

Meeting Minutes for the VQA Advisory Board Conference Call

April 16, 2021

Taken by Miranda Carper and Terese Camp, Duke VQA

Attendees:

VQAAB Voting Members			
☑ Bill Meyer (Chair)☑ Joan Dragavon (ACTG/HVTN)☑ Belinda Yen-Lieberman (ACTG)	☑ Robert Coombs (ACTG/HVTN)☑ Urvi Parikh (MTN)☐ Grace Aldrovandi	☑ Nicole Tobin (IMPAACT)☑ Jessica Fogel (HPTN)☑ Marco Schito	
VQAAB Non-Voting Members			
⊠Ron Bosch (SDAC) ☐ Saira Huq (NICHD)	☑ Diane Costello (IMPAACT/ACTG)	⊠ Fatima Jones (NICHD)	
HANC			
⊠Tyler Brown			
NIAID/DAIDS			
⊠Joe Fitzgibbon	⊠Keith Crawford		
Virology Quality Assurance / Duke Human Vaccine Institute			
☐Thomas Denny (PI)	⊠Raul Louzao	⊠Terese Camp	
⊠Bhavna Hora	⊠Sal Scianna	⊠Heidi Register	
☐Wes Rountree	⊠Katelyn Xiang	☐ Marcella Sarzotti-Kelsoe	
⊠ Kristen Skinner	☐ Chris Todd	⊠Miranda Carper	
⊠Zach Korzen-Varin	⊠Yunfei Wang		
Virology Quality Assurance Sub-Contractor			
⊠Mike Busch (Vitalant Research I	nstitute)		

Action Items

- VQA add number of labs participating for each PT to Mean Log₁₀ Recovery by Kit Type Figure
- VQA investigate if amplification of GEN2020_11 sample 4 is due to a kit lot issue. Send results of
 investigation to VQAAB and request their vote off-line

Meeting Minutes

- Opening Remarks
 - o Bill Meyer and Tom Denny
- Roll Call
 - o Miranda Carper
 - 8 voting members were present
- Review of Data
 - o RNA2021_02
 - 73 datasets from 53 laboratories were included in the analysis



- 3 datasets received a score of PC
- 1 dataset received a score of P
- Potential Issue Alerts were not released for this PT
 - Change in panel samples impacted labs; particularly those running RTM assays. Continue to monitor performance in upcoming PTs
 - RTM labs had higher standard deviation
- Change in Status
 - Two laboratories are reapproved for testing
 - One laboratory had a decline is status from approved to provisionally approved for testing
- VQA reviewed graphs showing the Mean Log10 Recovery by Kit Type for all VQA
 PT data and for VQA200 copy control data.
 - Action: Add number of labs for each kit type to graphs
- VQAAB voted and approved the scores calculated by the VQA
- o GEN2021_11
 - Call for vote to remove sample four from analysis
 - Several laboratories had issues amplifying sample 4 (HIV-1 Subtype C) for PR/RT; Majority of the labs that had issue were those using the ViroSeq assay (see Table 1 below)
 - Summary of Review of data and quality control testing
 - Samples were tested using in-house NGS, in-house Sanger and ViroSeq assay
 - VQA sequencing core did not have issues amplifying any of the samples
 - Phylogenetic tree analysis was performed on the data, no other issues were identified with sample 4
 - Advisory Board asked if the issues amplify sample 4 is a kit lot issue
 - Advisory Board Advice: If the problem is not due to the kit, then don't change the scores. If it is a kit lot issue the sample should be removed
 - Action VQA: investigate if this is a kit lot issue and provide the information to the VQAAB to vote via email
 - Post Meeting Note: All five laboratories that had issues amplifying sample 4 were using the same ViroSeq kit lot (50422). The VQAAB voted to remove sample for from analysis due to technical reasons (kit lot issue for ViroSeq). Sample 4 was removed for PR/RT and new reports were created and provided to the laboratories. Sample 4 remained in in INT analysis.
- Closing statement
 - VQAAB Annual Review
 - o On-site September 20, 2021



Table 1: Summary of laboratories that were unable to amplify sample 4

Number of PR/RT datasets	47
Number of labs unable to amplify sample #4 PR/RT	7
% of labs had problems with # 4 PR/RT	14.89%
Number of VS labs unable to amplify sample #4 PR/RT	5
Total number of VS labs PR/RT	9
% of VS labs had a problem with # 4 PR/RT	55.56%
% of labs had problems with # 4 PR/RT	14.89%
Number of labs that would have a score change if remove sample 4	
from analysis	2 PR/RT, 1 INT