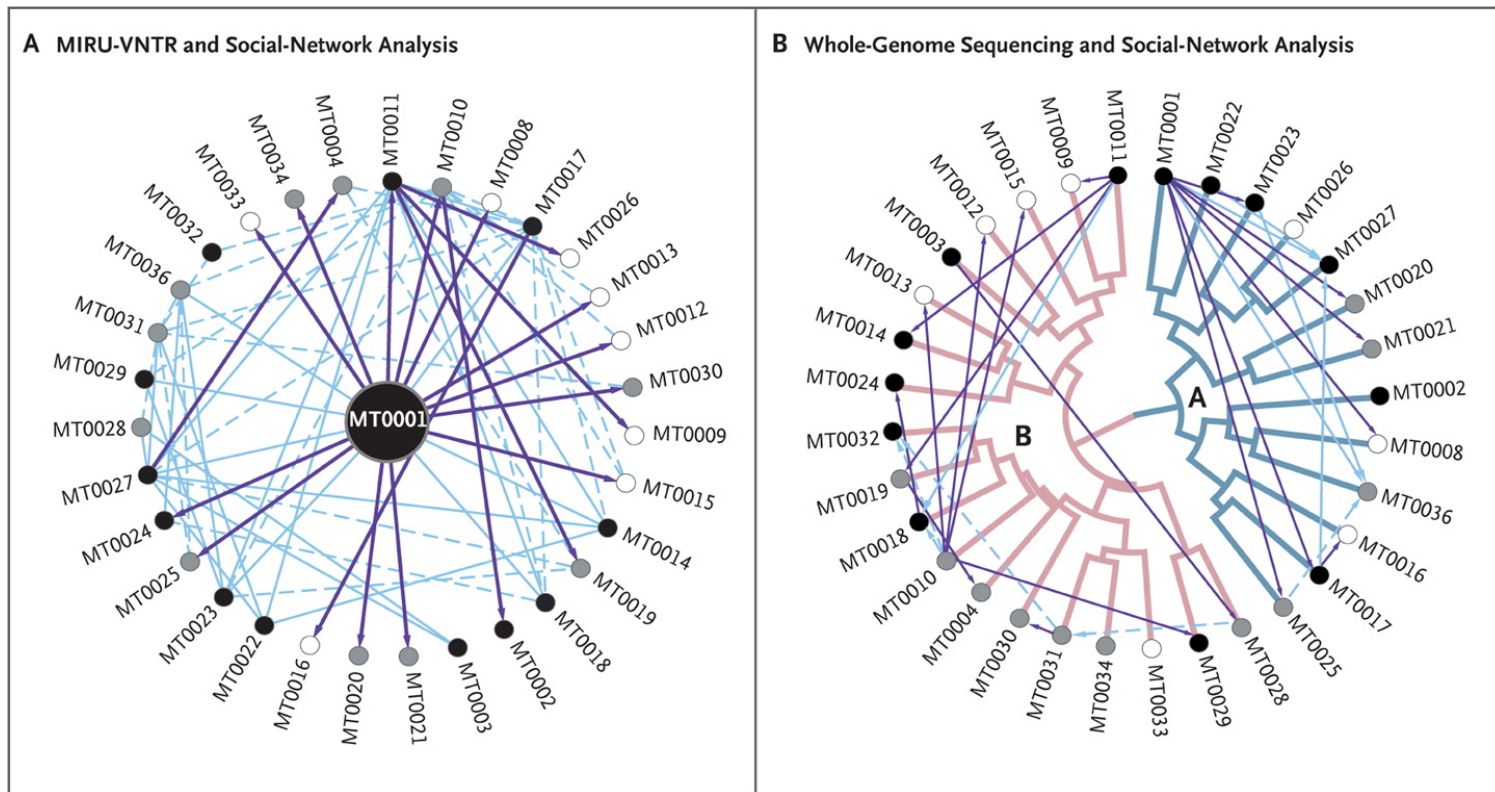


# Mycobacterium tuberculosis Outbreak





# Mycobacterium tuberculosis Outbreak



- Identified transmission pathways
- Revealed socioenvironmental factor that led to outbreaks

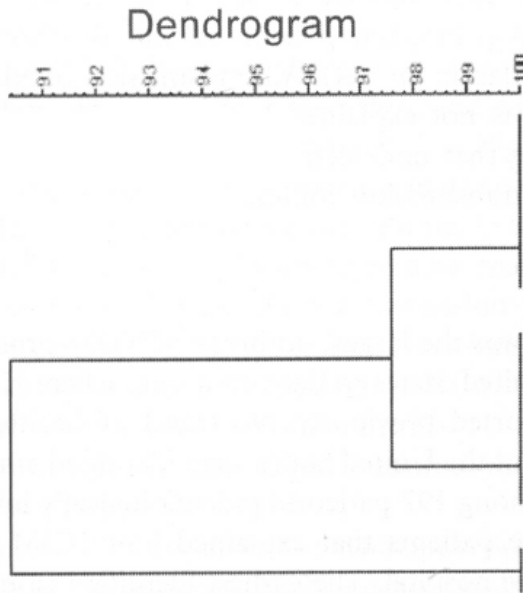


**Healthcare-associated infection**

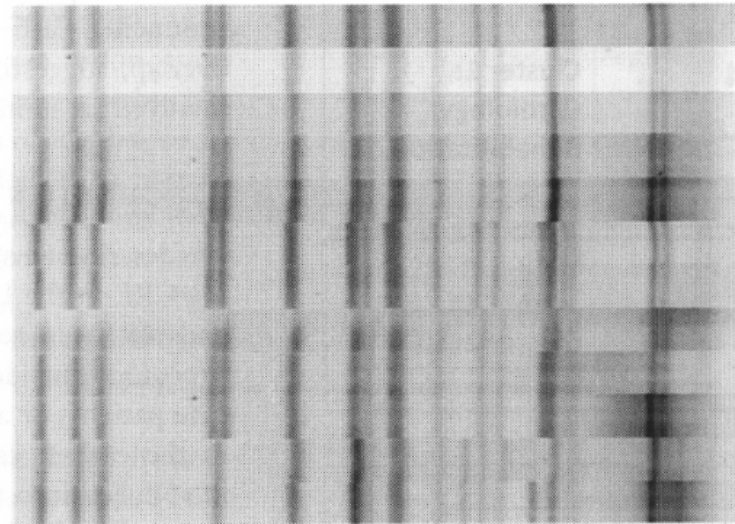
# Denver, Colorado - 2012

- 8 patients in a single acute care hospital
- Carbapenem-resistant *Klebsiella pneumoniae* producing New Delhi Metallo-Beta-Lactamase (NDM)
- Laboratory typing
  - Pulsed-field gel electrophoresis (PFGE)
  - Whole genome sequencing (WGS)

# Denver, Colorado - 2012



PFGE Pattern



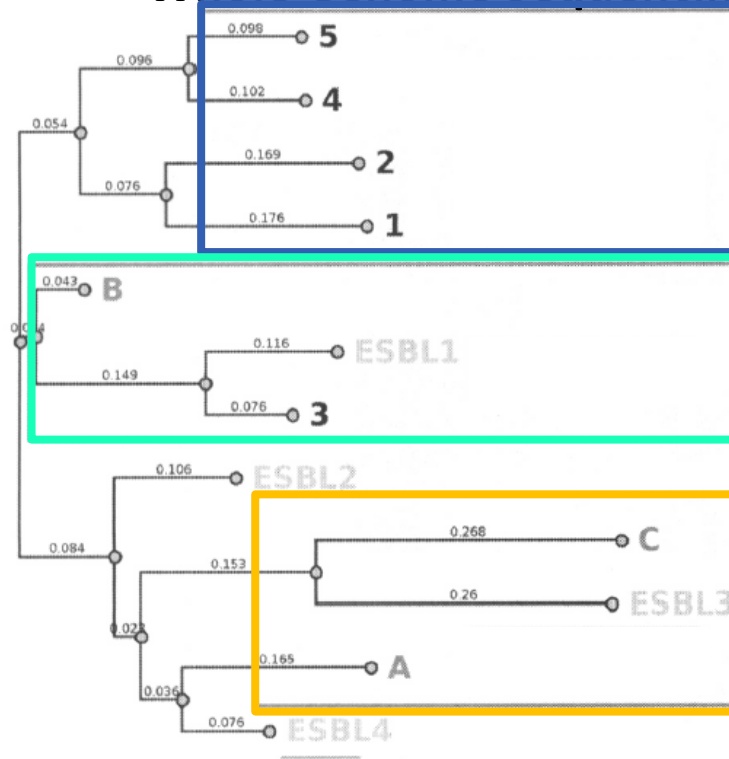
Isolate NDM

ESBL 4	-
C	+
ESBL 3	-
5	+
ESBL 2	-
A	+
B	+
4	+
2	+
1	+
3	+
ESBL 1	-



# Denver, Colorado - 2012

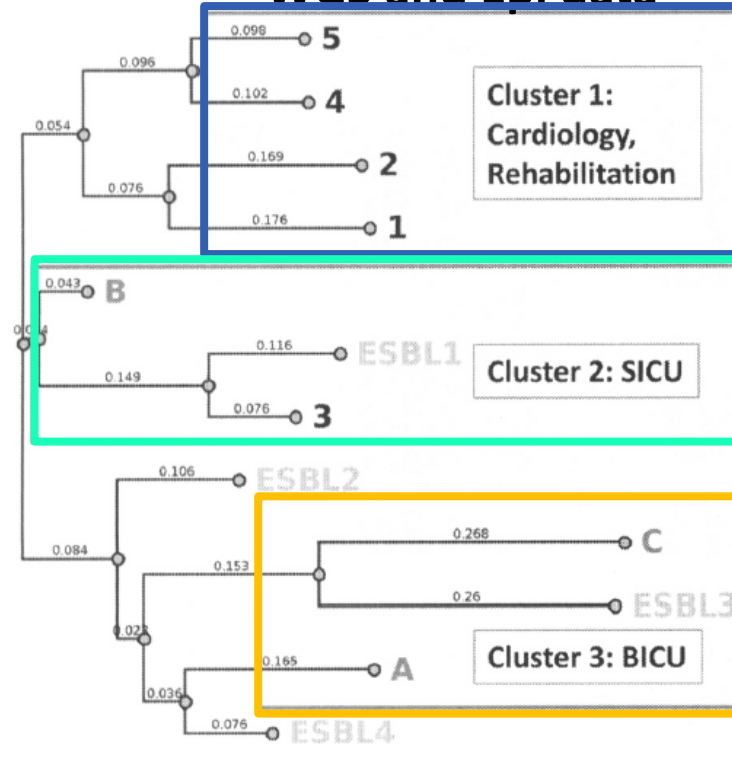
## Whole Genome Sequencing





# Denver, Colorado - 2012

## WGS and Epi data



Identified 3 areas where improvements in infection control were needed



# Infection Control and Prevention



# Standard Infection Prevention Strategies

## ■ Standard Precautions

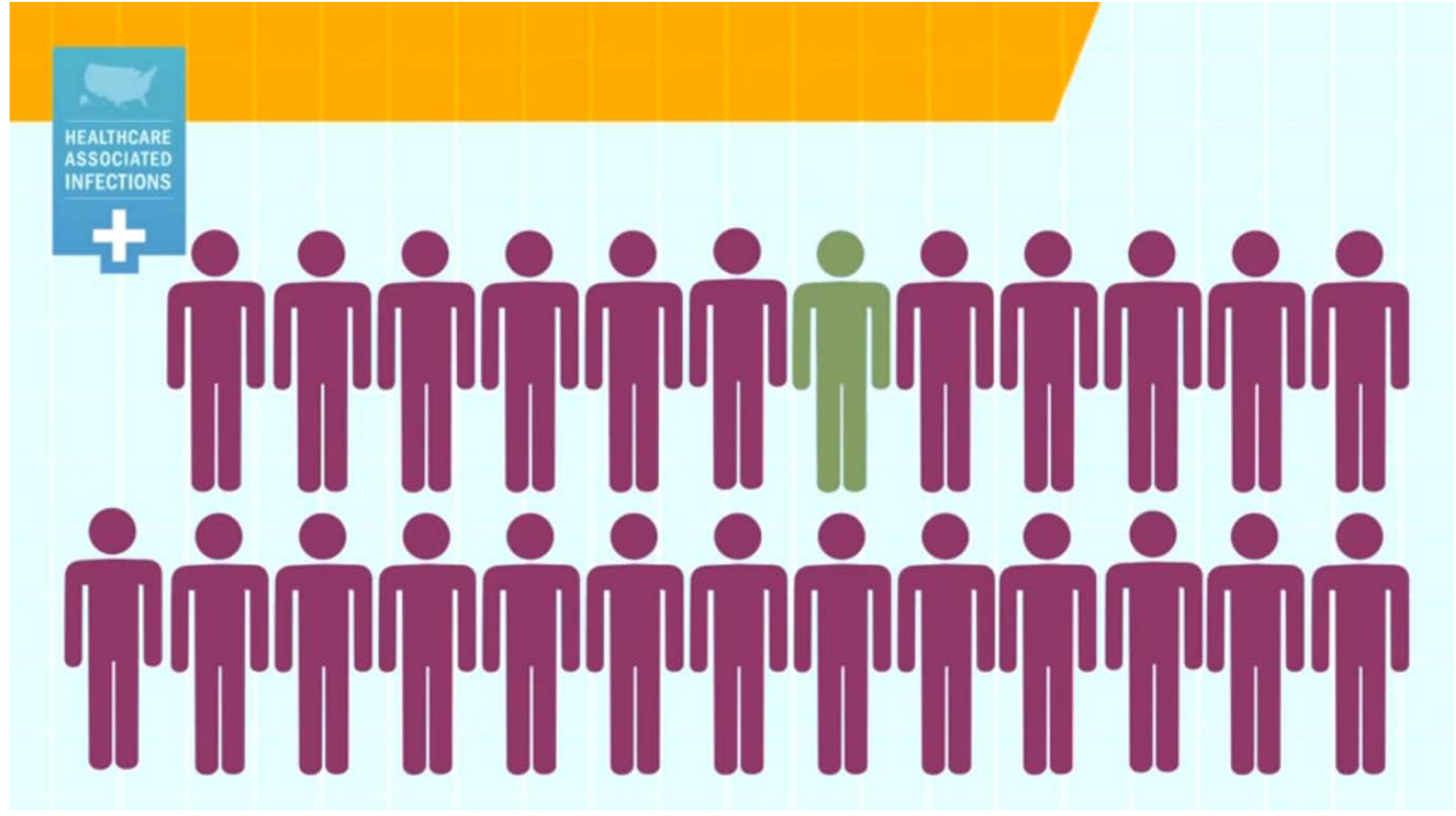
- Hand hygiene
- Personal Protective Equipment
- Respiratory hygiene
- Medication safety
- Injection safety
- Environmental/equipment cleaning and disinfection

## ■ Transmission-based Precautions

- Contact Precautions (e.g., CDI)
- Droplet Precautions (e.g., respiratory viruses)
- Airborne Precautions (e.g., TB, measles)

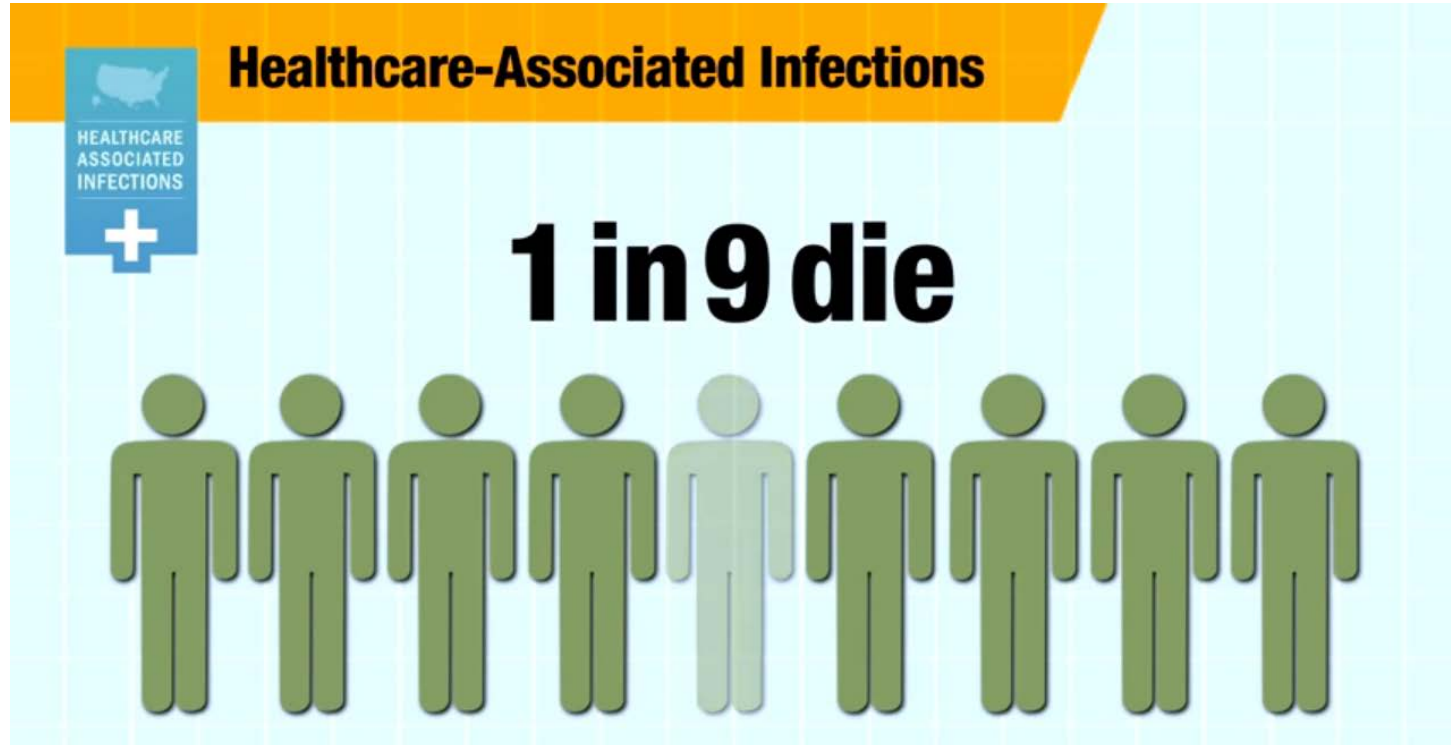
## ■ Antibiotic Stewardship

On any given day, about 1 in 25 hospital patients has at least one healthcare-associated infection (HAI)



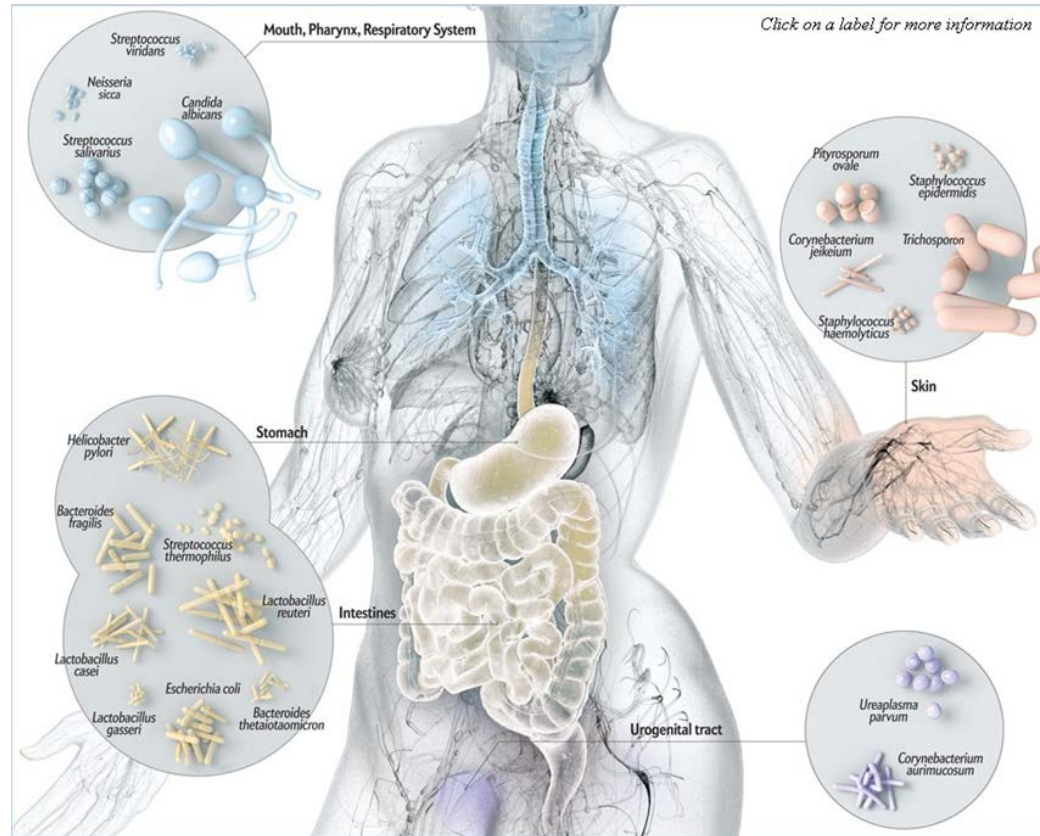


One in every 9 patients who gets a healthcare-associated infection will die during their hospitalization



- Vital to continue progress in healthcare epi and implementation research

# Protective Role of the Microbiota

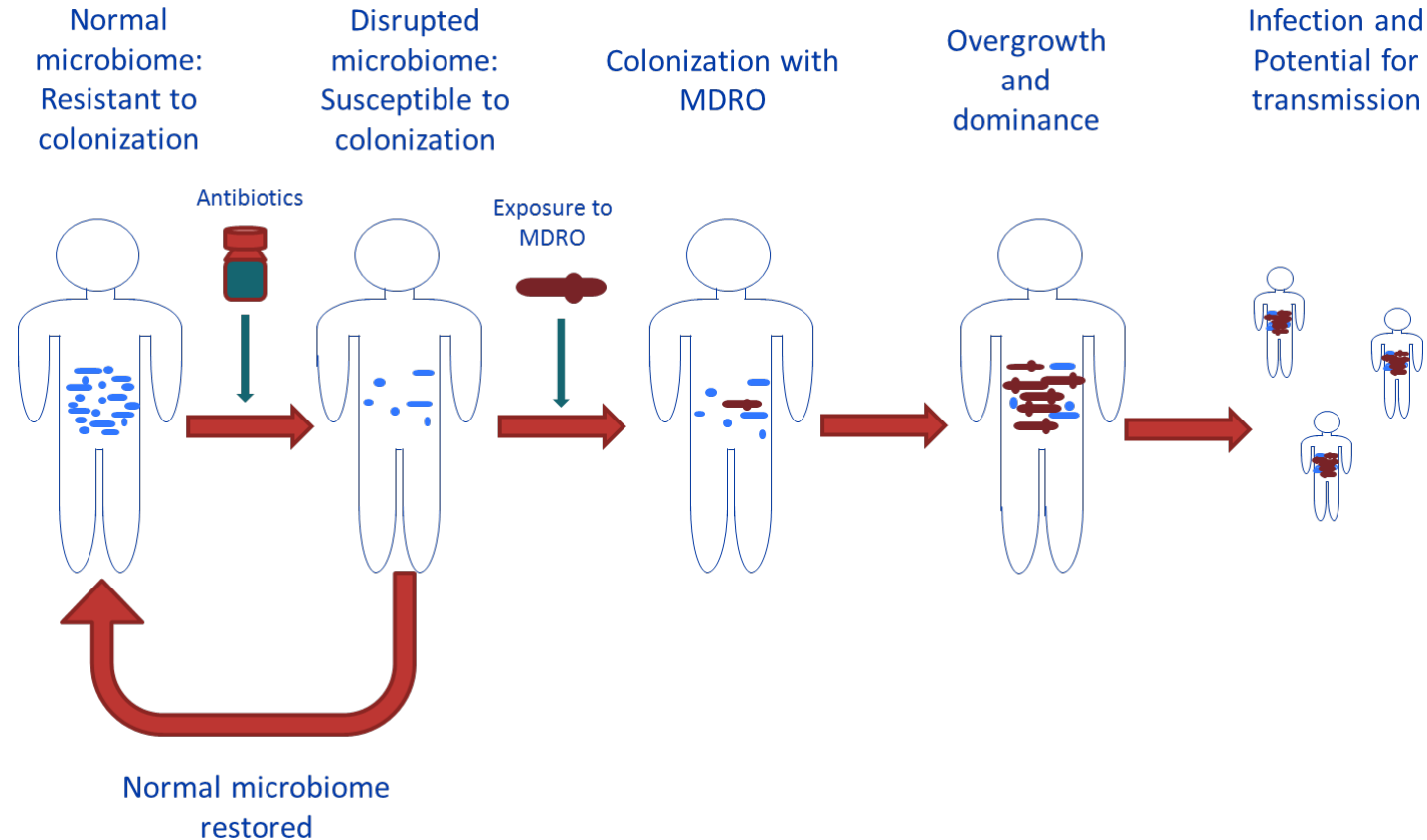




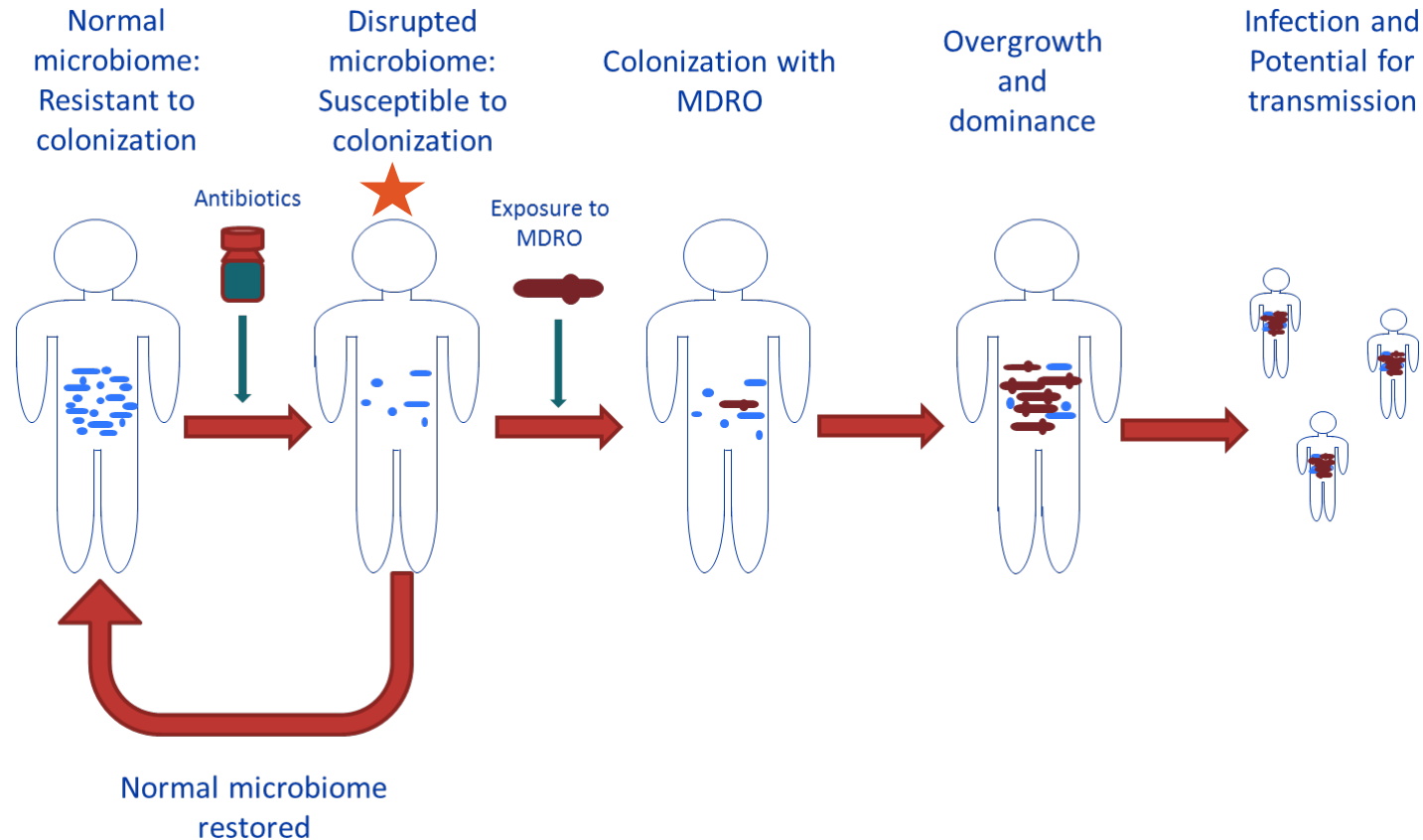
# Microbiome Disruption Indices (MDI)

- CDC working to develop
- Standardized criteria
- Characterize major human microbiomes
  - Lower intestinal microbiome

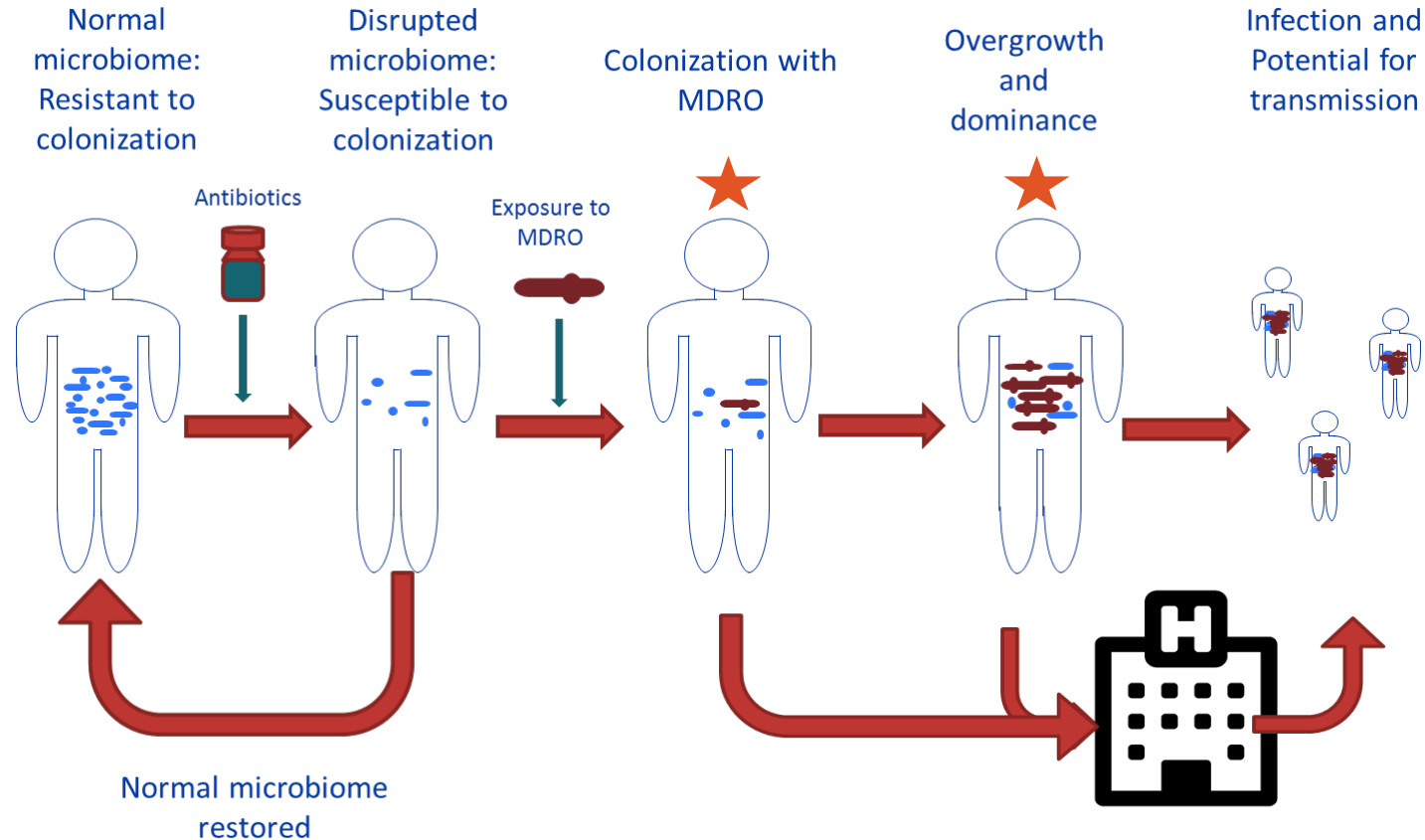
# Causal Pathway from Health to Disease



# Causal Pathway from Health to Disease



# Causal Pathway from Health to Disease



# Potential MDI

- Potential MDIs
  - Compositional diversity
  - Species richness
  - Presence/absence of protective species
  - Resistome
  - Functional status – metagenomics, metabolomics

# Uses for MDIs

## ■ Monitor microbiome

- When to take remedial action (e.g., fecal microbiota transplant)
- Antibiotic stewardship/selection
- Infection control decisions
  - Isolation precautions for when at increased risk for transmitting (colonized, dominated)
  - Reverse isolation precautions for those at high risk for colonization (disrupted)





## Uses for MDIs

- **Prepare for future FDA-approved microbiome remediation therapies**
  - e.g., FMT, synthetic stool, advanced/designer probiotics
- **Communicate disruptive potential of drugs, including antibiotics**
  - Rating system to gauge relative risks of different agents
  - MDIs determined during approval process and included in package insert

## Wrap Up

- Using WGS has already had a substantial impact on our ability to investigate outbreaks
- Collect timely and accurate epi data is crucial
- Interpret WGS in the context of available epi data

# Thank you

For more information, contact CDC  
1-800-CDC-INFO (232-4636)  
TTY: 1-888-232-6348 [www.cdc.gov](http://www.cdc.gov)

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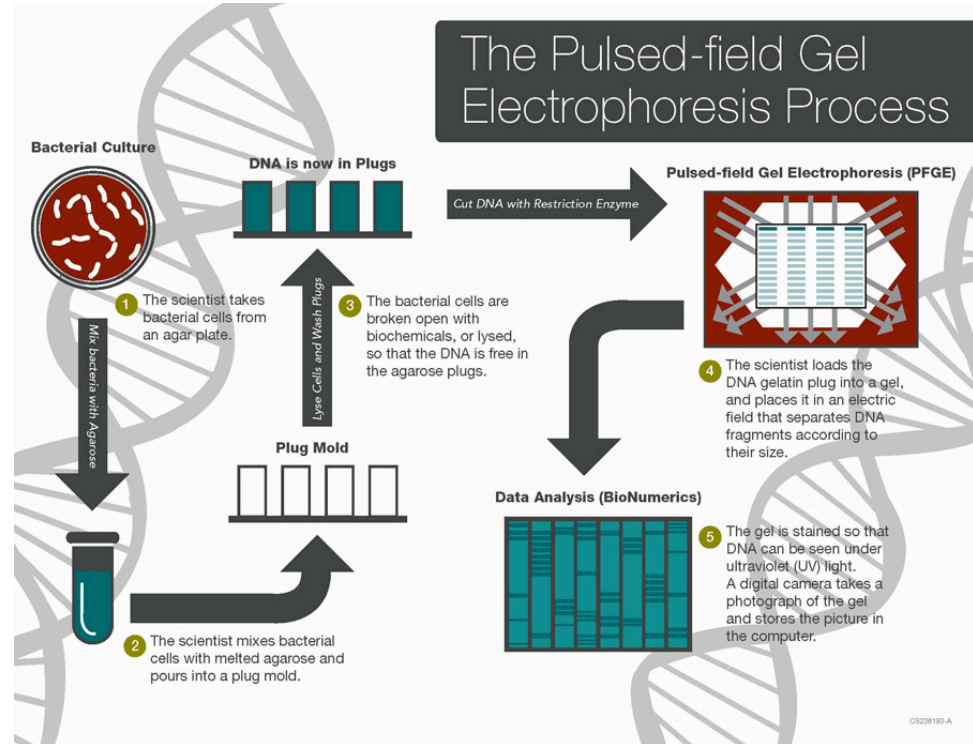


# Antibiotic Resistance Threat Report

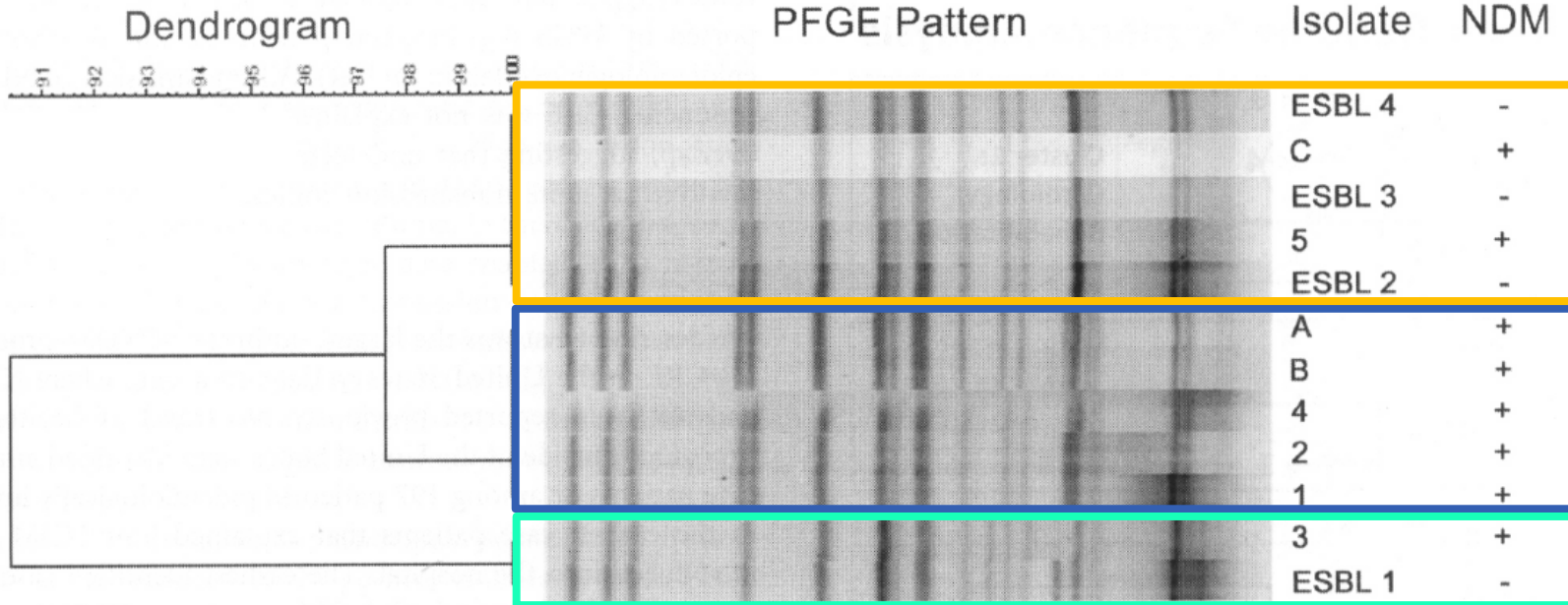
- 18 antibiotic resistant pathogens
  - >2 million infections
  - ~23,000 deaths
  - Billions in excess medical costs
- Half are healthcare-associated infections
- Vital to continue progress in healthcare epidemiology and implementation research

# Pulsed-Field Gel Electrophoresis (PFGE): Strengths

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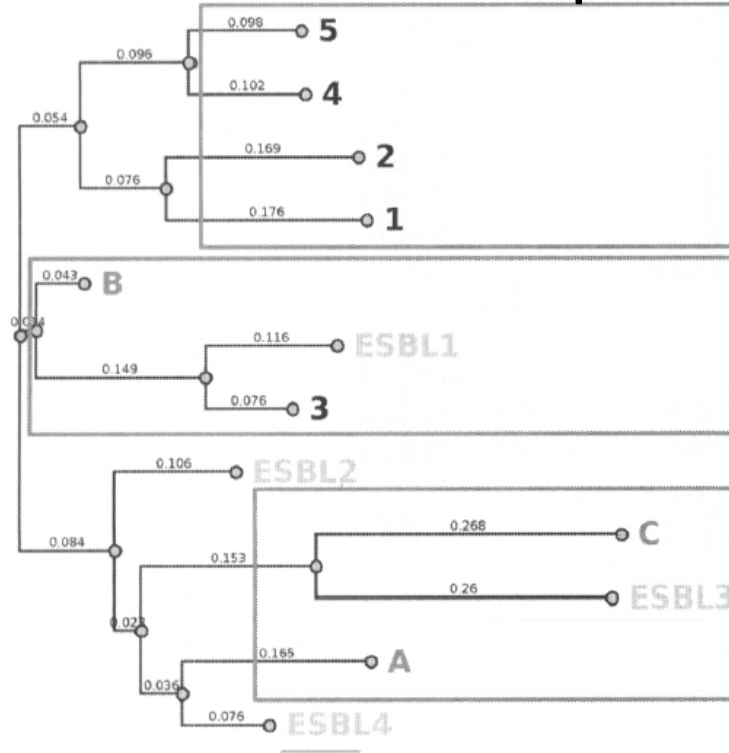
# Denver, Colorado - 2012





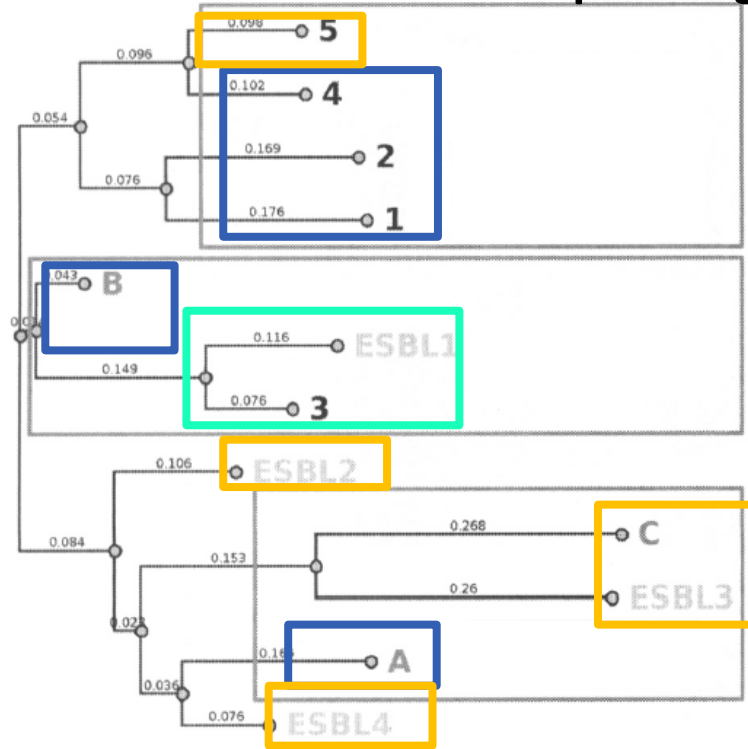
# Denver, Colorado - 2012

## Whole Genome Sequencing



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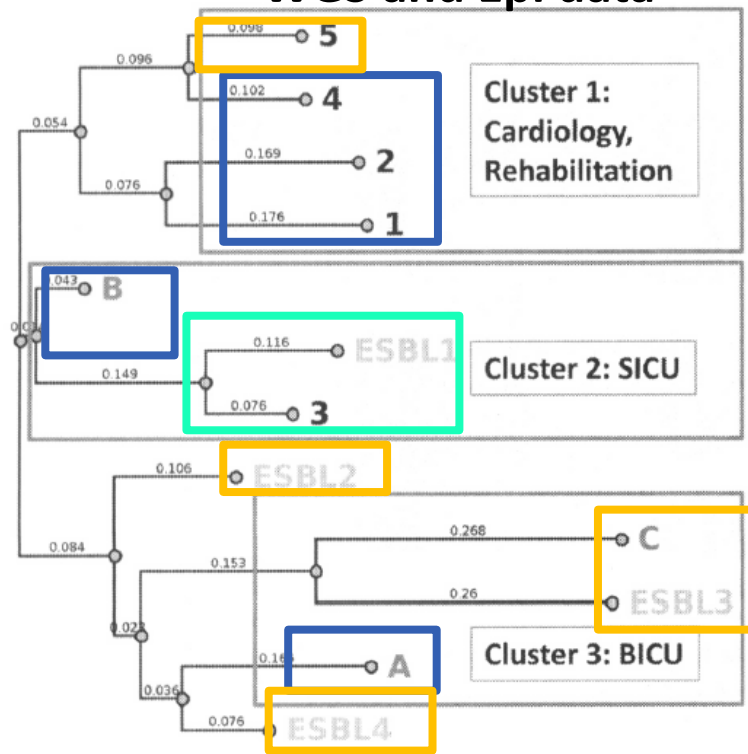
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