CDC PUBLIC HEALTH GRAND ROUNDS



Accessible Version: https://www.youtube.com/watch?v=jpXROWP6V74

The Emerging Role of Pathogen Genomics in Public Health

January, 21 2020



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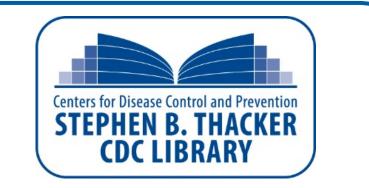
Additional Resources

Beyond The Data

"Take home" messages in a short podcast at: cdc.gov/grand-rounds





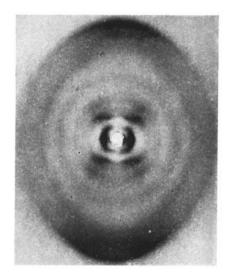


Scientific publications about this topic at: cdc.gov/library/sciclips

Email grandrounds@cdc.gov with any questions or for help locating the additional resources

Must Reads

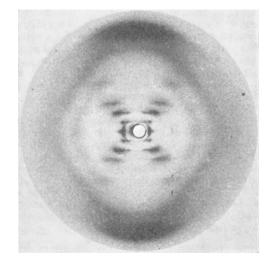
April 25, 1953 N A T U R E



MOLECULAR STRUCTURE OF NUCLEIC ACIDS

A Structure for Deoxyribose Nucleic Acid

WE wish to suggest a structure for the salt of deoxyribose nucleic acid (D.N.A.). This structure has novel features which are of considerable biological interest.



Today's Speakers and Contributors









Duncan MacCannell, PhD

Heather Carleton, MPH PhD

Lauren Turner, PhD

Gregory Armstrong, MD

Acknowledgments

Luciann Draper Luis Luque Paula Eriksen Steve Mann Ann Gallaty Alex Alvarez Charlotte Dugan Patrick Brady Nathelia Barnes Tom Race Muin Khoury William Thomas Marta Gwinn Elizabeth Neuhaus Scott Sammons

Upcoming Programs of Interest

February 18, 2020 Public Health Grand Rounds Measles

March 17, 2020 Public Health Grand Rounds Laboratory Response Network

CDC PUBLIC HEALTH GRAND ROUNDS



The Emerging Role of Pathogen Genomics in Public Health

January 21, 2020



CDC PUBLIC HEALTH GRAND ROUNDS



The Emerging Role of Pathogen Genomics in Public Health

January 21, 2020



Introduction to Advanced Molecular Detection and Pathogen Genomics

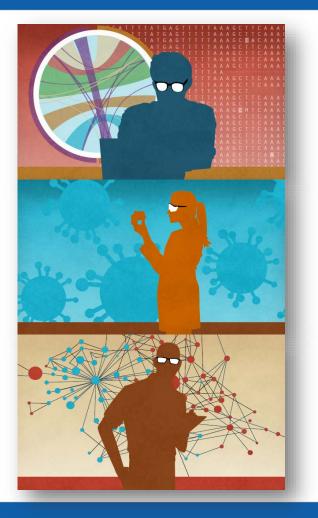


Duncan MacCannell, PhD

Chief Science Officer, Office of Advanced Molecular Detection Centers for Disease Control and Prevention



Advanced Molecular Detection (AMD)



>AMD program established by Congress in 2014

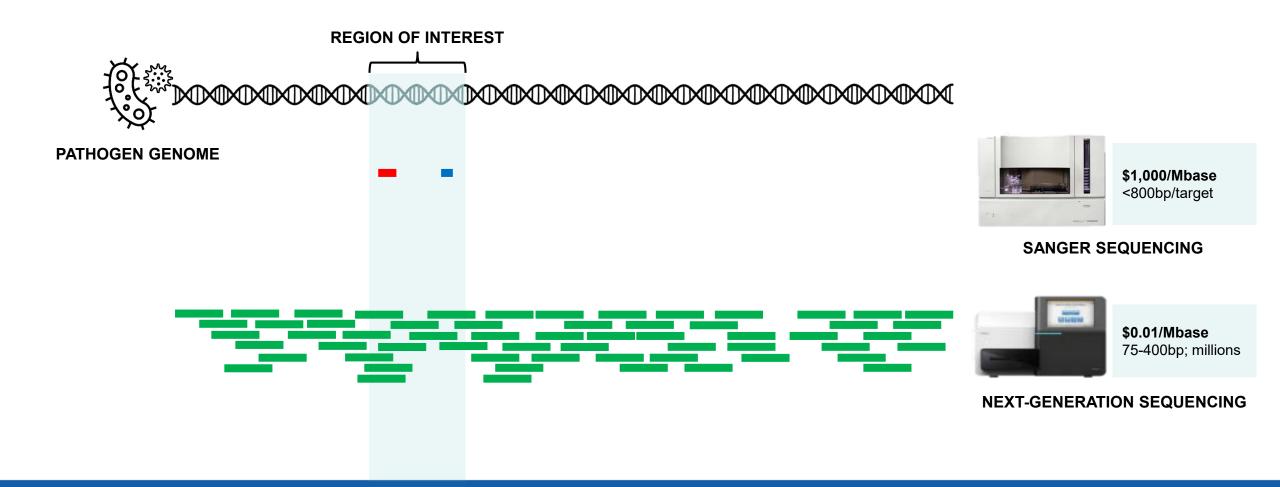
- \$30 million per year public health laboratory innovation program
- Focus on transformational laboratory tech and scientific computing

Principal objectives of AMD

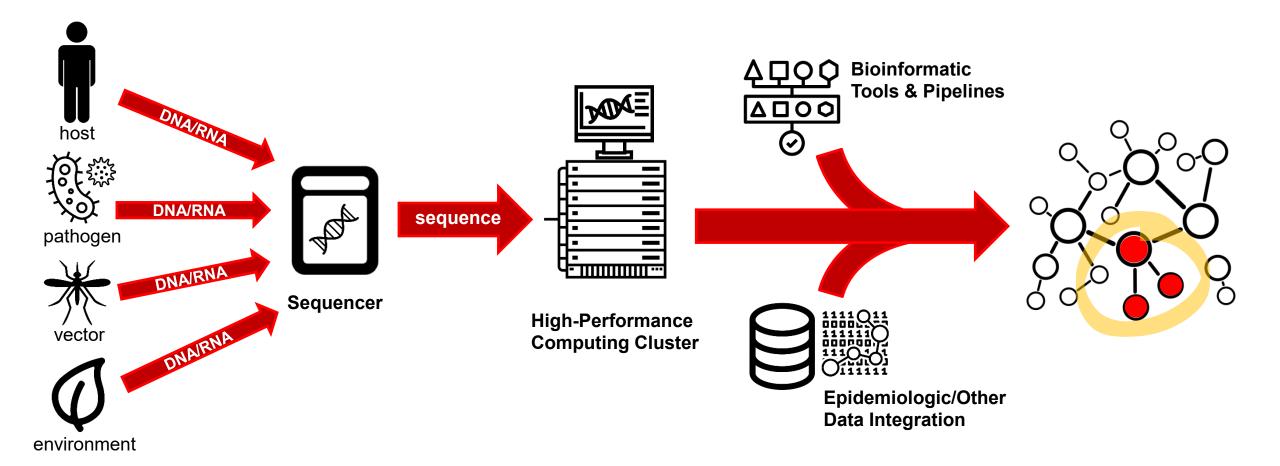
- Improve pathogen detection and characterization
- Enable new diagnostic methods to meet public health needs
- Support genomic and bioinformatics needs in the US public health system
- Implement enhanced, sustainable, integrated information systems
- Develop tools for prediction, modeling and early recognition of emerging infectious threats

www.cdc.gov/amd

Sequencing in Public Health

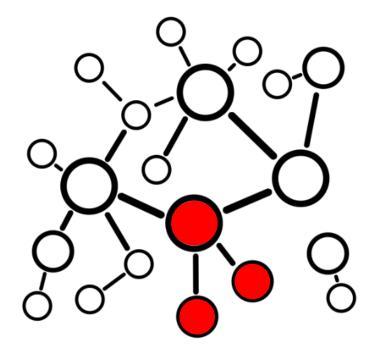


How is NGS Used in Public Health?



Armstrong et al, NEJM 2019

What Information can be Obtained from Pathogen Genomes?



Pathogen Genomics

- Reference identification
- High-resolution genotyping
 - Clustering and phylogenetics
- Functional annotation and characterization
 - Virulence, antimicrobial resistance, signatures
 Phenotype prediction
- Diagnostic development and validation
- Minor population variants, quasi-species
- Microbial ecology and diversity

AMD Projects and Applications: 2019



www.cdc.gov/amd/what-we-do/

On the Front Lines: Transforming Influenza Surveillance

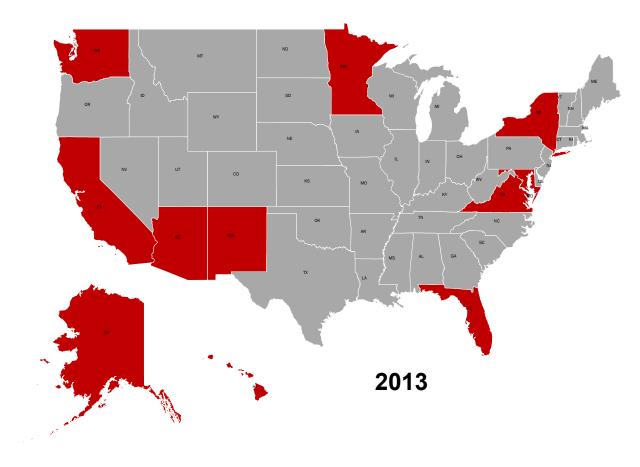


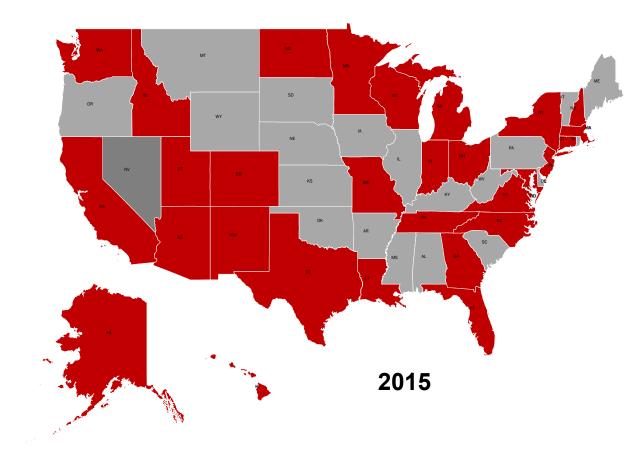


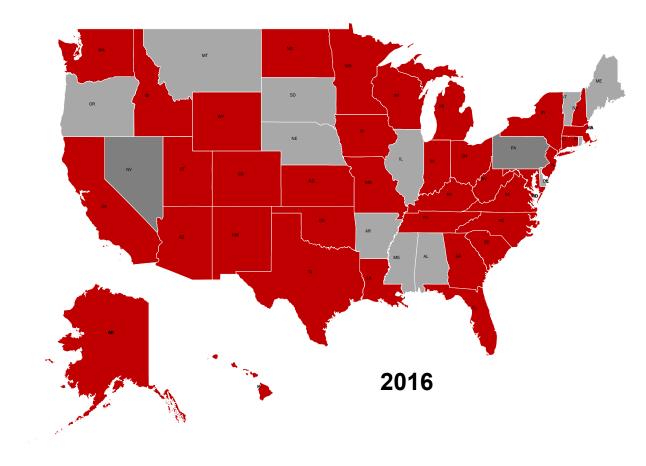


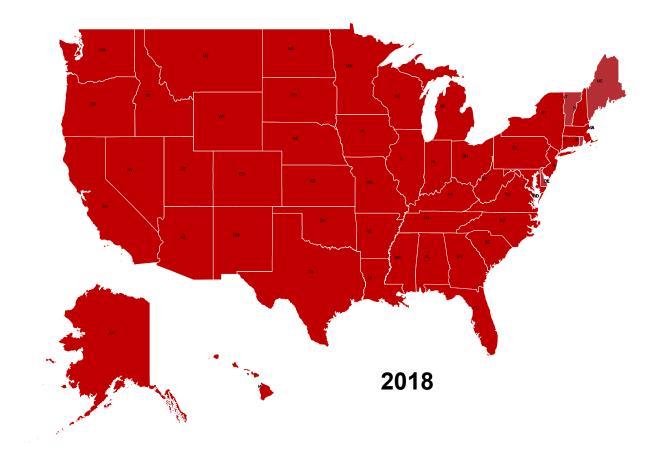
Sequencing instrument

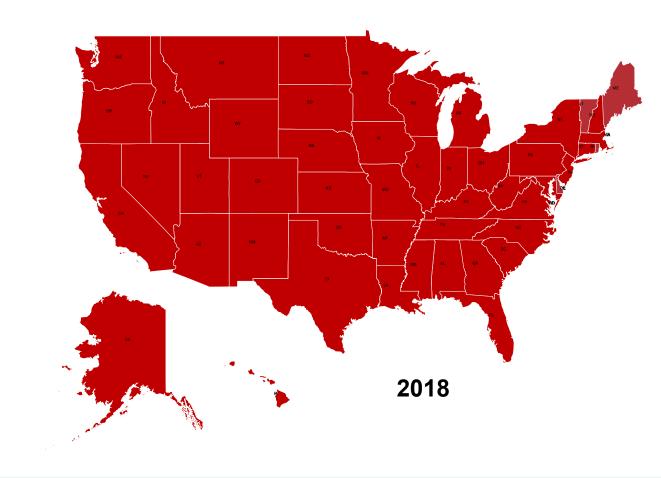
Rambo-Martin et al, bioRxiv 2019





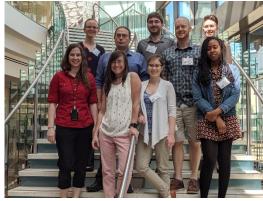






- > A growing list of applications
 - PulseNet
 - Healthcare-associated infections
 - Bacterial meningitis
 - Cryptococcus
 - Cyclospora
 - Viral hepatitis
 - Gonococcus
 - Influenza
 - Legionella
 - Malaria
 - Streptococcus
 - Tickborne diseases
 - Tuberculosis
 - To be determined

Building Sustainable Public Health Bioinformatics Capacity



CDC/APHL Bioinformatics Fellows, 2019 Cohort



AMD Bioinformatics Resource and Training Regions

Workforce development (recruitment and training)

- CDC and APHL Bioinformatic Fellowship Program
- AMD bioinformatics regional resources and training Leads
- Building community: state public health bioinformaticians (StaPH-B)
- Engage across the public health, academic, and private sector communities.
- Bridge the gap between existing open source bioinformatics software development and public health users and applications.

www.aphl.org/bioinformatics www.staphb.org www.cdc.gov/labguality/ngs-guality-initiative.html

AMD: In Summary

>AMD program

- Working to foster multidisciplinary laboratory innovation in public health
- Helping to drive shift toward open data, standards, methodological transparency and collaboration.
- With financial and technical support from CDC and key partners, NGS capacity has spread nationwide over the past 6 years.
- HPC infrastructure, bioinformatics workforce capacity, quality, and data management remain critical challenges.

Applications of Next-Generation Sequencing to Foodborne Disease Prevention



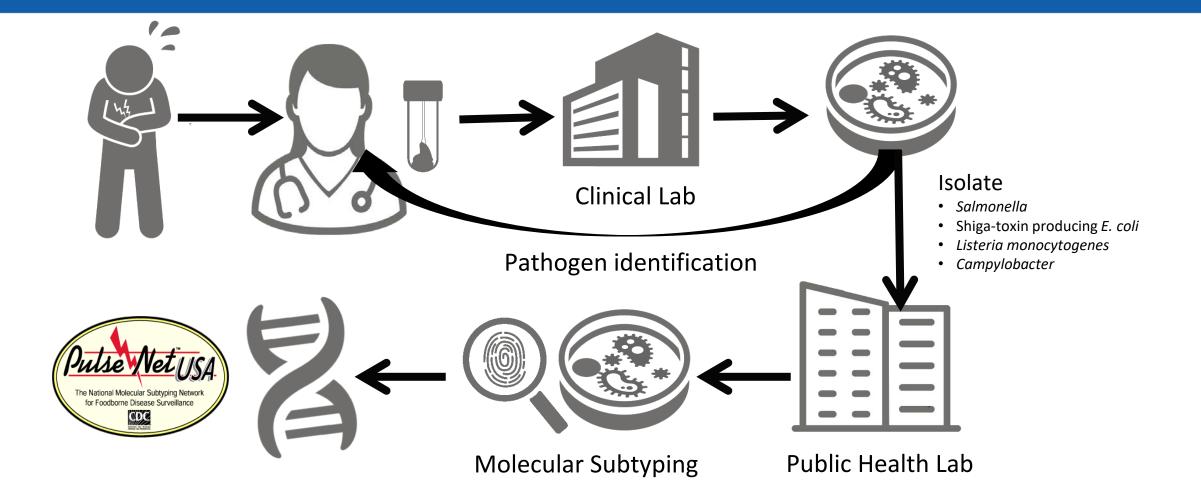
Heather Carleton, MPH, PhD

Bioinformatics and Metagenomics Team Lead

Enteric Diseases Laboratory Branch Division of Foodborne, Waterborne, and Environmental Diseases



Foodborne illness: From Patient to Public Health Surveillance



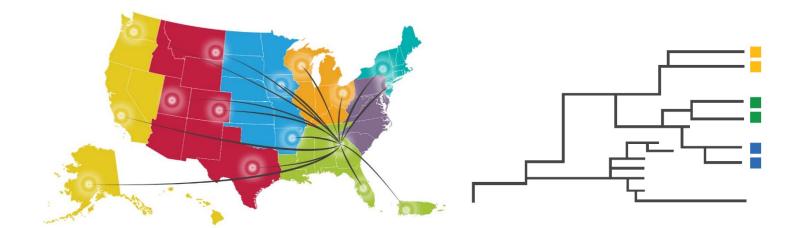
What is PulseNet?



TGGTATTTATGCTGCTGGAGGTTCCACAA GCCGCGTGCGGAGCCATATTTATTCCCCG CCGCATACGCCAGGATAATCCAGCATAAA GTTCTTTCCGTATGTTTTGCGTTCATTTA A GCTGCTGGAGGTTCGTGGTATTTCCACAA ITTTGCGTTGAGCCATATTTATTCCCCGTC CGCTTTATCGCCTAGACAAAACCTGCATAA GCCCCGCCGTGTCGGGA AATGTGTTCATT ACCTGCATAAATTCGCTTTATCGCCTAGA GGGA AATGTGTTCATTTA ACATTGCCCC

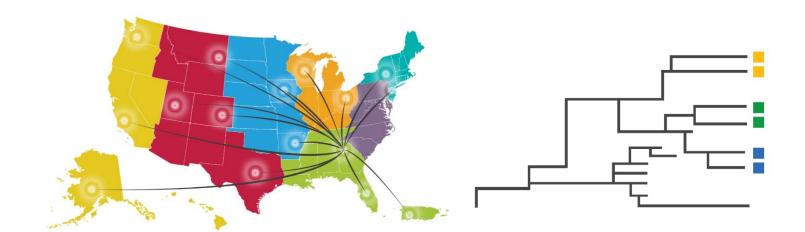
Public health labs generate molecular fingerprints using WGS

What is PulseNet?



Data from pathogens are transmitted to centralized database at CDC in Atlanta, Georgia PulseNet Monitors for clusters of illnesses with the same molecular "fingerprint"

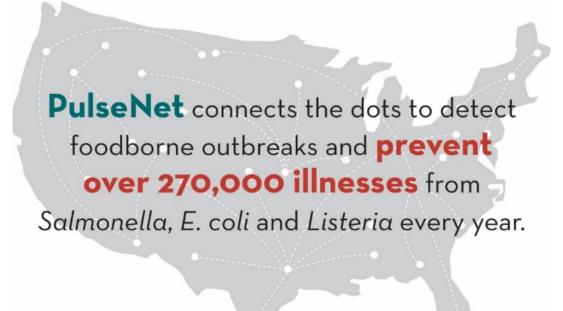
What is PulseNet?





Data from pathogens are transmitted to centralized database at CDC in Atlanta, Georgia PulseNet Monitors for clusters of illnesses with the same molecular "fingerprint" Then tells epidemiologists about clusters to investigate

PulseNet 1996-2020: Disease Prevention and Cost Savings



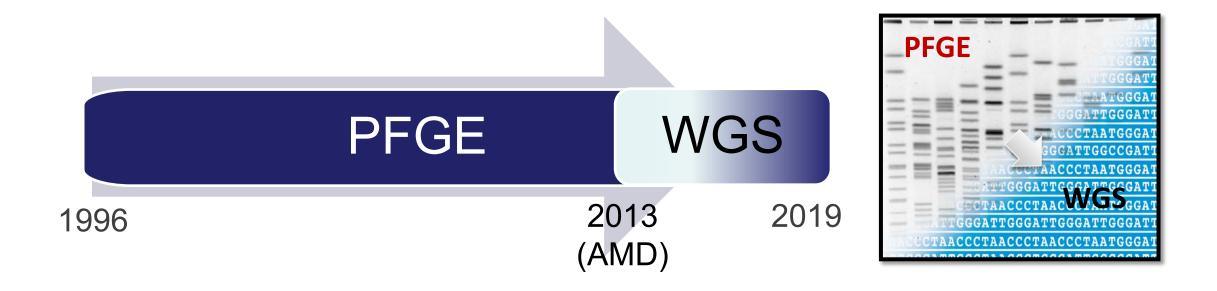
Every year **PulseNet** saves at least **half a billion dollars**

in medical costs and lost productivity.

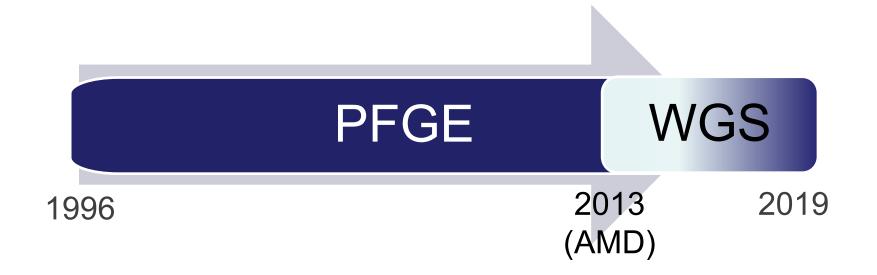


\$1 spent = \$70 saved!

Transition to Use of Whole Genome Sequencing in PulseNet



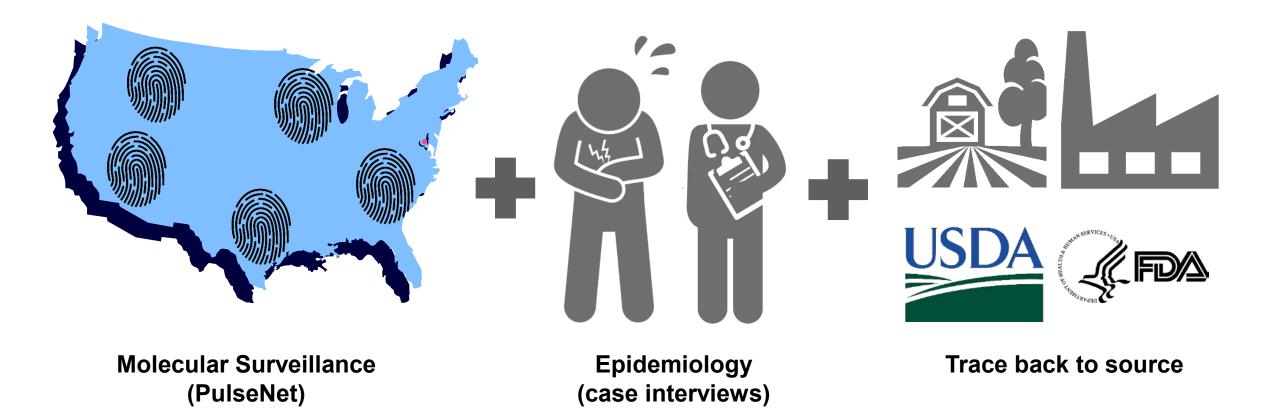
Transition to Use of Whole Genome Sequencing in PulseNet



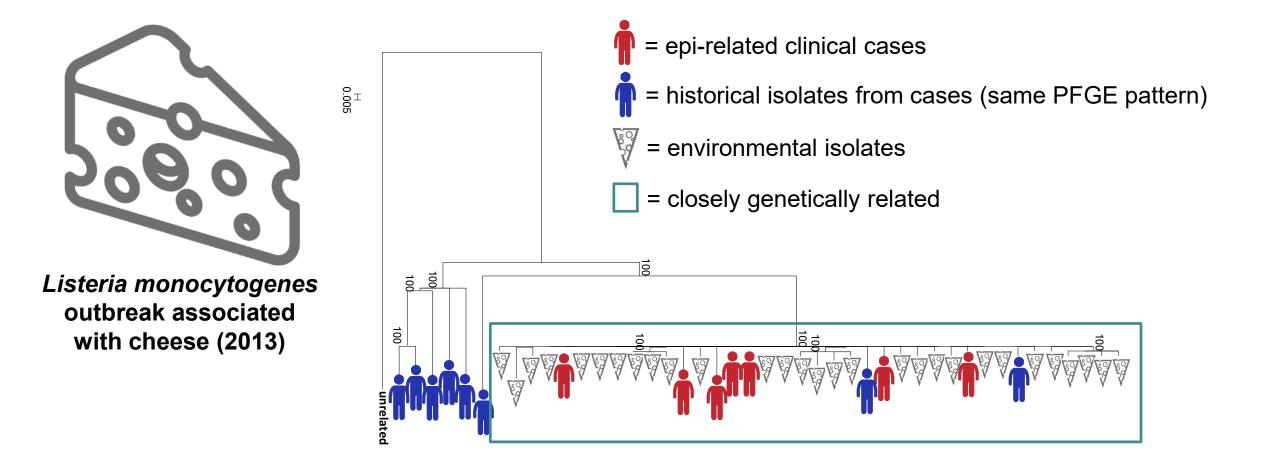
Utility of WGS Data:

- Detecting outbreaks with more precision
- More effective outbreak investigations and trace back

Putting Together the Picture of a Foodborne Outbreak

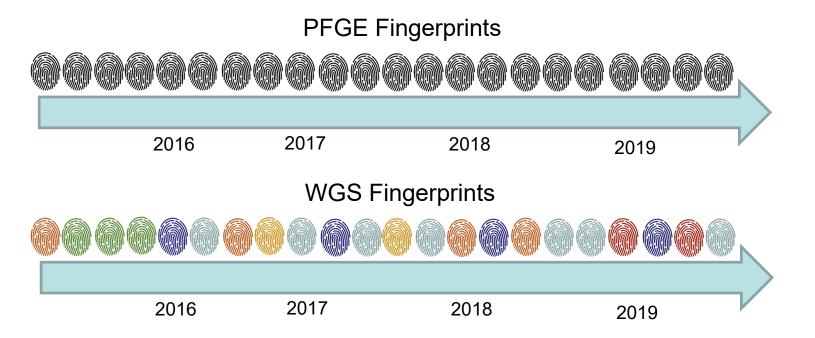


Using WGS to Link Clinical Cases with Outbreak Source



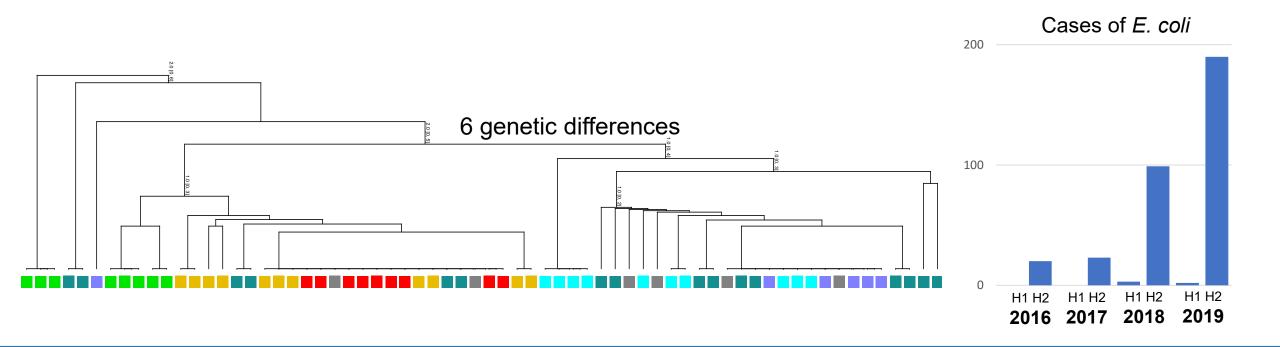
Using WGS to Track Reoccurring, Emerging, and Persisting Strains

WGS links cases that are associated with reoccurring, emerging, and persisting strains



Using WGS to Track Reoccurring, Emerging, and Persisting Strains

WGS linked cases of *E.coli* O157:H7 that occurred over a 4-year period Linked to a single growing region

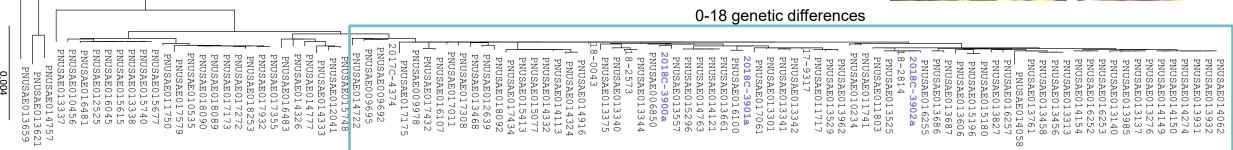


figures courtesy of Kane Patel

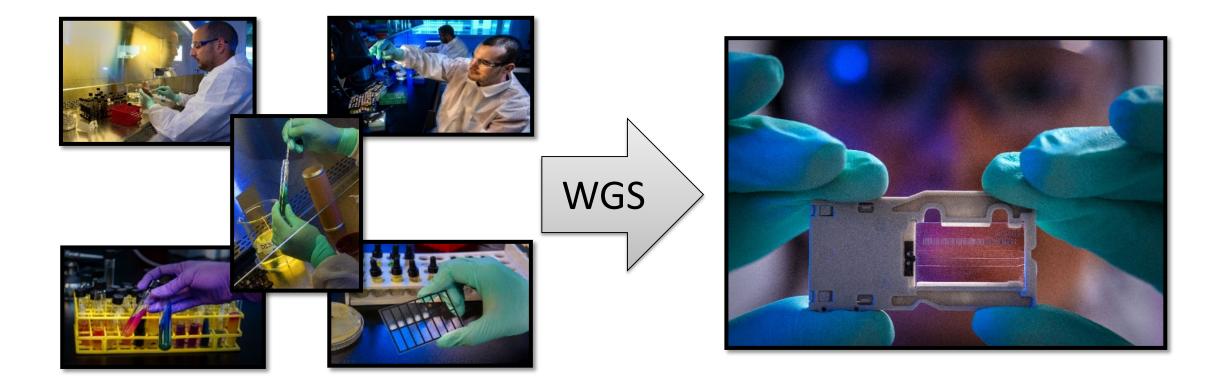
Using WGS to Help Outbreak Prevention Efforts

- E. coli O157 infections were linked to romaine lettuce from Yuma growing area
- *E. coli* O157 from water matched outbreak strains
- Led to new public health action
 - Requirement to sanitize water used for overhead irrigation during the past 21 days of production
 - Label bag with growing region and harvest date





WGS Provides More Information Than Just Relatedness

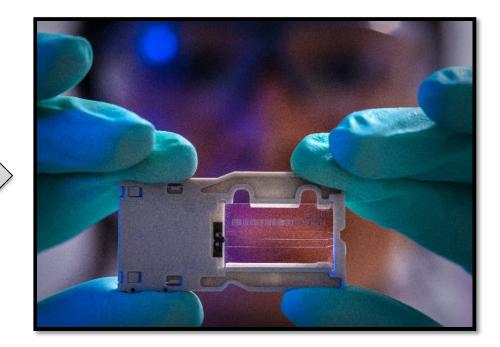


WGS Provides More Information Than Just Relatedness

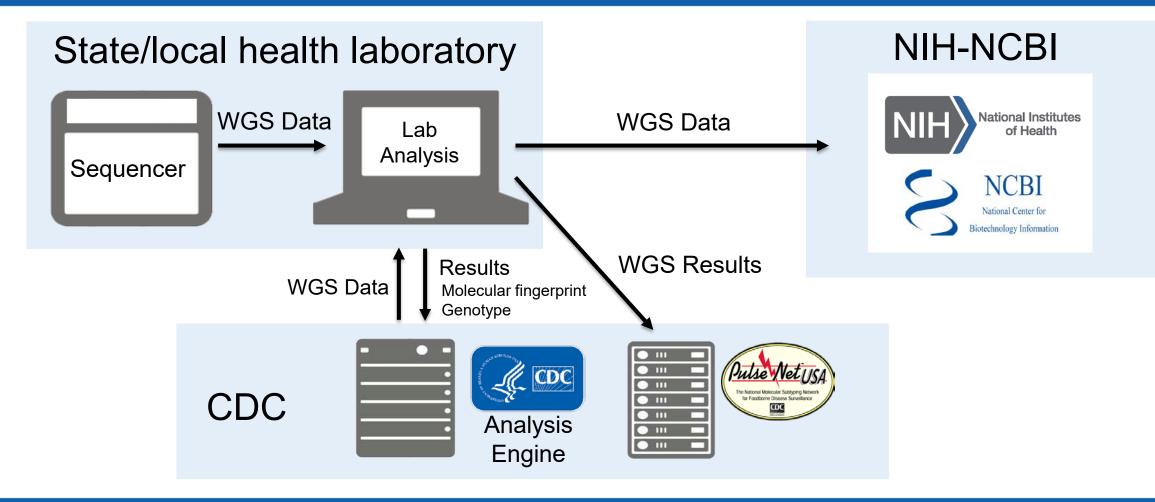
WGS

Genotyping provides

- Identification of organism
- Serotype (O and H genes; other surface marker genes)
- Antibiotic resistance genes and mutations – which can be associated with harder to treat infections
- Virulence genes can lead to more severe infections



PulseNet Uses a Distributed Analysis System to Extract Information from WGS

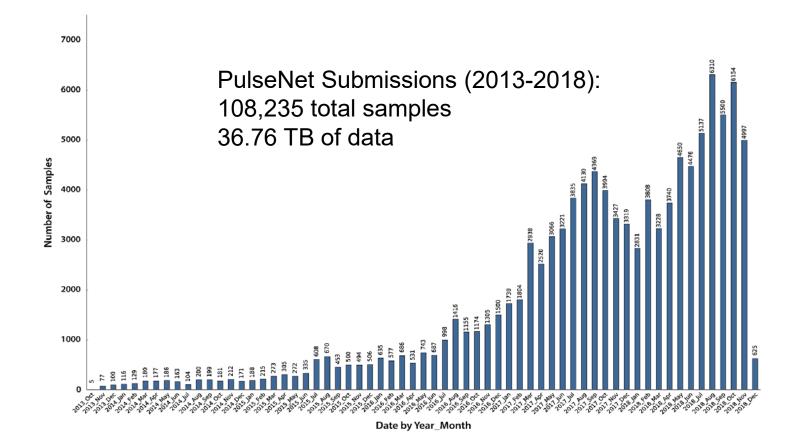


Challenges

- Amount of data
- Complex computing jobs
- Differing IT resources

Opportunities

- Detect foodborne outbreaks with more precision
- Receive genotyping and molecular fingerprint results for local cluster detection



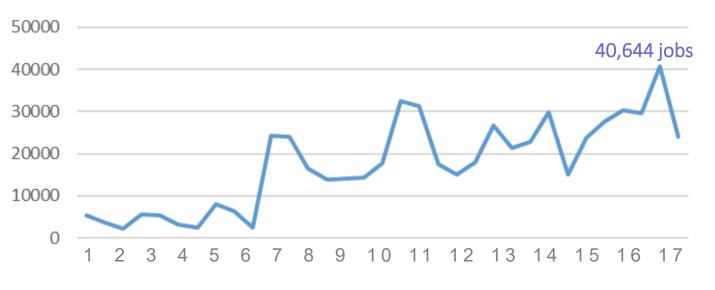
Mbps: megabits per second (1 Mbps = dialup internet, > 100 = "fast internet")

Challenges

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Jobs per Week in 2019

Mbps: megabits per second (1 Mbps = dialup internet, > 100 = "fast internet")

Challenges

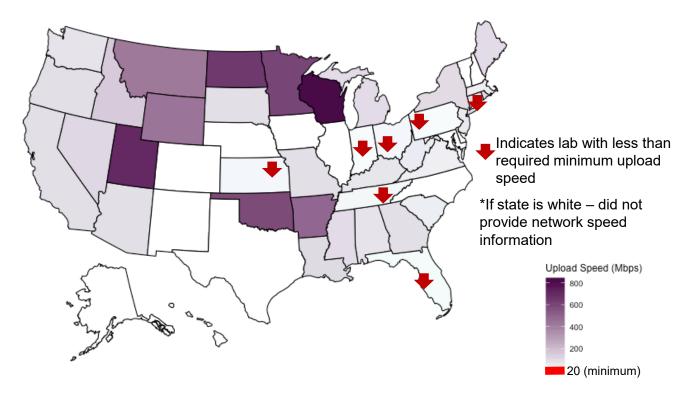
Amount of data

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Opportunities

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Public Health Lab Upload Speeds



Mbps: megabits per second (1 Mbps = dialup internet, > 100 = "fast internet")

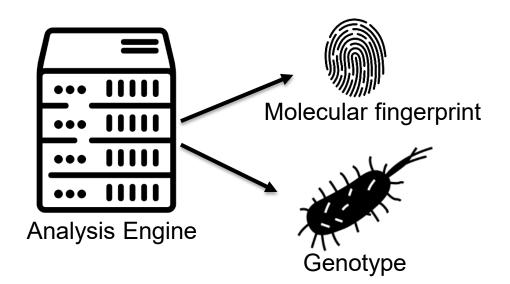
Challenges

- Amount of data
- Complex computing jobs
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Opportunities

44

- Detect foodborne outbreaks with more precision
- Receive genotyping and molecular fingerprint results for local cluster detection



Mbps: megabits per second (1 Mbps = dialup internet, > 100 = "fast internet")

The Future of PulseNet

- With the transition to WGS, PulseNet will be able to detect outbreaks with more precision.
 This transition was made possible with initial investments by AMD.
- Beyond WGS, PulseNet is continuing to innovate by applying novel technologies like metagenomics to characterize foodborne pathogens directly from patient samples.



State-level AMD Applications Across the Infectious Disease Spectrum



Lauren Turner, PhD

Foodborne and Advanced Pathogen Characterization Lead Scientist

Virginia Department of General Services, Division of Consolidated Laboratory Services



U.S. Department of Health and Human Services Centers for Disease Control and Prevention

Division of Consolidated Laboratory Services (DCLS) Virginia's State Laboratory

- The Commonwealth of Virginia's public health, environmental, agriculture, and consumer protection laboratory
- Serving local, state, and federal agencies
- Comprehensive testing services including
 - Neonatal screening
 - Drugs of abuse testing
 - Microbiology testing
 - □ Food and water contamination monitoring
 - Metal and pesticide analyses
 - Chemical analyses





Infectious Disease Surveillance The Not so Good Old Days

Pre-AMD surveillance and outbreak detection tools

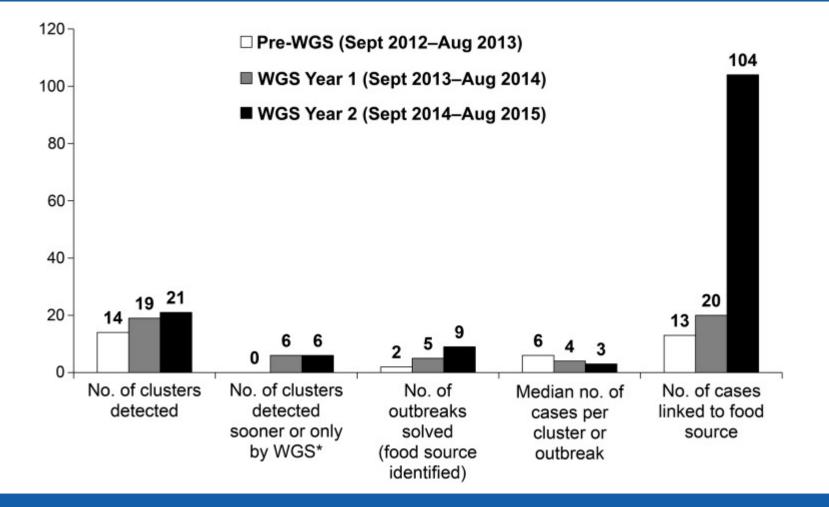
- Specimen culture and pathogen identification
- Serovar or serotype subgrouping
- Virulence factor genotyping and phenotyping
- DNA fingerprinting by pulsed-field gel electrophoresis (PFGE)
 - □ Foodborne, healthcare-associated and community acquired outbreaks; >25 different organisms

Constraints

- Costly and time intensive
- Loss of institutional knowledge with aging workforce
- Limited detection with targeted DNA-methods
- Limited resolution of pathogen genetic differences or similarity

Advancing Public Health Laboratory Infectious Disease Surveillance

- WGS improves accuracy
- Provides epidemiologists and regulatory officials with actionable results
- Improves use of laboratory resources
- Reduces turnaround time



Jackson BR et al, Clin Infect Dis 2016 WGS: whole genome sequencing Advancing Pathogen Surveillance through Next-generation Sequencing (NGS)

Goals of Virginia DCLS

- Pragmatic implementation of NGS to meet state and federal program needs
- Develop capacity for data analytics (bioinformatics)

Use pathogen genomics bioinformatics tools

- Pathogen identification
- Pathogen genotyping
- Early detection of possible outbreaks
- Enhanced evaluation of suspected outbreaks
- Outbreak tracking
- Source attribution





Emerging Applications of Next-generation Sequencing

Began as primarily foodborne enteric bacterial pathogens

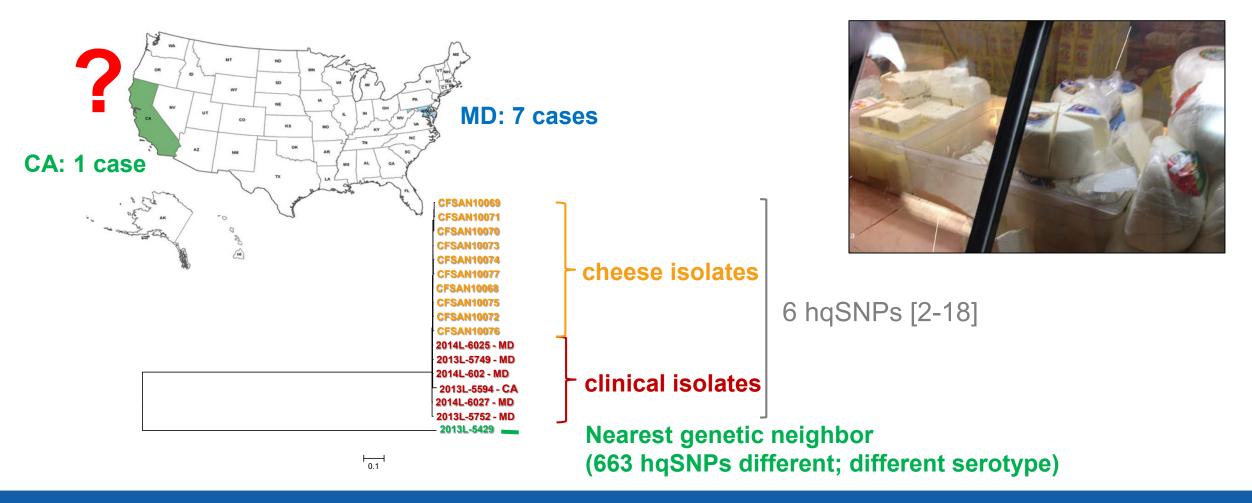
- Network-driven initiative (PulseNet)
- Surveillance and outbreak detection
- Transitioning from PFGE to WGS since 2014, beginning with *Listeria*
- Heavy reliance on federal partners for bioinformatics analysis
- Growing dependence on WGS data
 - □ Foodborne pathogen surveillance solely by NGS in June 2019
- Use of NGS expanding to tuberculosis, *Legionnaires' disease*, healthcare-associated infections, and more

DCLS' AMD Capacity Building: Number of Trained Staff, Samples, and Public Health Applications

	2013	2014	2015	2016	2017	2018	2019
Staff trained to sequence	Ŵ	Ē	Ŵ Ŵ	ŤŤ	ĦĦ ĦĦ	ĦĦ ĦĦĦ	ĦĦĦĦĦ ĦĦĦ
Staff trained to extract NA	Ť	Ť	ŤŤ	Ů Ů Ů Ů	<u></u>	ĦĦĦ ĦĦĦ	☆☆☆☆☆☆☆
Total samples sequenced	12	498	738	1063	2365	2252	2160
Instruments							
Bioinformatics staff				* _/	i		
GenomeTrakr	Ø	•	⊘	0	Ø	Ø	O
PulseNet		Ø	Ø	€	0	⊘	⊘
Tuberculosis					0	Ø	•
HAI						♥	Ø

NA: Nucleic acid HAI: Healthcare-associated infections

AMD Applications: Listeriosis Outbreak Source Tracking



hqSNP: high-quality single nucleotide polymorphism; median [range]

AMD Applications: Infection Transmission and Control in Healthcare Settings

Investigating transmission of pathogens

- □ Is transmission occurring between facilities?
- □ Is this strain regionally or locally persistent?
- □ Is there asymptomatic infection among healthcare providers?

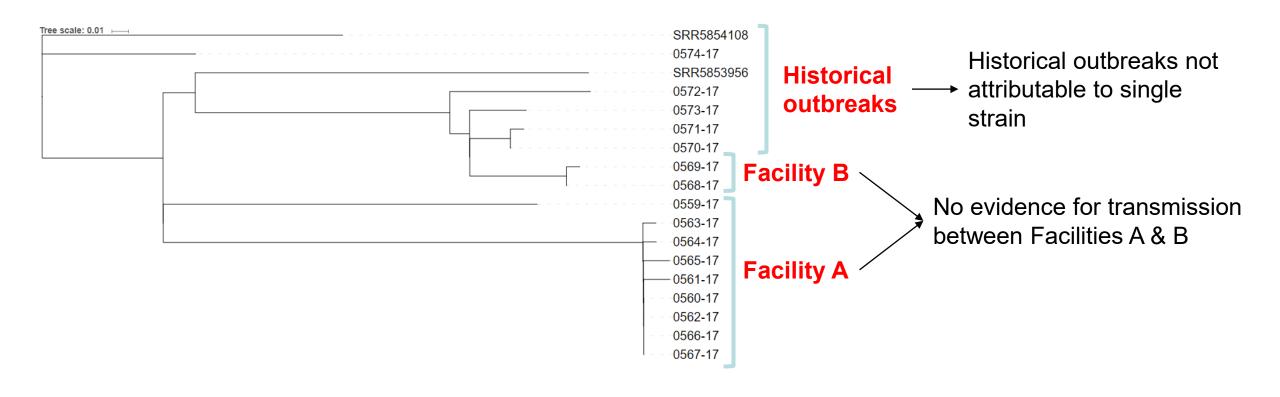
NGS to assess severe invasive Group A Streptococcus (GAS) outbreaks

- Multiple facilities
- Regional occurrence
- 2013-2017



AMD Applications: Infection Transmission and Control in Healthcare Settings

Group A Streptococcus (GAS) Isolates from Clinical Facilities



AMD Applications: Pathogen Identification and Genotyping

Applying predictive analytics for pathogen identification from sequence data

Reference database comparison

- Species identification
- Genotyping serotype prediction

sample	r1_q	r2_q	est_genome_length	est_cvg	number_contigs	species_prediction	subspecies_predictions
isolate001	35.61	31.92	5367997	76.32	256	Escherichia_coli	O26:H11
isolate002	34.09	28.99	5417894	49.7	253	Escherichia_coli	O103:H2
isolate003	36.36	33.63	2931612	130.08	27	Listeria_monocytogenes	NA
isolate004	36.86	35.43	4754353	77.82	31	Salmonella_enterica	Heidelberg
isolate005	36.67	34.53	1686219	120.42	15	Streptococcus_pyogenes	emm89
isolate006	36.42	34.93	1802069	94.48	24	Streptococcus_pyogenes	emm1

Challenges to AMD Integration

Development of scientist knowledge, skills, and abilities

- Next-generation sequencing (NGS) methods and instrumentation
- Bioinformatics analysis of NGS data

Building of NGS infrastructure

- Procurement, installation, and maintenance
- Data storage and transfer solutions for terabytes of NGS data
- Computational resources for data analysis
- Policies and procedures to support expanding use of NGS and efficient analysis of data
- Funding to sustain and capitalize on AMD investments

Opportunities Through AMD Integration

Many different approaches to achieve state capacity for AMD

Investment in different IT structures and software applications for data analysis
 Different avenues for computational resources



Opportunities Through AMD Integration

AMD Training & Bioinformatics Resource Support



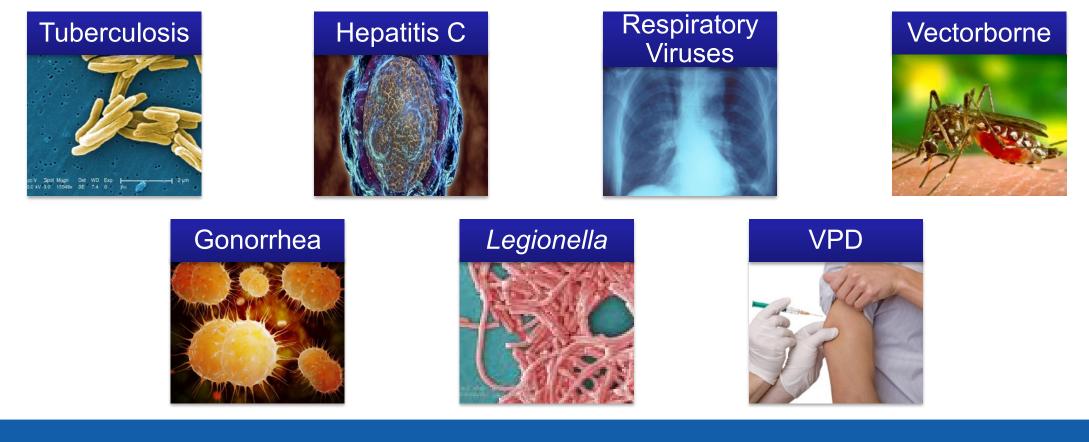
Modernization of public health lab testing

Novel partnerships to strengthen state programs

- Regional training and bioinformatics resource support
- University partnerships

Opportunities Through AMD Integration

States can tailor AMD to meet local disease control needs



VPD: vaccine-preventable disease

Future Virginia Laboratory Priorities and Needs

- Data integration and visualization
- **Data sharing agreements**
- Guidance for alignment with clinical laboratory requirements
- Applying AMD to detect pathogens in clinical samples
- Interdisciplinary workforce capacity building

Where Does Pathogen Genomics Go from Here?



Gregory L. Armstrong, MD

Director, AMD Program Centers for Disease Control and Prevention



U.S. Department of Health and Human Services Centers for Disease Control and Prevention

	AMD before 2013	AMD in 2020		
NGS capacity	One sequencer	Capacity in all infectious disease laboratories		
Bioinformatics capacity	<10 bioinformaticians	~60 bioinformaticians; widespread experience and expertise in microbial genomics		
Capacity in state/local health departments	NGS capacity in one laboratory	NGS capacity in all state and many local laboratories; 7 regional bioinformaticians and an increasing number of state-based bioinformaticians		

Continued adaptation of NGS to public health priorities Emerging areas of interest

- New sequencing technologies
- Metagenomics (sequencing directly from samples, rather than from cultured isolates)
- Data integration and modernization

cdc.gov/amd @cdc_amd

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.

Apply to Become a Bioinformatics Fellow

> Applications Due: February 29, 2020

The <u>APHL-CDC Bioinformatics Fellowship</u> aims to train and prepare bioinformaticians to apply their expertise within public health and design tools to aid existing public health personnel in the use of bioinformatics. Post-master's and postdoctoral-level professionals can apply their skills to a range of important and emerging public health problems, while gaining experience in their fields. Whether your specialty is metagenomics, algorithm/software development, microbial genomics, or another research area, we have a place for you!

www.aphl.org/fellowships/Pages/Bioinformatics.aspx