6 November 2020 Marc Salit, JIMB Director SLAC National Lab

SLAC National Lab Stanford University Harmonization Study Planning – Design and Recruiting 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.



Finalizing our Harmonization Study



Study design



Protocols



Samples



Labs







Logistics



Timeline

Reporting

Analysis

Purpose of Harmonization Study

The CSWG "Harmonization Study" will establish the equivalence of SARS-CoV-2 RNA target concentrations across a panel of materials and calibrate those results against the candidate WHO International Standard (IS) reference sample.

By calibrating with the NIBSC sample intended to establish the International Unit (IU), the values on the materials included in this study can be assert traceability to the IU when it becomes available.



Teams: Samples, Labs, Reporting, Analysis Plan



CSWG Harmonization Study Design



Open design and protocol questions



- How to address results from labs who are performing "Yes/No" qualitative tests
- Replication design to assess variability of lab measurements
 - do the 4 different samples cover this sufficiently?
- Do we include "No-template" controls
 - how do these fit in the replication design?
- Dilution scheme to get samples in range for lab's method
- Other questions?

Samples Team: Recruiting Plan



Need 64 identical samples at a single high nominal concentrations that can be diluted by the testing laboratory,

16 negative controls diluted in the same VTM

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Samples to include 3 suppliers each of...

Inactivated, cultured SARS-CoV-2 virus Recombinant virus Recombinant bacteriophage ('armored' RNA)

Unblinded, nominal values provided

Provide COA and Instructions for Use



When possible, use catalog products as they would be provided to a user



Manufacturers will send packaged materials to JIMB and JIMB will distribute to test laboratories

Labs Team: Recruiting Plan

- Lab will receive 4 tubes each of about 10 samples
 - full-process materials in VTM
- Samples clearly labeled
 - nominal abundance of viral RNA
 - "Certificate of Analysis" and "Instructions for Use"

- Lab will report results for each tube
 - reporting tool will be provided
 - results and a MIQE-compliant annotation & protocol
- Recruiting
 - Test Developers
 - Clinical Labs
 - National Measurement Labs



Logistics --David Catoe

Team formed for Design, Analysis, and Reporting



No embargo -- data released as soon as we've compiled it and established consistency Data reported with MIQE and emerging digital-MIQE annotation

> Dashboard-like interactive analysis to permit establishing traceability of new samples to WHO IS

Open data in open repository

Principles

Timeline



Discussion

What this study is not going to do



a comparison of tests



a comparison of labs



a survey of method performance (LOD, precision, repeatability)



an evaluation of commutability

We can make the standards to make molecular testing robust, reliable, and quantitatively comparable.



'Harmonization Kit" to establish comparability of a set of standards to put molecular testing results on a common scale "Benchmarking Kit" for turn-key evaluation of molecular testing platforms

just a few labs, NMIs



test developers



"Validation Kit" for blinded validation with a dashboard to form a "smart-grid" for testing

> routinely measured at testing labs

