



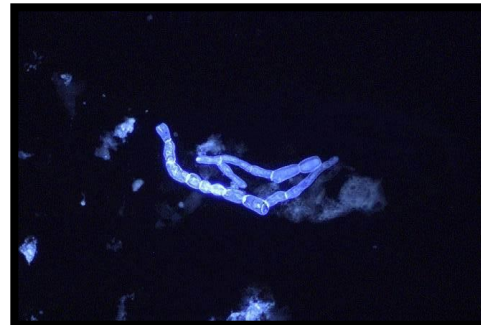
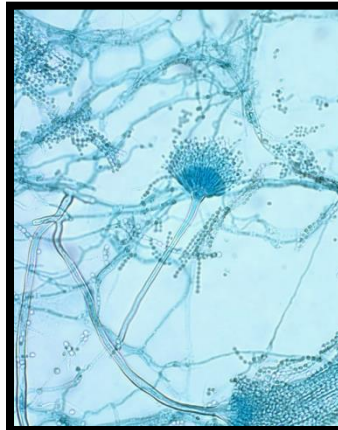
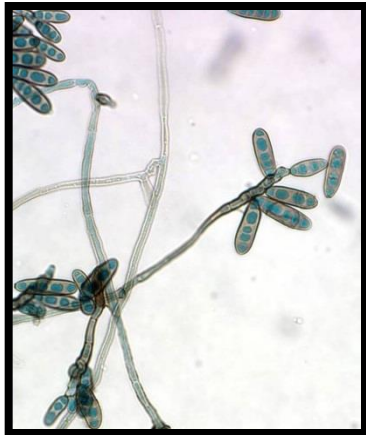
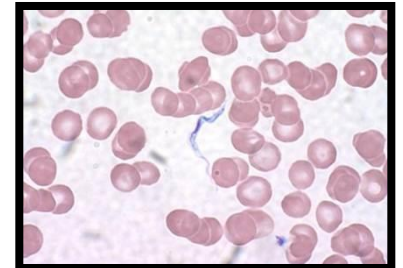
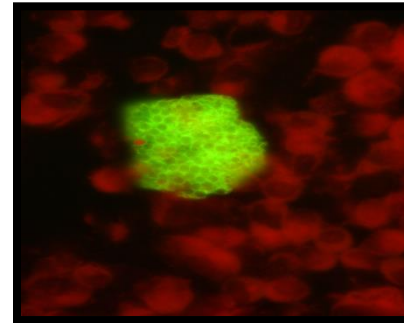
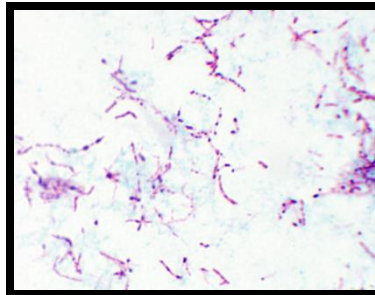
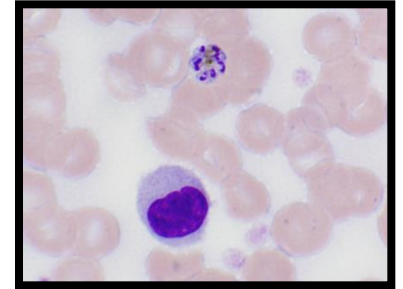
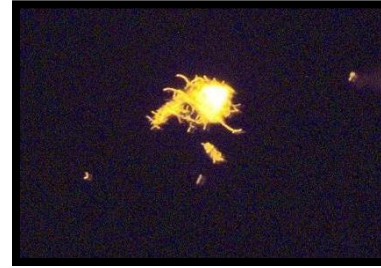
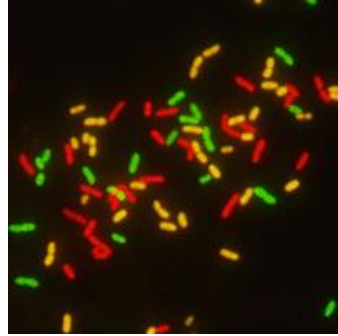
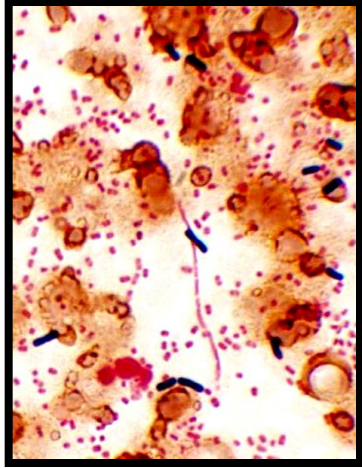
# Molecular Diagnostics: The Future of Clinical Microbiology

**Patrick R. Murray, PhD**  
**WW Director, Scientific Affairs**  
**BD Diagnostic Systems**

# Clinical Diagnostics

- Microscopy
- Culture
- Antigen Detection
- Antibody Detection
- Molecular Diagnostics

# Microscopy



# Microscopy

## **Techniques:**

- Contrasting stains
- Differential stains
- Immunofluorescent stains

## **Advantages:**

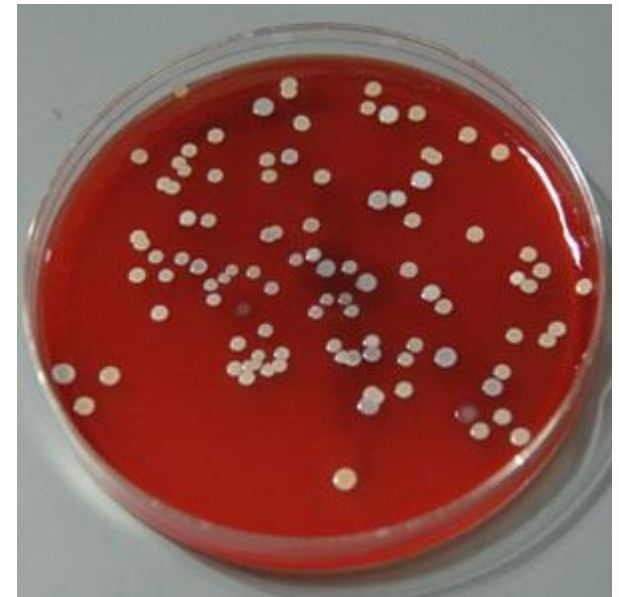
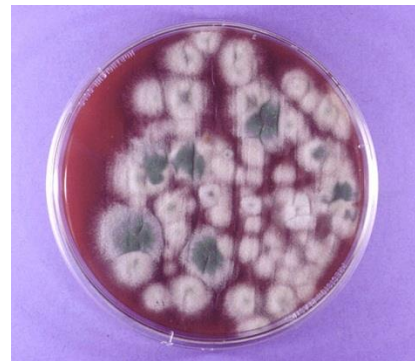
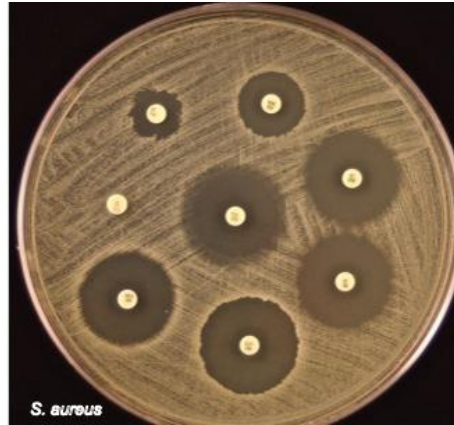
- Rapid assessment of specimens and associated pathogens
- Diagnostic test of choice for certain organism groups
- Specific identification of key pathogens

## **Disadvantages:**

- Relatively insensitive
- Subjective interpretation



# Culture



# Culture

## **Techniques:**

- Use of enriched, differential, and selective culture media, as well as tissue culture cell monolayers
- Used for all major groups of bacteria, fungi and viruses, as well as some parasites

## **Advantages:**

- Isolation of pathogens that can provide a definitive diagnosis
- Defines relative proportion of pathogenic and nonpathogenic organisms
- Organism available for ASTs

## **Disadvantages:**

- Slow time to results
- Only able to detect organisms that grow in culture

# Antigen Tests

- Antigen tests used for diagnosis of bacterial, fungal, viral and parasitic infections
- Useful for rapid results
- Inexpensive POC tests
- Not technically demanding and high volume tests performed on automated platforms
- Analytical performance (sensitivity, specificity, predictive value) improving but still inferior to culture and molecular diagnostics
- Provides clinical value but frequently must be confirmed by alternative diagnostic tests



# Antibody Tests

- Primarily used for screening patients for immunity to specific diseases or past exposure to pathogens
- Antibody tests used less commonly for diagnosis of acute infections; requires demonstration of elevated antibody levels or change in levels
- Improvements in culture techniques and introduction of molecular testing have decreased the value of serology for diagnosis of active infections





# Molecular Diagnostics: Historical

- Commercial assays performed on large platforms
- High volume assays, such as blood screening for viral agents and STD testing
- Microbiology lab developed tests were primarily restricted to research labs or academic settings
- Testing generally was expensive and required specialized technical expertise and testing facilities

# Molecular Diagnostics

## Dedicated Nucleic Acid Extraction



**bioMérieux easyMAG**



**Roche MagNA Pure LC**



**Hamilton Microlab Starlet**



**Qiagen  
QiaSymphony**

## Dedicated Real-time PCR



**Roche LightCycler**



**Qiagen  
Rotor-Gene**

# Integrated Extraction & Amplification Systems

## BD Viper XTR



- Closed Systems
- Floor standing

## Hologic/GenProbe

### Panther



- Closed Systems
- Floor standing

### Tigris



# Integrated Extraction & Amplification Systems

## BioFire Film Array



## Nanosphere Verigene



## Cepheid GeneXpert



## BD MAX™ System





# Integrated Extraction & Amplification Systems

## BD MAX™ System

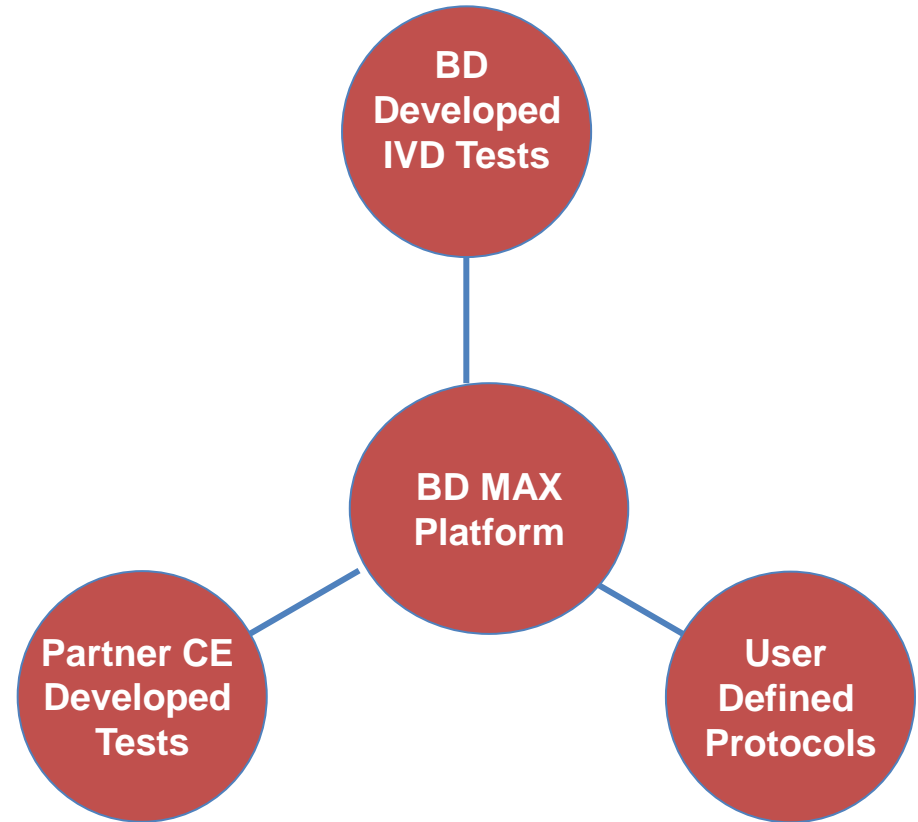


- **Broad IVD and LDT menu**
- **Open system**



# A New Model for Molecular Assay Development

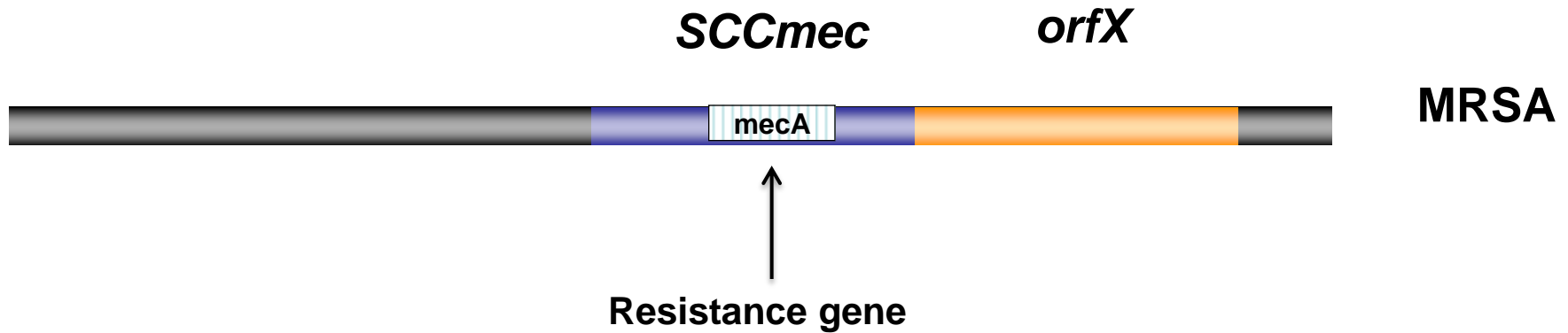
- **BD R&D**
  - 6 teams of 8-10 scientists currently work on assay development
  - 1 team of 6-8 support open system assay development
- **Commercial partners**
  - Diagenode
  - BioGX
  - Others
- **Clinical partners**
  - EU MAX Expert User group
  - US MAX Expert User group
  - Research groups



# MRSA: Making a Good Assay Better



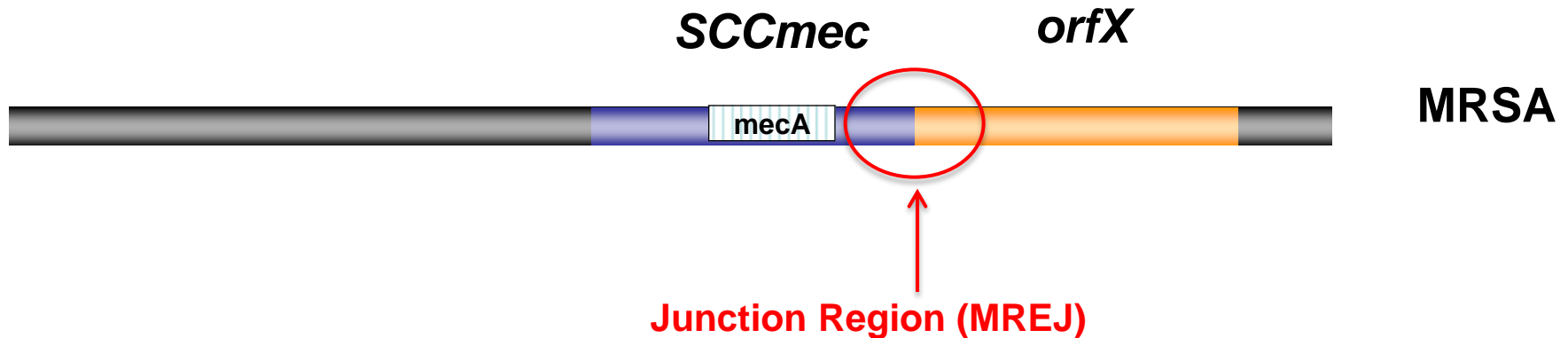
# MRSA: *mecA* Resistance Gene



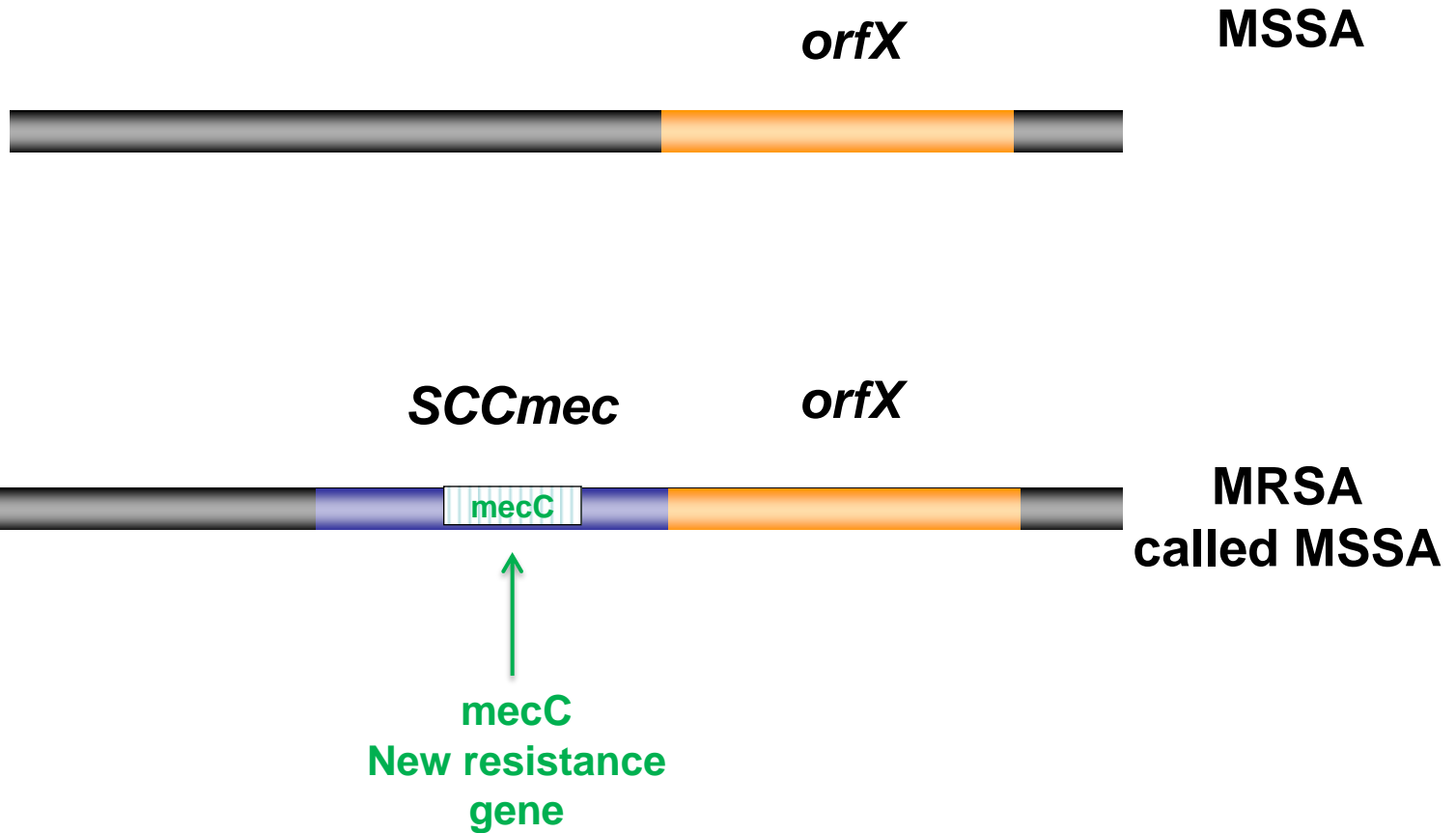


# MRSA: mecA Resistance Gene

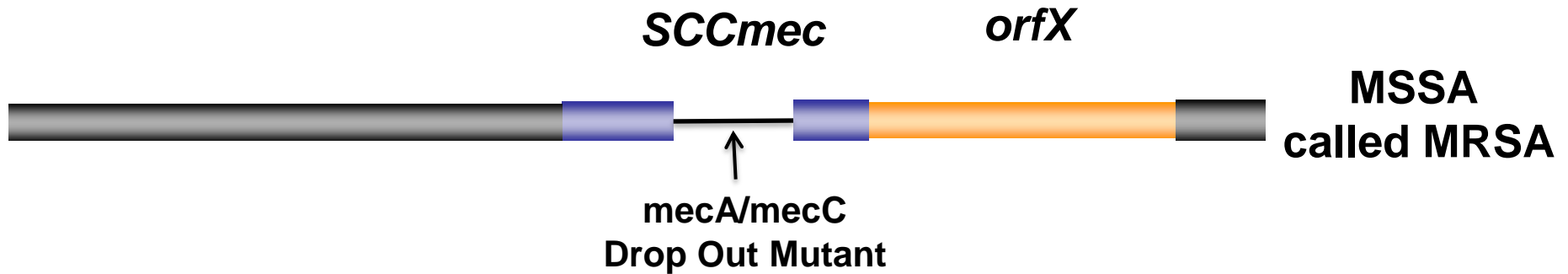
## MRSA: Junction Region Target



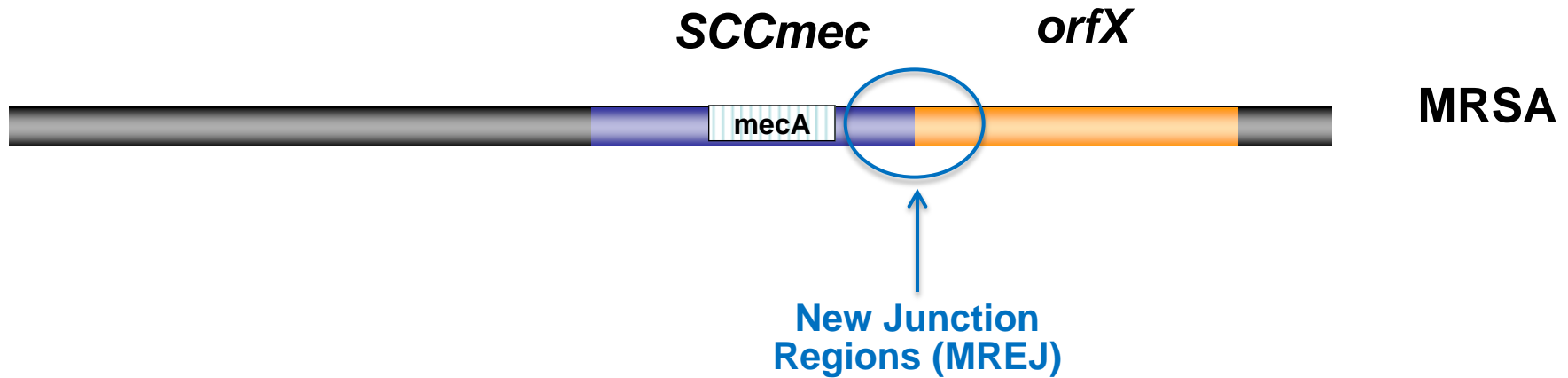
# MRSA: New Resistance Genes



# MRSA: Drop Out Mutants



# MRSA: New Junction Region Targets





# Next Generation MAX StaphSR and MRSA-XT will detect a broader range of MRSA variant strains

## MREJ Types detected

Current MAX MRSA

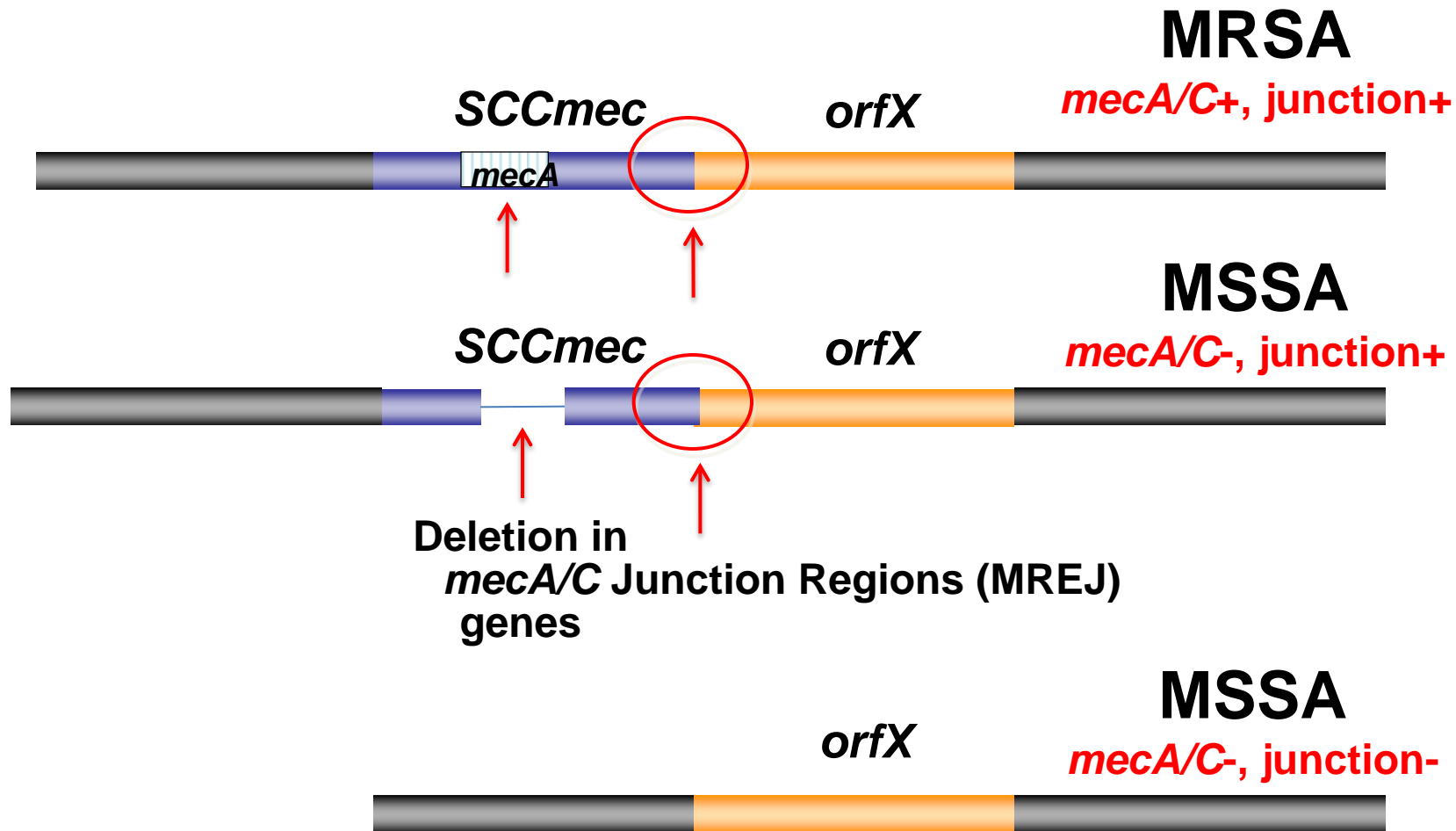
i ii iii iv v vii

Next Gen MAX StaphSR  
and MAX MRSA-XT

i ii iii iv v **vi** vii **ix** **xiii** **xiv** **xxi**

- Detects both *mecA* and *mecC* genes
- Detects *mecA* and *mecC* dropout mutants
- Detects newly discovered MREJ strains

# New MAX MRSA XT and StaphSR Assays Are More Sensitive and Specific



# Development of Syndromic Menu

- Clinical approach to menu development is – develop comprehensive syndromic solutions
- Enteric diseases
  - Clostridium difficile (CE/IVD)
  - Bacterial panel – Salmonella, Shigella, Campylobacter, STEC (CE/IVD)
  - Viral panel – Norovirus, Rotavirus (CE)
  - Parasites – Cryptosporidium, Giardia, E. histolytica (CE)
  - Uncommon enteric bacterial pathogens
- Assay development
  - Detect pathogens not previously assayed
  - Improve detection over existing assay methods



# Clostridium difficile Diagnostic Tests

Diagnostic Method	Target	Sensitivity (%)	Specificity (%)	Turnaround Time
Cell cytotoxicity culture assay	Toxin	70-80	90-95	Days
Glutamate dehydrogenase	Common antigen	70-90	<90	Hours
Enzyme immunoassay	Toxin	40-70	>97	Hours
Toxigenic culture	Toxin	>95	95-97	Days
PCR	Toxin	>95	>97	Hours



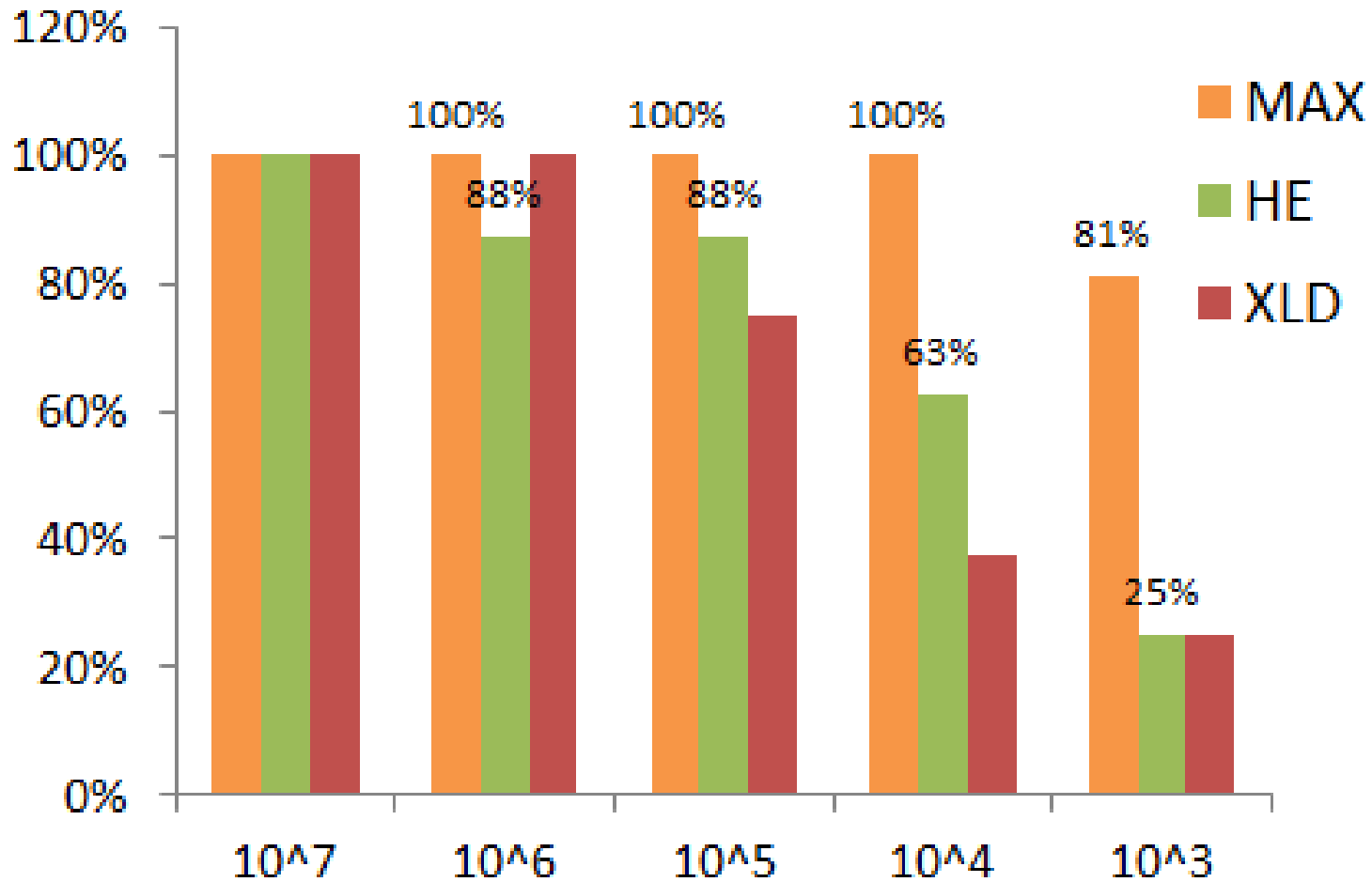
# BD MAX Limit of Detection

## Salmonella spp.



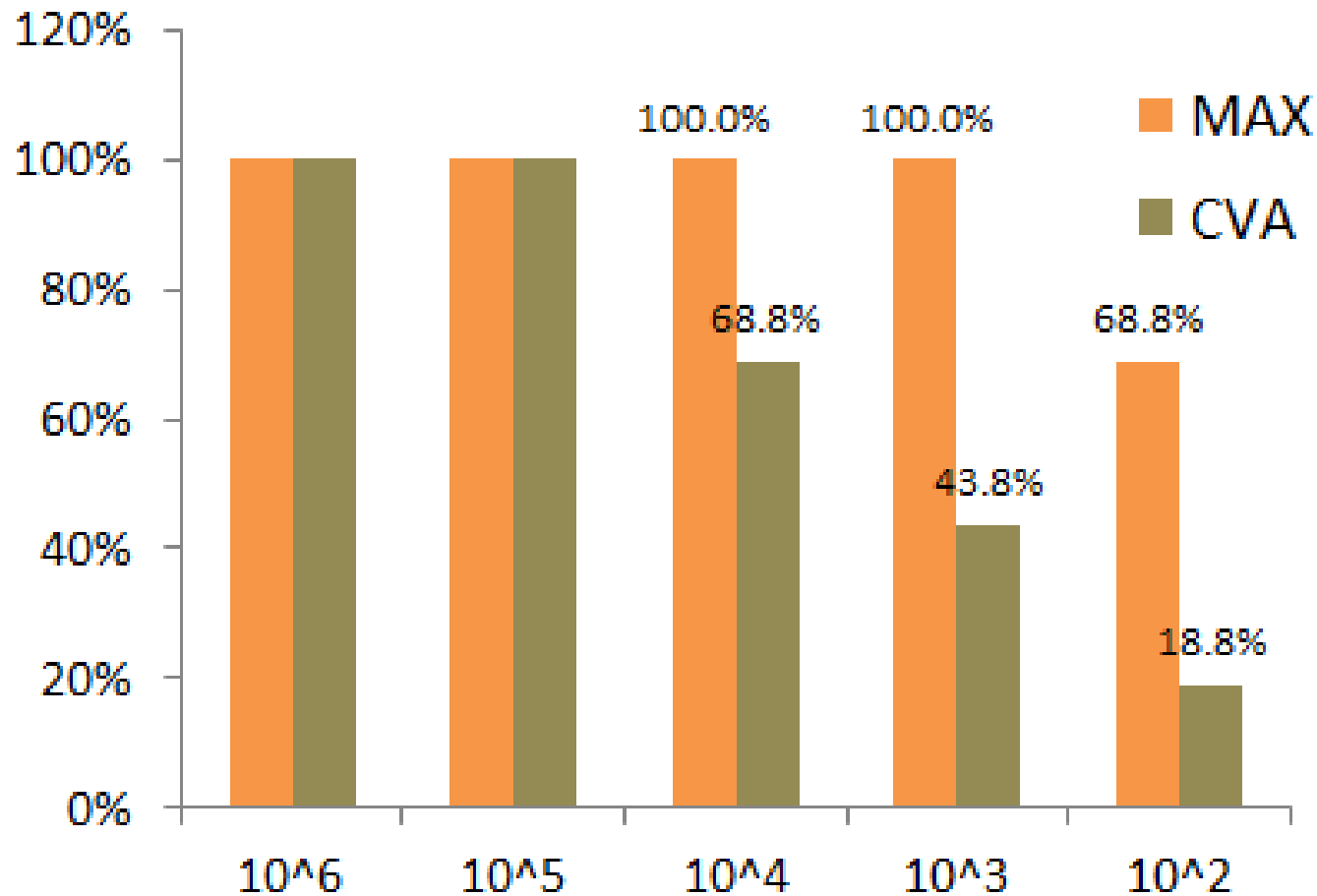
# BD MAX Limit of Detection

## Shigella spp.



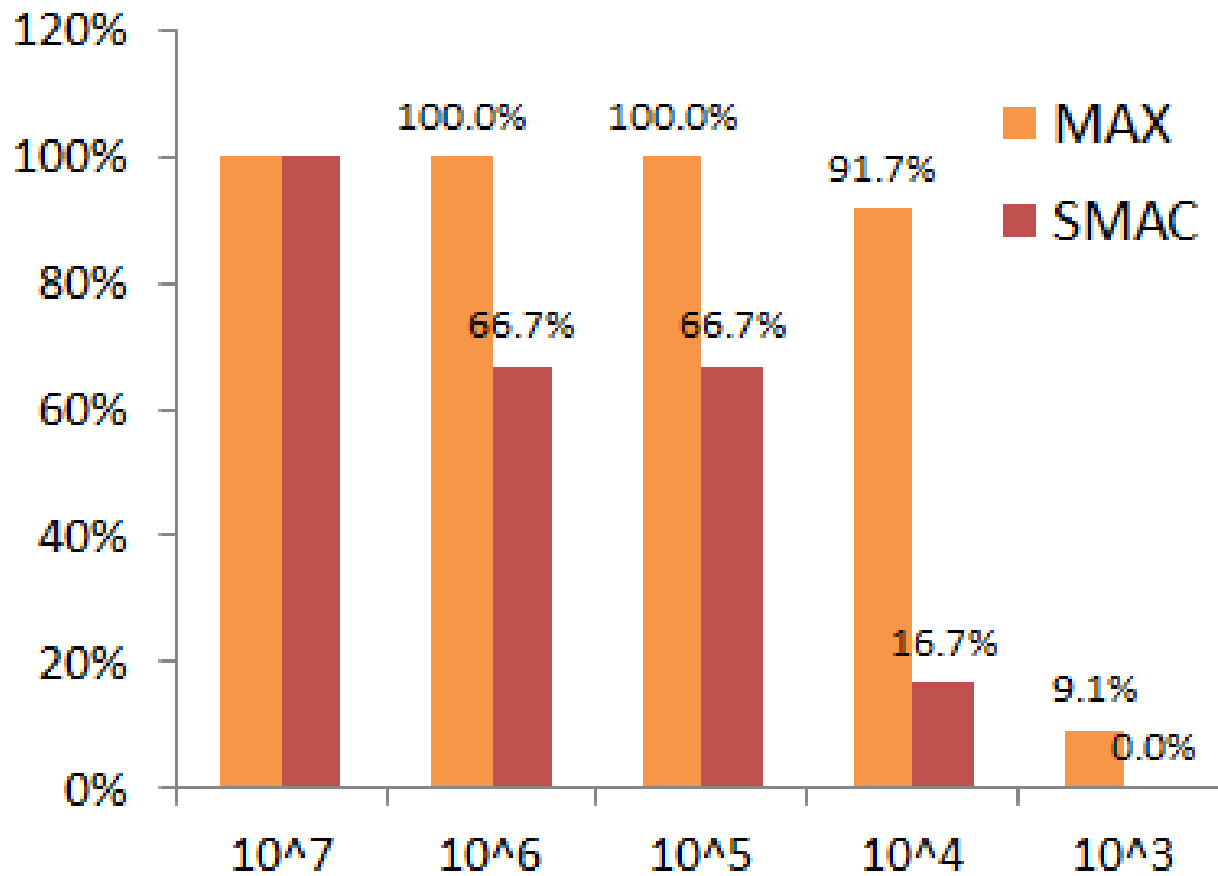
# BD MAX Limit of Detection

## Campylobacter spp.



# BD MAX Limit of Detection

## EHEC (0157)



# Global Medical Problems

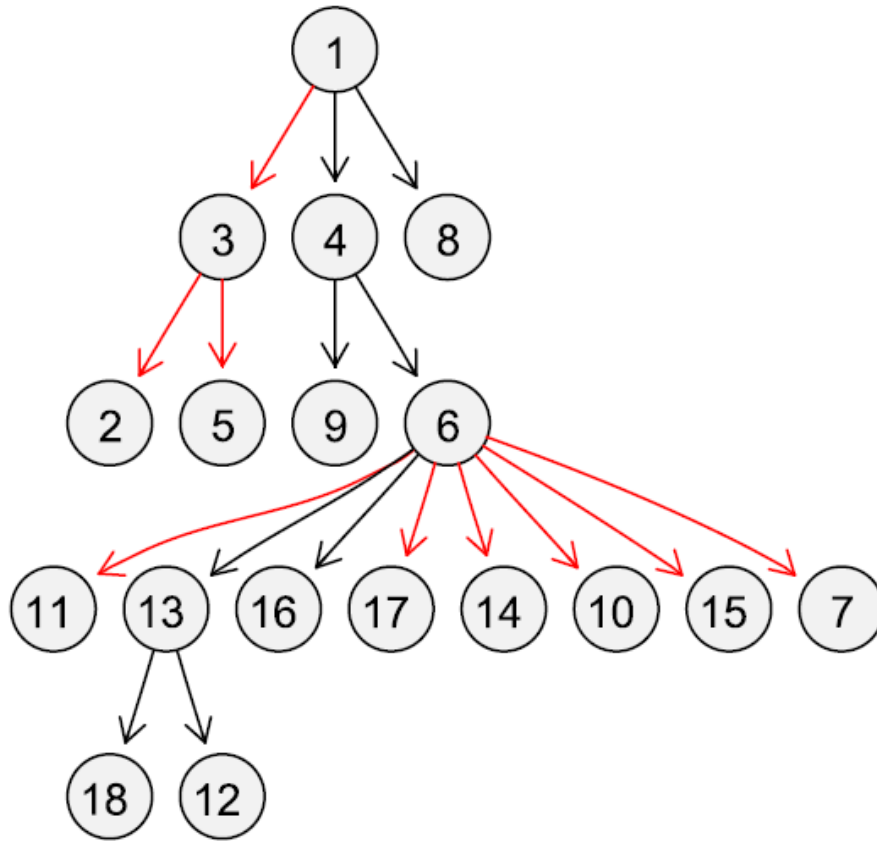
Problem
Outbreaks with carbapenem-resistant bacteria
Bacterial meningitis outbreak in Chile
Coronavirus outbreak in Middle East



# 2011 Outbreak with Multidrug Resistant Gram-Negative Bacteria at the NIH Clinical Center

- On June 13 a patient from NYC was admitted into the NIH Clinical center ICU. She was colonized with a MDR *A. baumannii* and a KPC strain and was immediately placed on enhanced contact isolation in a private room. She was discharged 1 month later.
- On August 5 a second patient was found to be infected with the same strain of KPC.
- A total of 18 patients became infected, with 11 deaths including 7 patients whose deaths were directly attributed to this organism.
- This outbreak illustrates the difficulty in controlling, by traditional infection control practices, essentially untreatable multidrug resistant gram-negative bacteria.

# Tracking a Hospital Outbreak



- Traditional epidemiology approaches underestimated transmission
- Successful intervention required a combination of comprehensive screening and rigorous infections control practices.

# Carbapenemase Assay

Developer	Assay Targets
BD R&D	KPC, NDM-1, Oxa-48
Dalpke et al	Panel 1: KPC; VIM-2 IMP-1,-2; GES Panel 2: NDM; OXA-23,-48; VIM-1
CheckPoint	KPC; NDM; Oxa-48; VIM
BioGX	KPC; NDM; OXA-48; VIM/IMP

## 2012-13 *Neisseria meningitidis* Outbreak in Chile

- On Sunday Dec 16 a woman traveling on LAN flight from Santiago, Chile to Punta Cana, Dominican Republic developed symptoms of meningitis and died of an overwhelming infection with *Neisseria meningitidis*. All 200 passengers on this flight were given prophylactic antibiotics.
- By April 2013, 178 cases of meningococcal meningitis were reported in Chile.
- Most infections were with serogroup W135 and were localized in the Metropolitan Region (includes Santiago) and the adjoining Valparaíso Region.



# BD MAX™ Open System Group (OSG)

- A multiplex bacterial meningitis assay for *N. meningitidis*, *S. pneumoniae*, and *H. influenzae* was developed in 1 month
- Based in Baltimore MD R&D Systems Engineering:
  - Composed of Manager, Senior Scientists and Engineers
- Objective: Provide advanced applications support to customers and MAX partners
- Offers scientist-scientist collaboration for customers developing LDTs, performs technical feasibility testing in-house

## 2012-13 New Coronavirus Outbreak in Middle East

- Between April 2012 and May 2013, 49 cases of human infection with a new coronavirus originating in the Middle East (Middle East Respiratory Syndrome Coronavirus; MERS-CoV) were reported with 26 deaths (53%) due to severe acute respiratory disease.
- Cases were reported from Saudi Arabia (37), Jordan (2), as well as the UK (4), France (2), Tunisia (2) and Germany (2).
- MERS-CoV is related to bat viruses; however, no animal reservoir has been identified. Until recently, limited person-to-person spread was documented in close contacts and healthcare personnel.
- Most coronavirus infections are mild respiratory disease but MERS-CoV and SARS-CoV cause severe acute respiratory syndromes with an associated high mortality.



# MERS-CoV Assay

- Developed by clinical partners (Hendrickson and Bose, Medical College of Wisconsin)
- Primers, probes, and plasmid control designed for RNA-dependent RNA polymerase gene using 3 published genome sequences for MERS-CoV.
- Clinical testing performed with 20 spiked positive samples and 56 negative clinical samples. Assay with 100% sensitivity and specificity.
- Development and validation time: 2 weeks

# Global Problems – Molecular Solutions

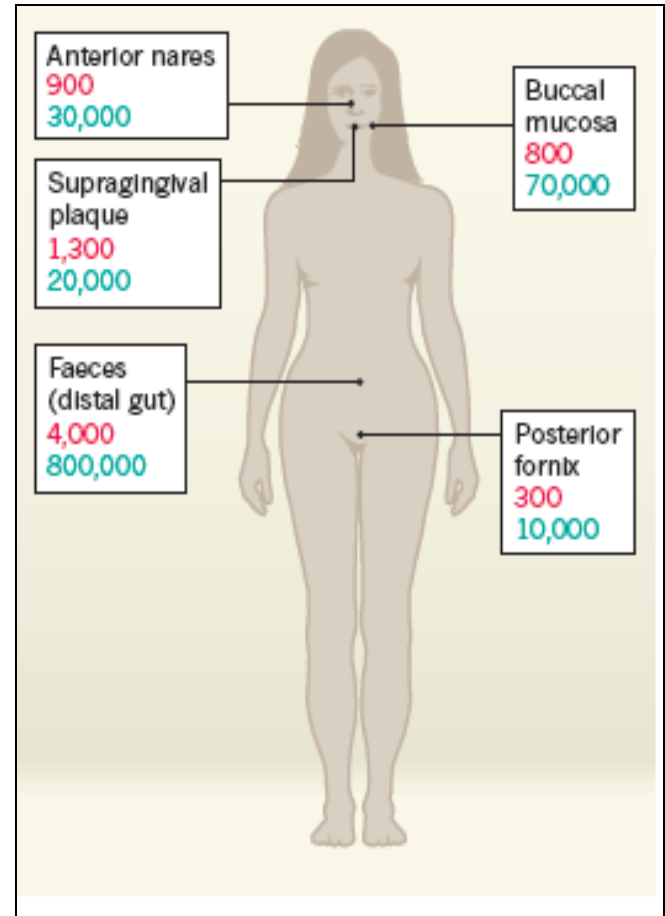
Problem	Solution
Outbreaks with carbapenem-resistant bacteria	Multiplex assay for KPC, NDM-1, Oxa-48
Bacterial meningitis outbreak in Chile	Multiplex assay for <i>N. meningitidis</i> , <i>S. pneumoniae</i> , <i>H. influenzae</i>
Coronavirus outbreak in Middle East	Screening assay and confirmatory assay for MERS-CoV

# Summary

- BD MAX, BioFire Film Array, Cepheid GeneXpert, Nanosphere Verigene and Hologic-GenProbe Panther and Tigris are examples of integrated systems of nucleic acid extraction, amplification, and analysis.
- BD MAX is a unique open system for simultaneously performing multiplex assays developed by BD, commercial partners, and clinical partners.
- The ability to develop novel assays (such as for the diagnosis of multidrug-resistant bacteria, bacterial meningitis, MERS-CoV) enables diagnostic labs to respond rapidly to emerging infectious diseases

# Human Microbiome

- Human Genome Project (1990-2003) – 13 year project to map the 22,000 genes in the human genome
- Human Microbiome Project (2005-2010) – 5 year project to map the 8 million unique genes in the bacteria that populate our bodies
- Most bacteria on the body are critical for our health (“bacteria are good not bad”)
- Although there is **taxonomic heterogeneity** (most of us have a unique population of bacteria), there is **functional redundancy** (we share common gene functions)



# Microbiome and Intestinal Health and Disease

- The intestinal microbiome is characteristically diverse and functions in close balance to:
  - protect against colonization with pathogens
  - detoxify potential poisons
  - produce energy and nutrients by digestion of food
  - maintain mucosal and systemic immunity
- **Clostridium difficile diarrhea** – under the influence of antibiotics or disease, the microbiome shifts from a complex flora to a predominance of toxin-producing *C. difficile*.
- **Crohn's disease** and **inflammatory bowel disease** – distinct, characteristic changes of microbiota that alter regional metabolism and mucosal immunity.

# What Impact Will Microbiome Research Have on the Clinical Microbiology Laboratory?

- **Antibiotic susceptibility testing:** genome sequencing can identify known resistance genes but cannot determine expression or previously unrecognized resistance markers.
- **Microbial identification:** sequencing is unlikely to replace mass spectrometry for organism identification but can identify the composition of a population of organisms, virulence potential of microbes, and be used as the definitive typing method.
- **New approach to laboratory diagnostics:** laboratory testing could shift from diagnosis of disease to prediction of disease by monitoring shifts in the microbiome species or metabolic function.
- **Technical challenges:** preanalytical processing, sequencing, and data analysis are daunting but success will revolutionize diagnostics.





# Development of Syndromic Menu

- Syndromic menu
  - Comprehensive multiplex assay
  - Targeted multiplex assays
- Comprehensive assays
  - Broad menu of analytes – but
  - Expensive
  - Decreased analytical performance
- Targeted assays
  - Requires clinical assessment of patient and development of differential diagnosis
  - Faster assay development and regulatory clearance