Clinical Laboratory COVID-19 Response Call Monday, February 8th, 2021 at 3:00 PM ET

Welcome

- Jasmine Chaitram, CDC Division of Laboratory Systems (DLS)
- CDC Virtual Reality Laboratory Training Course
 - Joe Rothschild, CDC Division of Laboratory Systems (DLS)
- Sequencing Efforts of Public Health Laboratories
 - Heather Blankenship and Marty Soehnlen, Michigan Department of Health and Human Services
- SARS-CoV-2 Variants Update
 - Vivien Dugan, CDC Laboratory and Testing Task Force for the COVID-19 Response
- Vaccine Breakthrough Case Investigations
 - Leisha Nolen, CDC COVID-19 Response Vaccine Breakthrough Team
- FDA Update
 - Tim Stenzel, U.S. Food and Drug Administration (FDA)

COVID-19 Resources for Laboratories

- LOINC In-Vitro Diagnostic (LIVD) Test Code Mapping for SARS-CoV-2 Tests
 https://www.cdc.gov/csels/dls/sars-cov-2-livd-codes.html
- IVD Industry Connectivity Consortium https://ivdconnectivity.org/livd/
- Antigen Testing Guidance
 https://www.cdc.gov/coronavirus/2019-ncov/lab/resources/antigen-tests-guidelines.html
- Frequently Asked Questions about COVID-19 for Laboratories https://www.cdc.gov/coronavirus/2019-ncov/lab/faqs.html

- Interim Guidance for Collecting, Handling, and Testing Clinical Specimens
 - https://www.cdc.gov/coronavirus/2019nCoV/lab/guidelines-clinical-specimens.html
- Diagnostic Tools and Virus
 https://www.cdc.gov/coronavirus/2019-ncov/lab/tool-virus-requests.html
- Emergency Preparedness for Laboratory Personnel https://emergency.cdc.gov/labissues/index.asp
- CDC Laboratory Outreach Communication System (LOCS) https://www.cdc.gov/csels/dls/locs/

CDC Preparedness Portal

https://www.cdc.gov/csels/dls/preparedlabs/covid-19-clinical-calls.html

Find CLCR call information, transcripts, and audio recordings on the CDC Preparedness Portal

Division of Laboratory Systems



Schedule for Clinical Laboratory COVID-19 Response Calls

The next call will be on **Monday, February 22**nd from **3:00 PM to 4:00 PM ET**



We Want to Hear From You!

Training and Workforce Development

Questions about education and training?

Contact LabTrainingNeeds@cdc.gov



How to Ask a Question

- Using the Zoom Webinar System
 - Click the Q&A button in the Zoom webinar system
 - Type your question in the Q&A box and submit it
 - Please do not submit a question using the chat button





- For media questions, please contact CDC Media Relations at media@cdc.gov
- If you are a patient, please direct any questions to your healthcare provider

Center for Surveillance, Epidemiology, and Laboratory Services

CDC Virtual Reality Laboratory Training Course

Joe Rothschild

Training and Workforce Development Branch
Division of Laboratory Systems

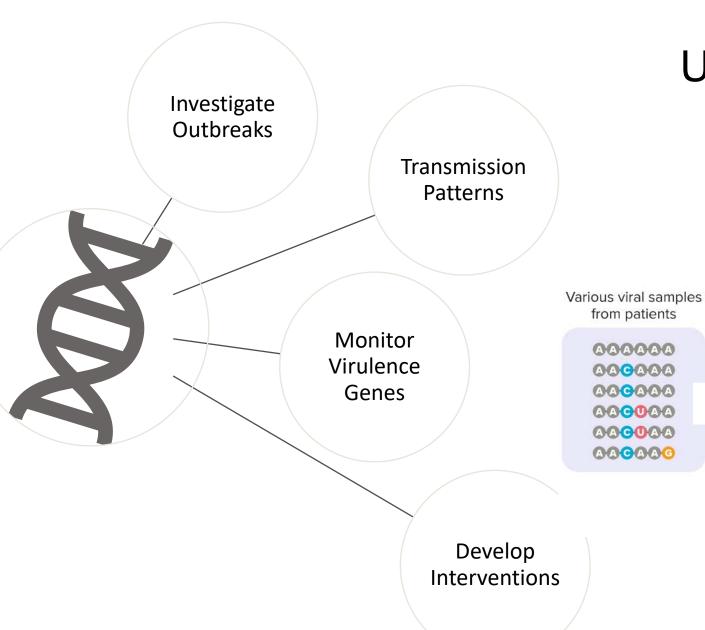




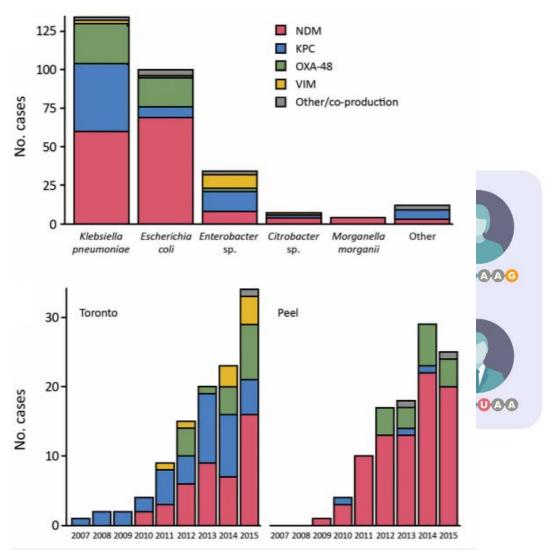
CDC DLS Presentation 8 February 2021

Marty Soehnlen, PhD MPH PHLD(ABB)
Director of Infectious Disease
Michigan Dept of Health and Human Services

Heather Blankenship, PhD Bioinformatics/Sequencing Section Manager Michigan Dept of Health and Human Services

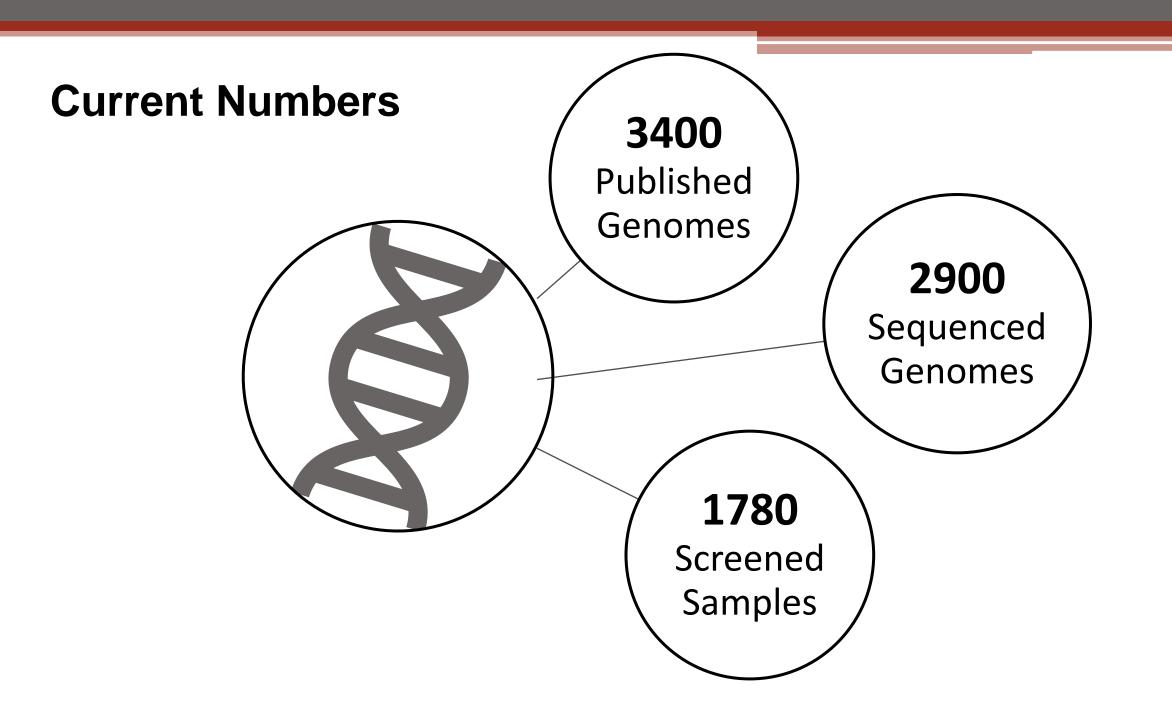


Use of Pathogen Genomics



SARS-CoV-2 Response

- 1 Identify baselines for the virus
- 2 Examine transmission dynamics
- Identify clinically important variants
- Contribute to the viral biology knowledge
- 5 Examine cluster outbreaks



SARS-CoV-2 Response

Sequencing

- Wet Lab
- Sequencer

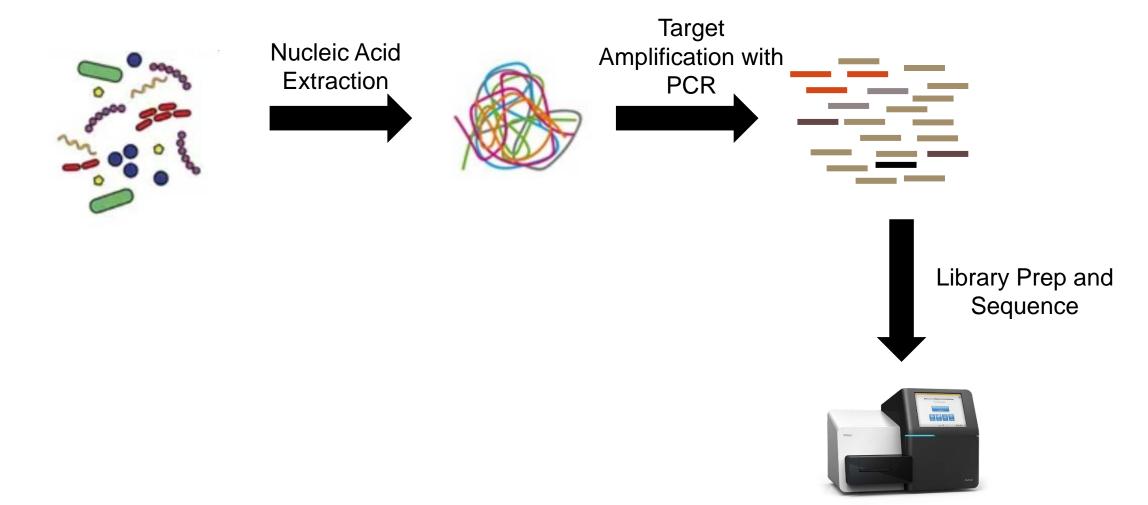
Bioinformatics

- Pipelines
- QC
- PublicRepositories

Communicate

- Share with epi partners
- Nextstrain
- Visualizations

Sequencing



Nucleic Acid Extractions

- QIAGEN EZ-1
- QIAGEN QIAcube
- MagnaPure 24
- MagnaPure Compact
- ThermoFisher Kingfisher
- Omega BioTek**

** by end of February

No difference among extraction methods

What do we sequence?

- Anything with a Ct < 30-32
 - PCR screen: CDC primers or ThermoFisher
- Epidemiological interest: within a reasonable Ct value
- Representative sample distribution

Sequencing Preps

- ARTIC v3 primers -> Nextera XT -> NextSeq
- ARTIC v3 primers -> Nextera Flex -> MiSeq
- QIAGEN QIAseq -> Nextera Flex -> MiSeq
- QIAGEN QIAseq -> Nextera XT -> NextSeq

SARS-CoV-2 Response

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Bioinformatics

- Pipelines
 - computational space and resources available
 - command line v packages (CLC Workbench, BioNumerics, etc)
- QC
 - at what thresholds will you make for sharing data public or utilize for analysis
 - our thresholds:
 - read coverage: 50X
 - genome coverage: >94% coverage

Public Repositories

GISAID – consensus sequence

- NCBI raw sequencing data and/or consensus sequence
 - requires confirmation that human reads are cleaned from the data

SARS-CoV-2 Response

Sequencing

- Wet Lab
- Sequencer

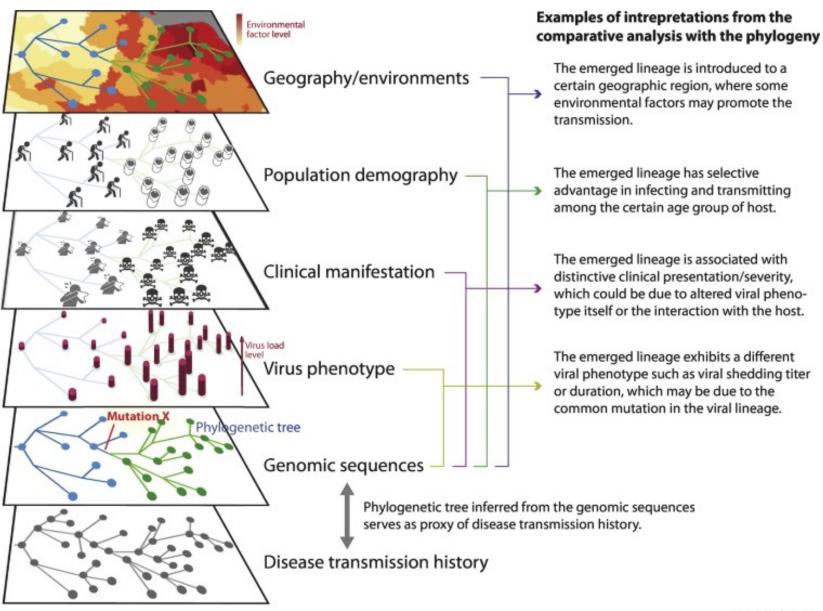
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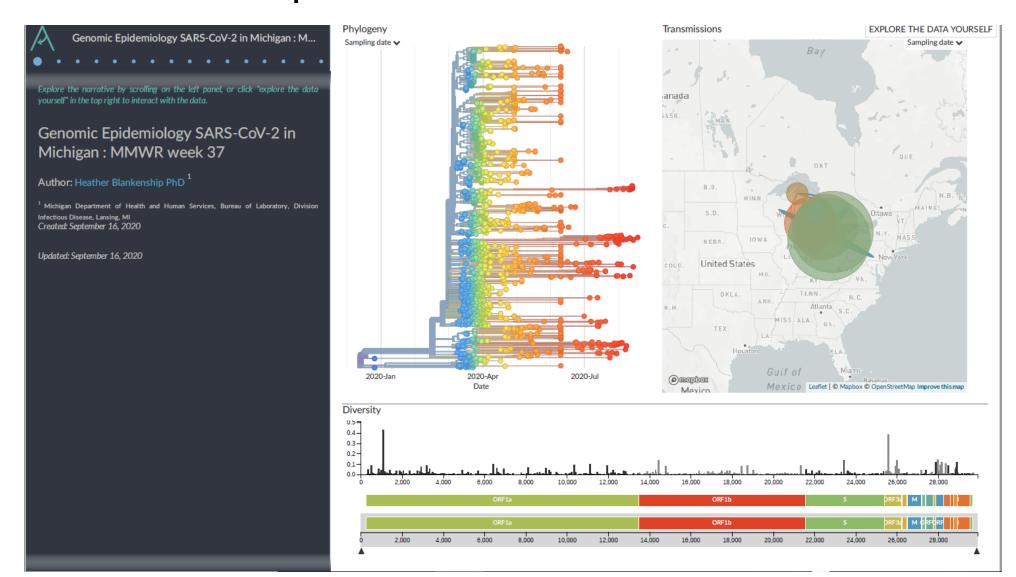
- Share with epi partners
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How can we use the data for public health action?



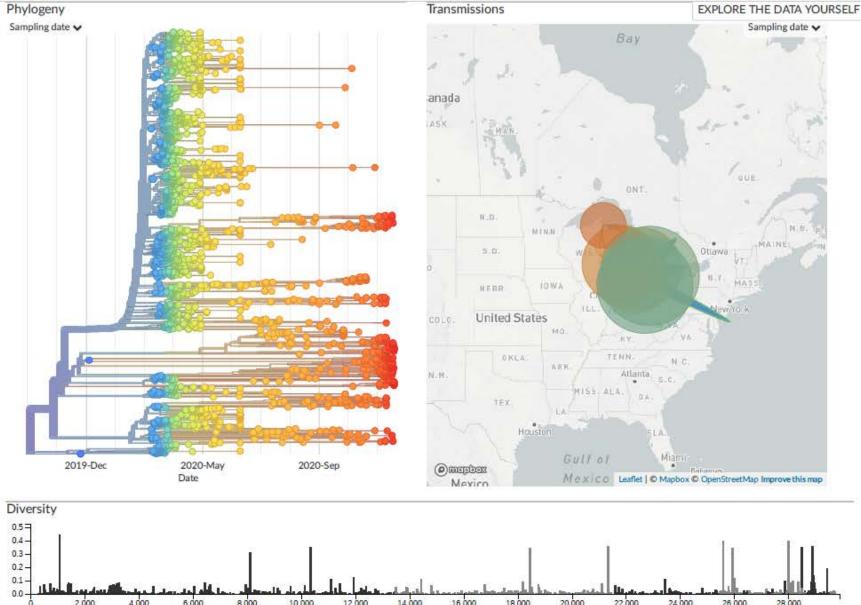
Overlaying Epidemiological and Genomic Data

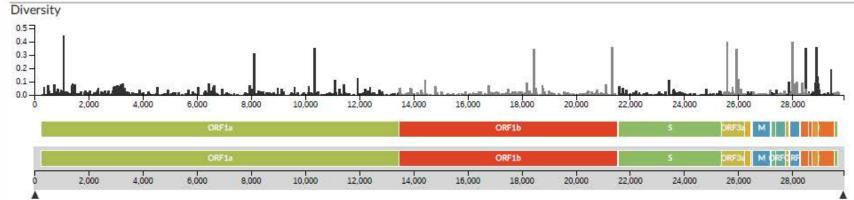
SARS-CoV-2 Report



Summary

We have currently sequenced 2528 SARS-CoV-2 genomes since the first case was detected in Michigan. The first few pages will include the background information of SARS-CoV-2 sequencing. Updates and new genomic information is presented starting on page 6.

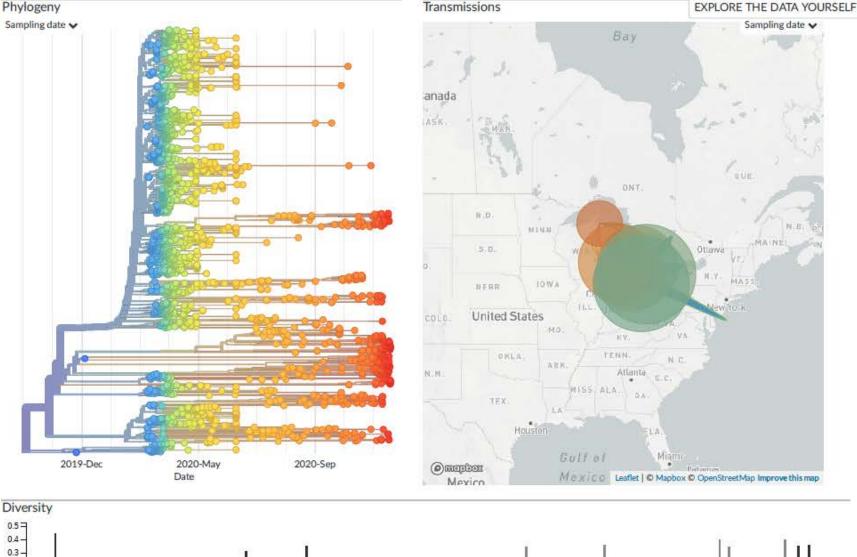


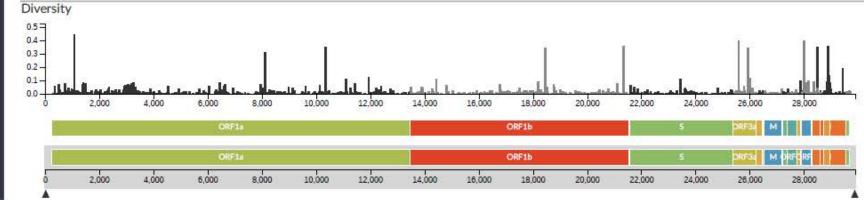


SARS-CoV-2 Sequencing Purpose

The purpose of analyzing the viral genomes are to:

- To understand the viral biology to help inform future scientific paths (potential vaccine development, identify environmental host and source of the virus, and general viral dynamics).
- To monitor for clinically important variants (assess association of clinical outcomes with different strain variants or identify variants that may disrupt diagnostic test targets).
- To inform public health action for future "flareup" or potential seasonality (transmission networks).
- To establish national and regional baselines to assess the effectiveness of response strategies.
- To examine local and national transmission dynamics of the virus for immediate public health tracking in site specific areas.





SARS-CoV-2

SARS-CoV-2 is classified as a coronavirus and the etiological agent responsible for COVID-19 infections. Coronaviruses are not a novel family of viruses and includes other known viruses such as, MERS and SARS, in addition to the common cold viruses that circulate each year. The nucleic acid material for these viruses are RNA encoded (instead of DNA).

In order for viruses to continue to spread and replicate, they have to infect human cells so that they can utilize the enzymes and machinery of the host cell for replication. The machinery (RNA polymerase) that is responsible for copying the RNA is error prone and lacks the ability to proofread and edit the mistakes. This step can lead to the accumulation of mutations in the viral genomes.

Not all mutations that are generated will be beneficial for the virus. Three events can occur from a mutation that is generated:

- · the mutation causes the virus to no longer be viable these viruses will most likely never be isolated
- the mutation has no effect on the virus these viruses may be isolated from multiple people, redundancy in the genetic code allows for mutations to occur and not impact the protein structure or activity
- o the mutation will allow the virus to spread among the population and may result in a virus that is more pathogenic

For some of the mutations that occur, we are still unsure of the associations between these mutations and clinical outcomes or presentation of symptoms. We are also unable to identify whether a specific mutation will result in a more pathogenic strain or change the disease severity without performing the additional molecular epi analysis using a complete study population, even then, we will face limitations that may impact the associations that we see with various mutations and outcomes.

Demographics

Discrepancies may be seen for the total metadata and genomic sequences, due to sequences that pass QC but are lower quality being filtered out of the phylogenetic analysis or duplications of samples.

Gender

total (n=2528)	gender
1186	female
1314	male
26	unknown

Age

total (n=2528)	age
75	0-18 years
1499	19-65 years
950	>66 years
4	unknown

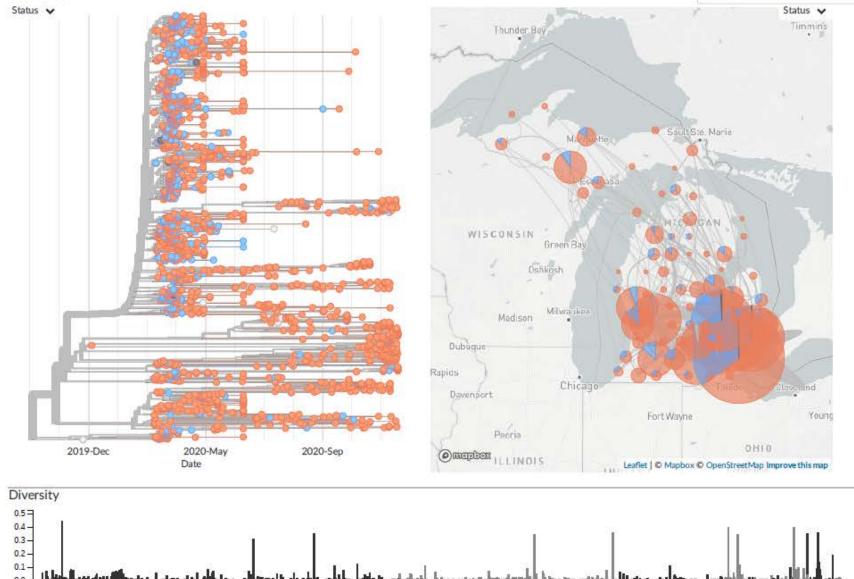
Regional Distribu on

total (n=2224)	region
254	Region 1
593	Region 2N
555	Region 2S
389	Region 3
150	Region 5
349	Region 6
68	Region 7
126	Region 8

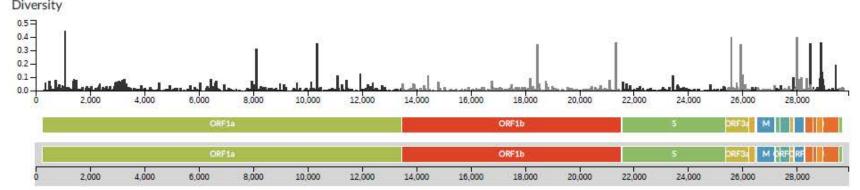
Phylogeny

Status of Individual

Overlaying the patient status on the phylogenetic tree gives no indication of a specific clade or cluster associated with deceased patients.



Transmissions



EXPLORE THE DATA YOURSELF

- Questions?

Center for Surveillance, Epidemiology, and Laboratory Services

SARS-CoV-2 Variants Update

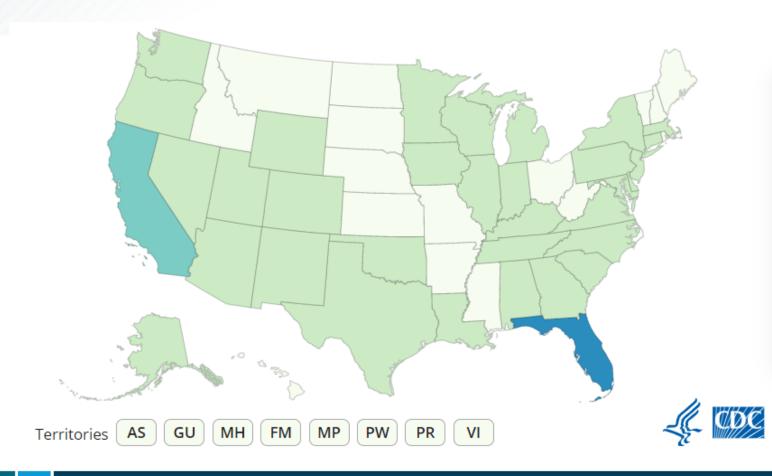
Vivien Dugan

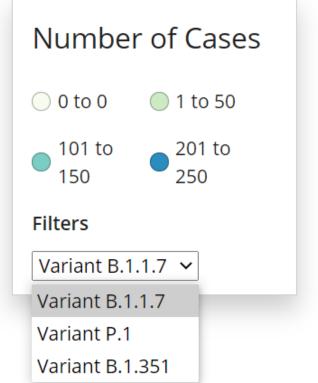
CDC Laboratory and Testing Task Force for the COVID-19 Response



US COVID-19 Cases Caused by Variants

https://www.cdc.gov/coronavirus/2019-ncov/transmission/variant-cases.html





Center for Surveillance, Epidemiology, and Laboratory Services

Vaccine Breakthrough Case Investigations

Leisha Nolen

CDC COVID-19 Response Vaccine Breakthrough Team



These slides were shared during the call, but are not available for public distribution.

Center for Surveillance, Epidemiology, and Laboratory Services

FDA Update

Tim Stenzel

U.S. Food and Drug Administration (FDA)



U.S. Food and Drug Administration (FDA)

COVID-19 Emergency Use Authorization (EUA)
 Information for Medical Devices

https://www.fda.gov/medical-devices/emergencysituations-medical-devices/emergency-useauthorizations

COVID-19 In Vitro Diagnostic EUAs

https://www.fda.gov/medical-devices/coronavirus-disease-2019-covid-19-emergency-use-authorizations-medical-devices/vitro-diagnostics-euas

COVID-19 Frequently Asked Questions

https://www.fda.gov/emergency-preparedness-and-response/coronavirus-disease-2019-covid-19/coronavirus-disease-2019-covid-19-frequently-asked-questions

COVID-19 Updates

https://www.fda.gov/emergency-preparedness-and-response/mcm-legal-regulatory-and-policy-framework/emergency-use-authorization#2019-ncov

FDA Townhall Meetings

https://www.fda.gov/medical-devices/workshopsconferences-medical-devices/virtual-town-hall-seriesimmediately-effect-guidance-coronavirus-covid-19diagnostic-tests-06032020

 Independent Evaluations of COVID-19 Serological Tests

https://open.fda.gov/apis/device/covid19serology/



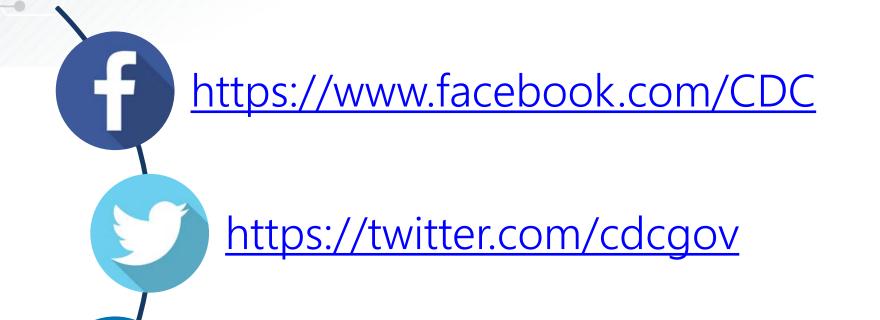
U.S. Food and Drug Administration (FDA)

- COVID-19 Diagnostic Development
 CDRH-EUA-Templates@fda.hhs.gov
- Spot Shortages of Testing Supplies: 24-Hour Support Available
 - 1. Call 1-888-INFO-FDA (1-888-463-6332)
 - 2. Then press star (*)
- FDA MedWatch

https://www.fda.gov/safety/medwatch-fda-safety-information-and-adverse-event-reporting-program

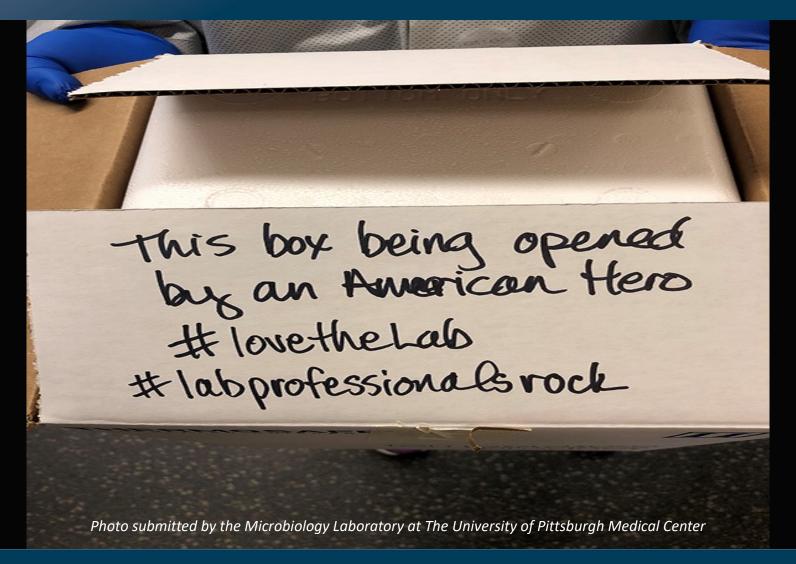


CDC Social Media



https://www.linkedin.com/company/cdc

Thank You For Your Time!



Division of Laboratory Systems Excellent Laboratories, Outstanding Health