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Viral RNA Standards Harmonization:

Data Review: Trends, Anomalies, Summaries

Coronavirus Standards Working Group

# What should a Coronavirus Standards Working Group do?



Assure development and availability of standards, controls, interlab testing, knowledge to support successful rollout & scaling of 2019-nCoV testing



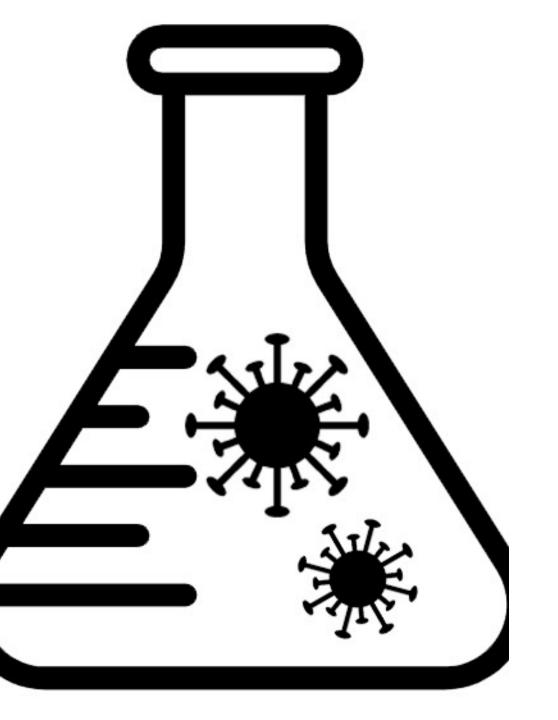
Identify and develop critical infrastructure to support...

confidence in test results interoperability scale-up long-term capacity



Identify best practices that should be institutionalized

Learn what we need to so next time we have a global network in place ready to make standards.



## Agenda

## **Updates**

 Standards for Genomic Surveillance

# Viral RNA Harmonization Study

• Data Review

### Viral RNA Harmonization Study Data

- Data in from 13/14 labs
- Analysis Dashboard pretty refined
- Feedback from a few folks has been really helpful

#### CSWG Viral RNA Harmonization Study Status

	NIST	NML/LGC	NIB (Slovenia)	Bio-Rad	Western	MUSC	Мауо	Labcorp	Quest	Biogazelle	MassCPR Diagnostics	Stanford Medicine	Los Alamos	biodesix
Panel Received	<b>~</b>	~	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>	$\checkmark$	<b>~</b>	$\checkmark$	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>
Lab Metadata Entry Initiated	<b>~</b>	~	$\checkmark$	$\checkmark$	<b>~</b>	$\checkmark$	$\checkmark$	~	<b>~</b>	<b>~</b>	~	$\checkmark$	~	<b>✓</b>
Lab Metadata Entry Complete	<b>~</b>	~	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	<b>✓</b>		<b>✓</b>	$\checkmark$	$\checkmark$	<b>~</b>	$\checkmark$	$\checkmark$
Lab Data Received	<b>~</b>	~	$\checkmark$	$\checkmark$	<b>~</b>	<b>~</b>	<b>✓</b>		<b>✓</b>	$\checkmark$	~	<b>✓</b>	<b>✓</b>	$\checkmark$
Data Summarized	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>		<b>~</b>	<b>~</b>	<b>~</b>	<b>✓</b>	<b>~</b>	<b>~</b>
Data Analyzed	<b>~</b>	~	~	<b>~</b>	~	<b>~</b>	<b>~</b>		<b>~</b>	~	<b>~</b>	<b>~</b>	<b>~</b>	<b>~</b>

Pending analysis refinements I know about...



is there a concentration value for results, or is it simply IU (or IU/unit volume)?



how do we summarize multiple targets from a laboratory?

does it matter?



can we identify trends in results

extraction effects from different labs?

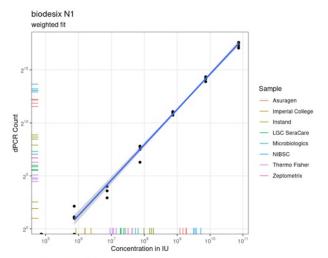


how do we estimate measurement uncertainty?

#### CSWG RNA Harmonization Study

#### Calibration Curves





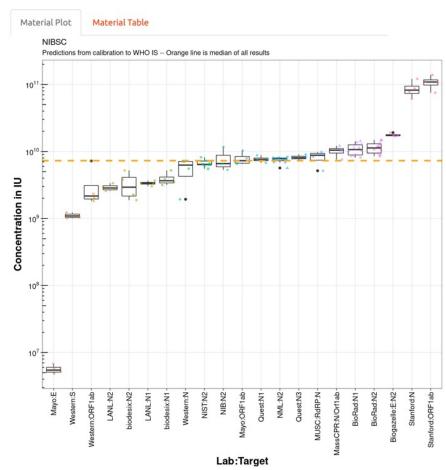
#### Calibration Curve Fit

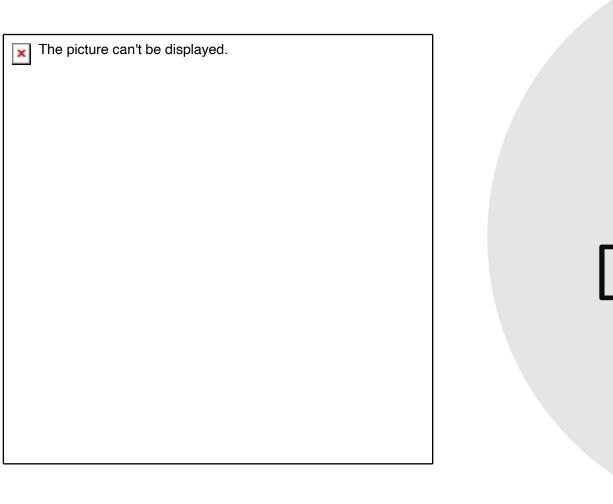
	logSig						
Predictors	Estimates	CI					
(Intercept)	-18.71	-19.94 – -17.49					
logConc	3.33	3.20 - 3.46					
Observations	20						
R <sup>2</sup> / R <sup>2</sup> adjusted	0.994 / 0.994						

#### Material Results









# Discussion