

# Everything in Sequence: Listeriosis Outbreak Investigations in the Era of WGS

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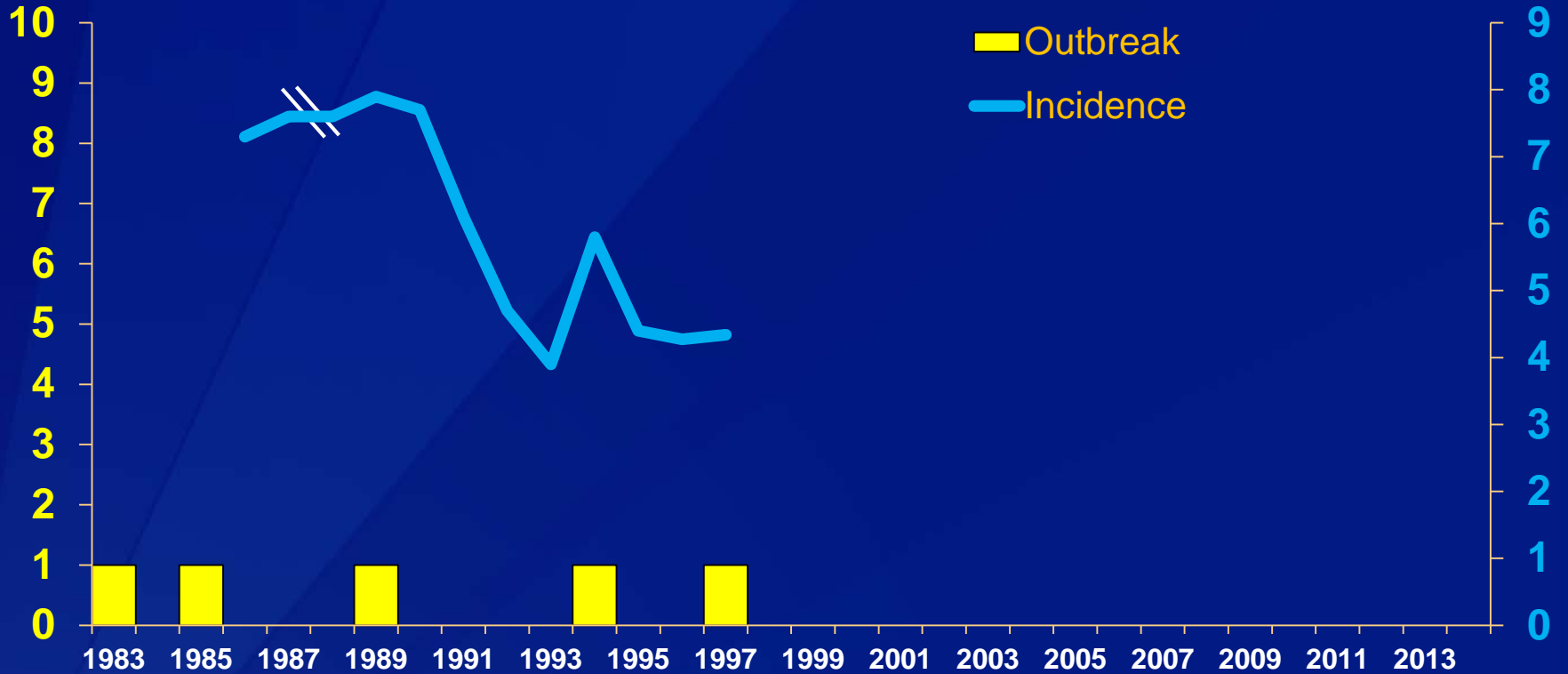
## Disclaimers

- *The findings and conclusions in this presentation are those of the authors and do not necessarily represent the official position of the Centers for Disease Control and Prevention.*
- *No financial conflicts of interest to report.*

# Listeria Outbreaks and Incidence, 1983-2014

No. outbreaks

Incidence  
(per million pop)



Era

Pre-PulseNet

Outbreaks per year

0.3

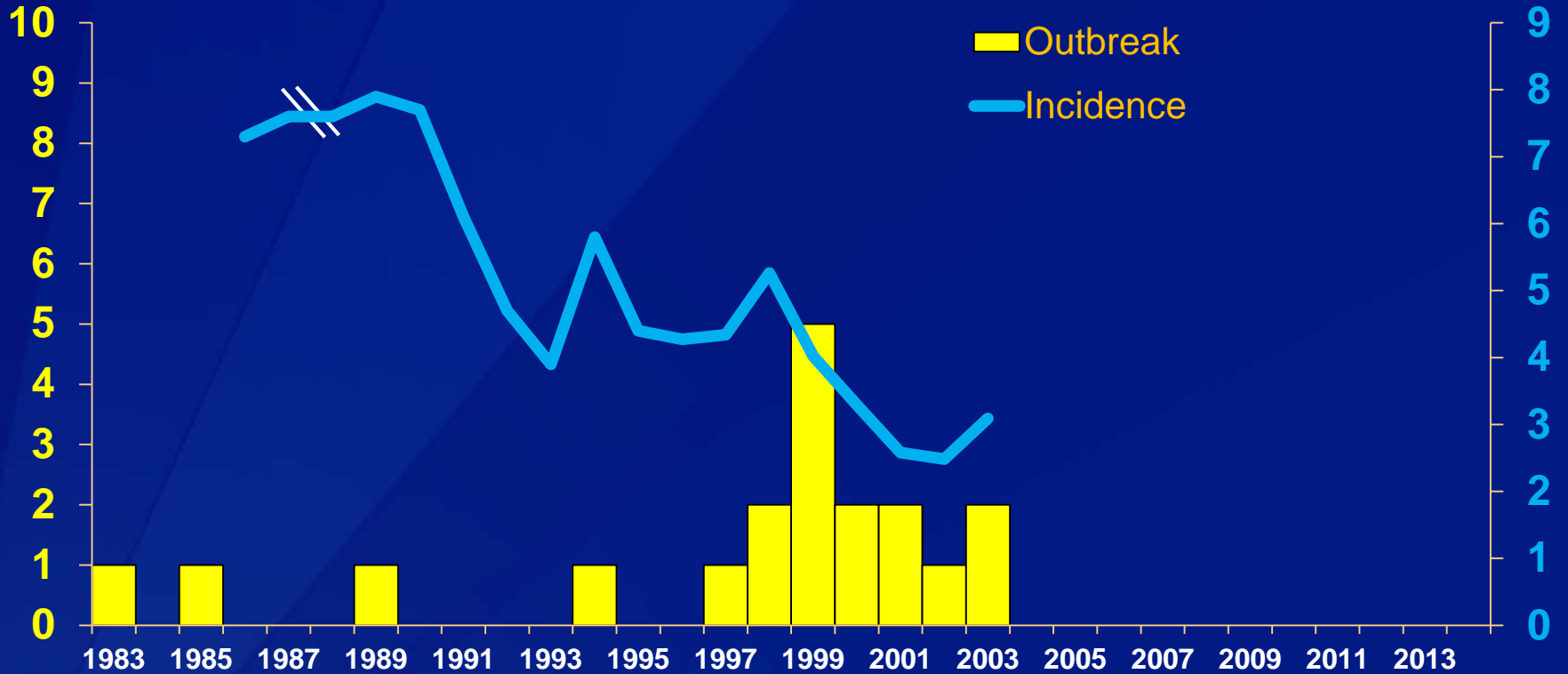
Median cases per outbreak

69

# Listeria Outbreaks and Incidence, 1983-2014

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Incidence  
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Era

Pre-PulseNet

Early PulseNet

Outbreaks per year

0.3

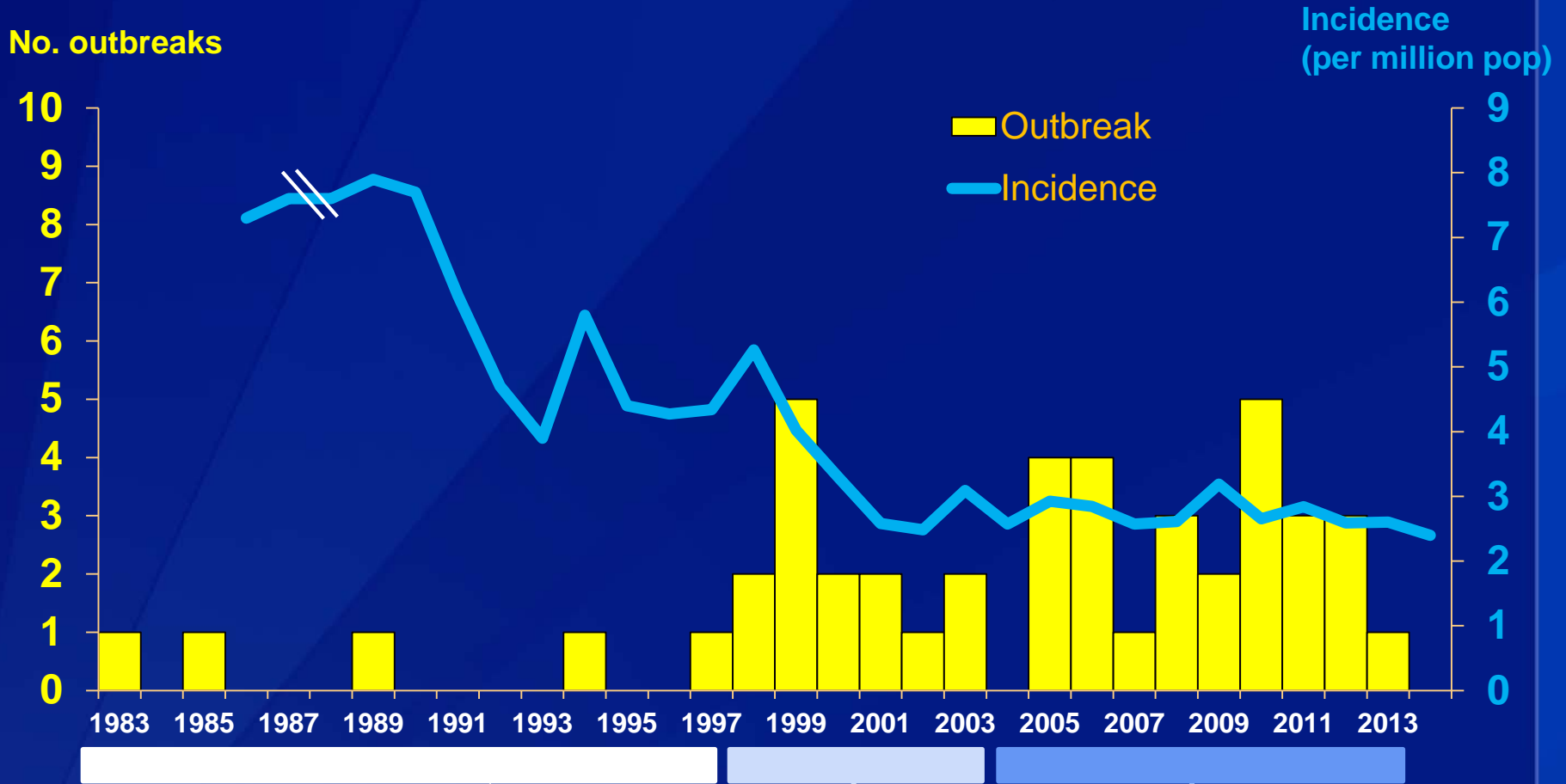
2.3

Median cases per outbreak

69

11

# Listeria Outbreaks and Incidence, 1983-2014



<u>Era</u>	<u>Pre-PulseNet</u>	<u>Early PulseNet</u>	<u>Listeria Initiative</u>
Outbreaks per year	0.3	2.3	2.6
Median cases per outbreak	69	11	5.5

# Listeria WGS Pilot Project

- ❑ Started September 2013
- ❑ Goal: Sequence all *Listeria monocytogenes* isolates
- ❑ Near real-time (<1 week for patient isolates)



Public Health Agency of Canada



# Listeria WGS Analysis Methods

- ❑ Kmer
- ❑ High-quality single nucleotide polymorphism (hqSNP)
- ❑ Whole-genome multilocus sequence typing (wgMLST)

The screenshot shows the NCBI BioSample database entry for the sample SAMN02950476. The search results are displayed for the sample ID PNUSAL000870. The pathogen is identified as *Listeria monocytogenes*. The entry includes various attributes such as strain, collection date, isolation source, host, geographic location, and submission information.

NCBI Resources How To

BioSample BioSample PNUSAL000870  
Save search Advanced

Display Settings: Full

**Pathogen: clinical or host-associated sample from *Listeria monocytogenes***

Identifiers BioSample: SAMN02950476; Sample name: PNUSAL000870; SRA: SRS672874

Organism [Listeria monocytogenes](#)  
cellular organisms; Bacteria; Firmicutes; Bacilli; Bacillales; Listeriaceae; Listeria

Package Pathogen: clinical or host-associated; version 1.0

Attributes

strain	PNUSAL000870
collected by	CDC
collection date	missing
isolation source	missing
host	missing
geographic location	<a href="#">USA</a>
latitude and longitude	missing
host disease	missing

Submission CDC, Heather Carleton; 2014-08-01

ID: 2950476  
[BioProject](#) [SRA](#)

# Relationship Between SNPs and wgMLST Alleles

SNPs

gene 1

gene 2

gene 3

wgMLST  
allele  
differences

1



1



# Relationship Between SNPs and wgMLST Alleles

SNPs

gene 1

gene 2

gene 3

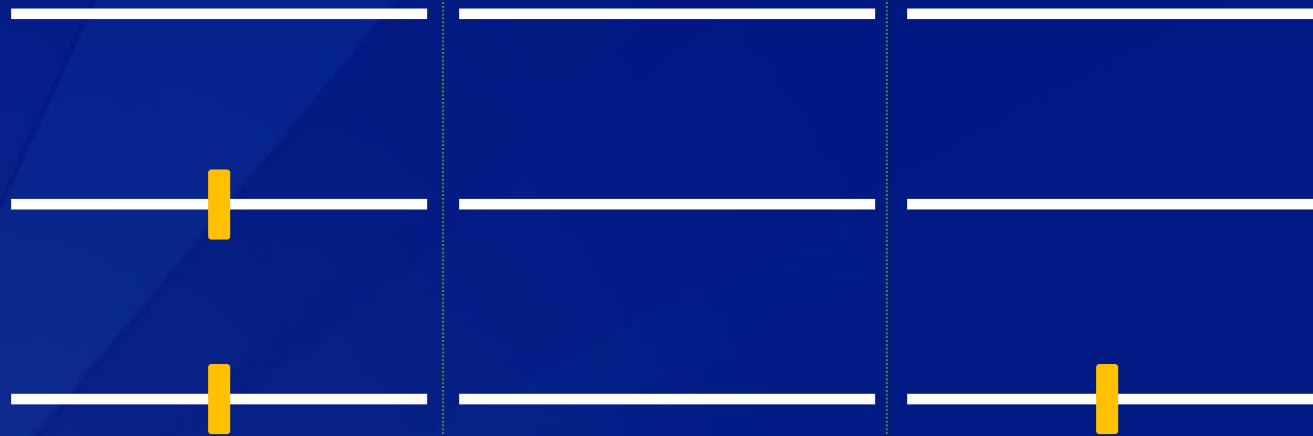
wgMLST  
allele  
differences

1

1

2

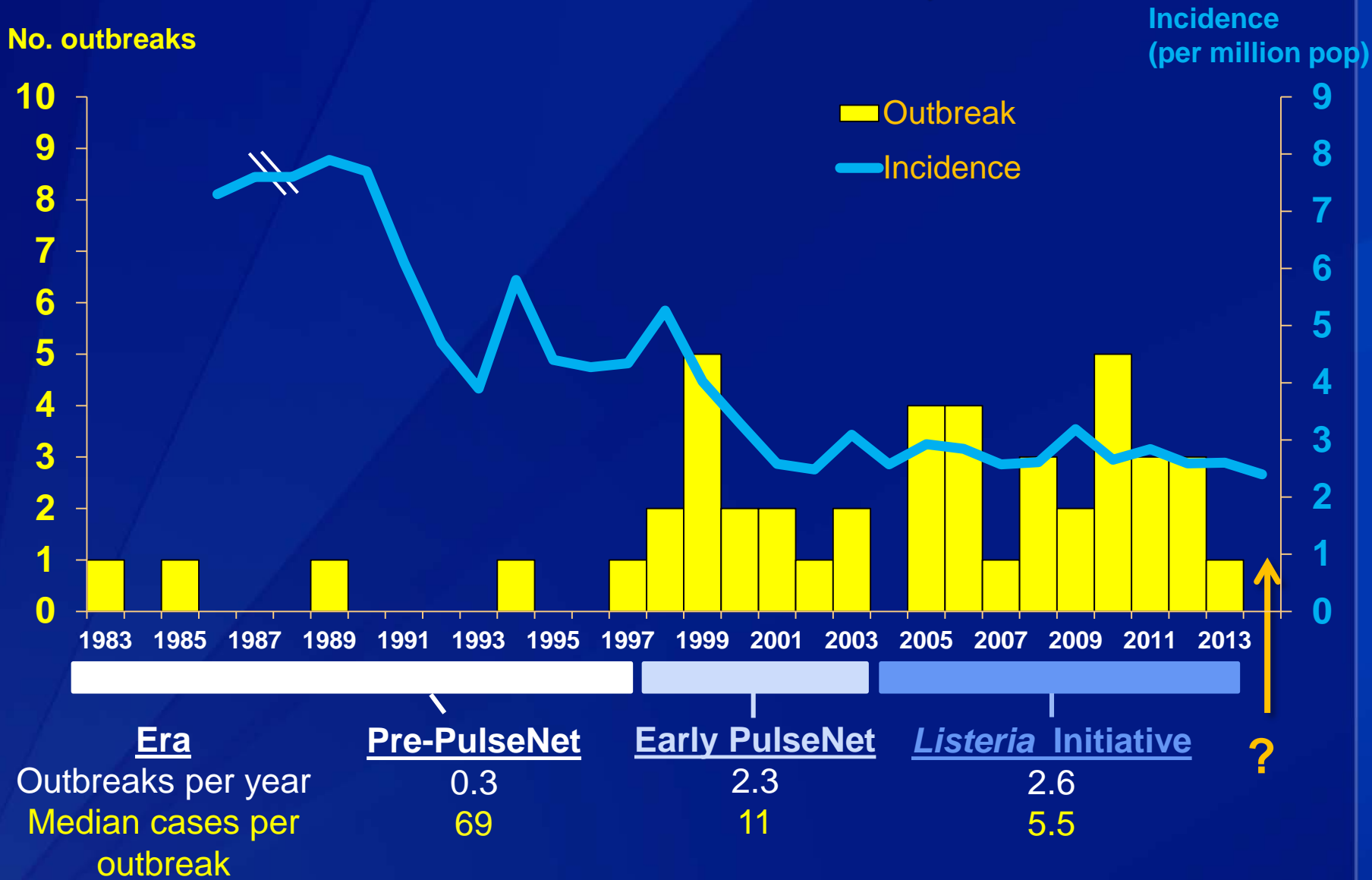
2



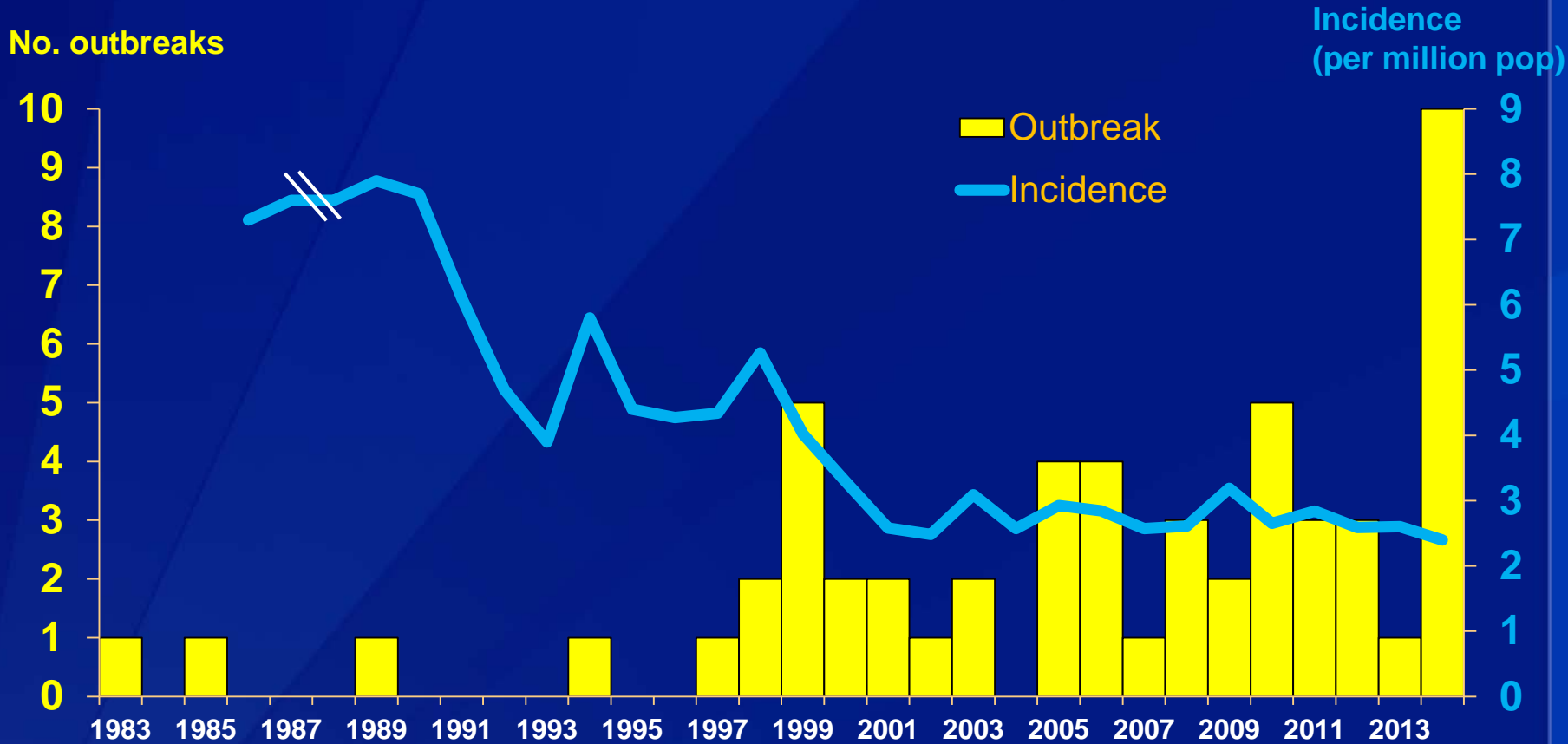
# Relationship Between SNPs and wgMLST Alleles



# Listeria Outbreaks and Incidence, 1983-2014



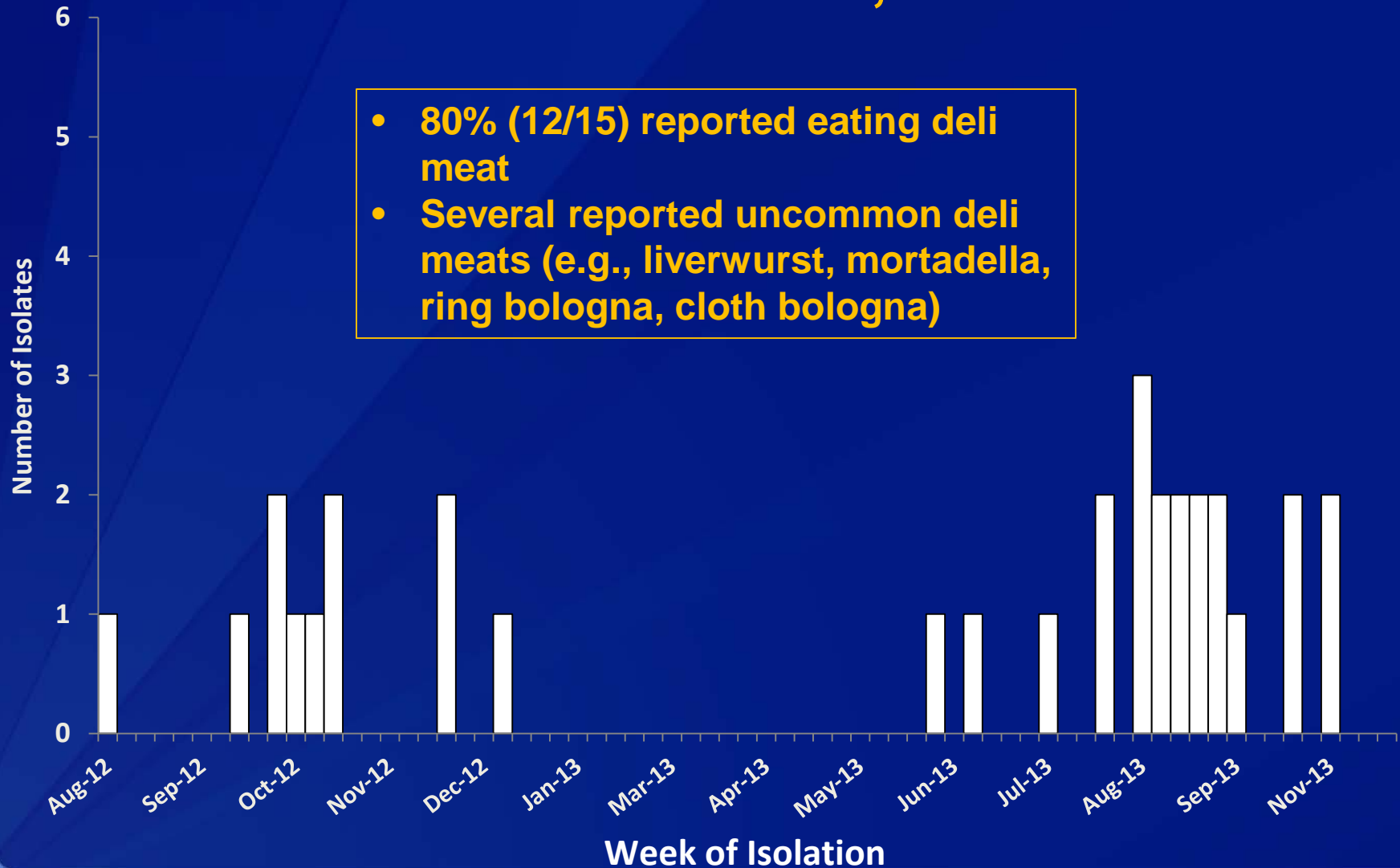
# Listeria Outbreaks and Incidence, 1983-2014



Era	Pre-PulseNet	Early PulseNet	Listeria Initiative	WGS
Outbreaks per year	0.3	2.3	2.6	10
Median cases per outbreak	69	11	5.5	3

# **HOW DID WGS AFFECT INVESTIGATIONS?**

# PFGE Cluster Possibly Associated with Uncommon Deli Meats, 2012–2013









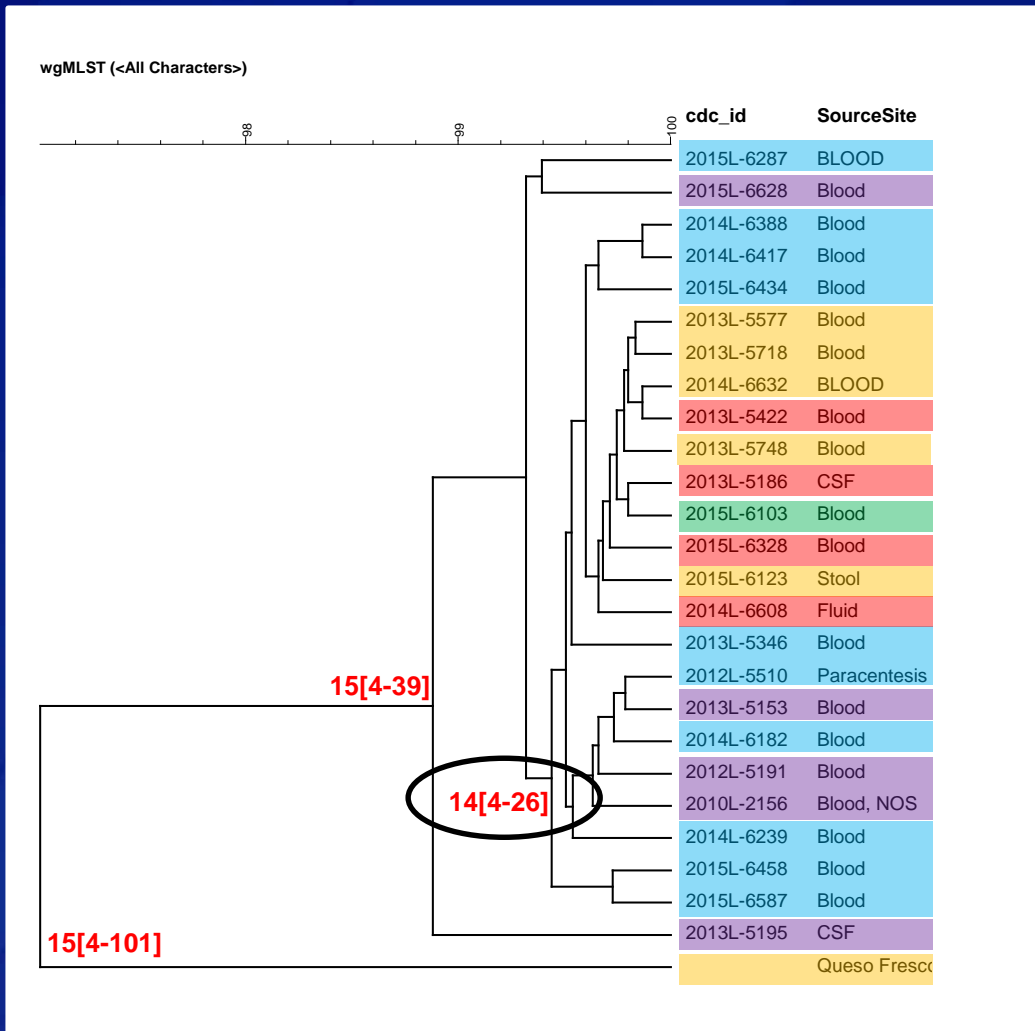
## **Cluster Investigation December 2013**

- ❑ **PulseNet identifies PFGE cluster of 4 cases with rare pattern**
  - California (2), Illinois (1), Massachusetts (1)
  - Specimen collection dates July–December 2013
- ❑ **All 4 patients seemed to be of Middle Eastern descent (Armenia, Syria, and Yemen)**
- ❑ **2 patients interviewed with supplemental form**
- ❑ **Unable to identify common food**
- ❑ **Cluster closed in January 2014**

# Cluster Investigation August 2015

- ❑ **PulseNet identifies cluster of 12 cases with different PFGE pattern, also rare**
  - California (7), Massachusetts (2), New York (2), Colorado
  - Specimen collection dates 2013–2015
  - 10 sequenced isolates  $\leq 25$  alleles by wgMLST
- ❑ **More WGS analysis shows isolates from 2010–2015 with 4 other PFGE patterns are also highly related, including isolates from 2013 cluster**

# wgMLST Phylogeny as of Sept. 16



- 5 different PFGE patterns
- All isolates from patients except “unrelated” food isolate at bottom

## Investigation Unfolds

- **With more cases**
  - Middle Eastern connection still present
  - Soft cheese association identified (OR 26,  $p < 0.001$ ), mostly Middle Eastern and Mediterranean style
  - Suspect Company A based on brands reported
  
- **PulseNet finds matching 5 environmental isolates from Company A cheese production facility in 2010**



# One dead, 24 sick from multi-state Listeria outbreak linked to soft cheeses

By Debra Goldschmidt, CNN

Updated 1:39 PM ET, Fri September 18, 2015





## Unlikely to Be Implications of WGS

*[Person A] drew the “yes” side of the first debate question [at International Association for Food Protection annual meeting], “Is shoe leather epidemiology dead in the era of whole-genome sequencing?”*

**-Food Safety News. July 27, 2015**

*“You catch things far earlier” with sequencing, said [Person B]. “It can be two cases. If you see a match, Bam! You've got em.”*

**-Reuters. August 27, 2015**

## **A New Era *Listeria* WGS**

- ❑ **Helped detect more clusters and solve more outbreaks**
- ❑ **Focused epidemiological resources**
- ❑ **Identified new food sources of listeriosis**
- ❑ **Epidemiologic data remain essential**
- ❑ **WGS not just for *Listeria* anymore**



# Acknowledgments

**State, Local, and Territorial Health Departments**

**Advanced Molecular Detection Initiative**

**Food and Drug Administration**

**US Department of Agriculture's Food Safety and Inspection Service**

**National Center for Biotechnology Information**

**Enteric Diseases Laboratory Branch**

**Enteric Diseases Epidemiology Branch**

**Outbreak Response and Prevention Branch**

**Epidemic Intelligence Service**

For more information please contact Centers for Disease Control and Prevention

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Visit: [www.cdc.gov](http://www.cdc.gov) | Contact CDC at: 1-800-CDC-INFO or [www.cdc.gov/info](http://www.cdc.gov/info)

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# Public Interest

## U.S. Google Search Trends for “Listeria”

