



National Laboratory for HIV Reference Services
Sexually Transmitted and Bloodborne Infections
National Microbiology Laboratory
Public Health Agency of Canada

HIV Serology Quality Assessment Program Summary for Panel HIVSER 2022Nov14

2022Nov14 HIV Serology Panel		
Panel Sample	True Status	Labs Reporting Incorrect Status
A	HIV-1 Ab Positive	HV31
B	HIV-1/2 Ag/Ab Negative	
C	HIV-2 Ab Positive	HV31
D	HIV-1/2 Ag/Ab Negative	
E	HIV-2 Ab Positive	HV31

Summary of findings observed for the 2022Nov14 panel:

- 1) Participant HV31 incorrectly selected "HIV-1 Positive" for Sample A, C and E (reactive result using only the bioLytical HIV-1/2 INSTI Rapid test).
- 2) Participant HV23 was not able to return results by the submission due date.



National Laboratory for HIV Reference Services
Sexually Transmitted and Bloodborne Infections
National Microbiology Laboratory
Public Health Agency of Canada

HIV Serology Quality Assessment Program

Final Report for Panel HIVSER 2022Nov14

Issued 2023-February-07

Introduction

The NLHRS distributed the 2022Nov14 panel and the 2023Apr19 panel on October 26, 2022. This final report is specific to the 2022Nov14 panel only and is publicly available; however, the identity of participants are not disclosed. The deadline for results submission was November 14, 2022. The preliminary report was issued on December 02, 2022.

Panel Samples, HIV Test Kits, and Data Entry

- *Panel Composition:*
 - The 2022Nov14 panel consisted of five samples: two HIV negative (B, D), one HIV-1 Ab positive (A) and two HIV-2 Ab positive (C, E). Samples A and E were diluted 1 in 2 with defibrinated human plasma (Basematrix 53, Seracare Life Sciences). Testing and characterization of the panel by the NLHRS prior to shipment are presented in Appendix 2. Panels were sent to 42 participants including the NLHRS on October 26, 2022.
 - The metrological traceability and uncertainty is not applicable for this panel.
- *HIV Test Kits*
 - Ten different assays were used by 42 participants (including the NLHRS) who returned results (Appendix 3).
- *Data entry*
 - Results entry for this panel utilized an NML developed website.

Homogeneity and stability

- The homogeneity and stability of the 2022Nov14 HIV serology panel was assessed by comparing the participants' results with the panel characterization results obtained by the NLHRS prior to the panel send-out.
- There was no indication of heterogeneity or instability of the panel samples as the data submitted by the participants was consistent with the expected results from the NLHRS characterization of each panel member (Figures 1, 2, and Appendix 2).

Results

- *Evaluation Criteria:*

- Negative samples: HIV non-reactive/negative in the final HIV serology interpretation with assay results supporting the final serology interpretation.
- Positive samples: HIV reactive/positive in the final HIV serology interpretation with assay results supporting the final serology interpretation. Participants must provide a recommendation for further action for samples that they could not determine the true serology status for based on the assay used in their testing.

- *Qualitative Group Analysis (Figures 1 and 2)*

- *Sample A (HIV-1 Ab Positive)* – 41/42 participants (including NLHRS) provided either a correct serology status and/or recommendation.
- *Sample B (HIV-1/2 Ag/Ab Negative)* – 41/42 participants (including NLHRS) provided either a correct serology status and/or recommendation.
- *Sample C (HIV-2 Ab Positive)* – 41/42 participants (including NLHRS) provided either a correct serology status and/or recommendation.
- *Sample D (HIV-1/2 Ag/Ab Negative)* – 41/42 (including NLHRS) participants provided either a correct serology status and/or recommendation.
- *Sample E (HIV-2 Ab Positive)* – 41/42 (including NLHRS) participants provided either a correct serology status and/or recommendation.

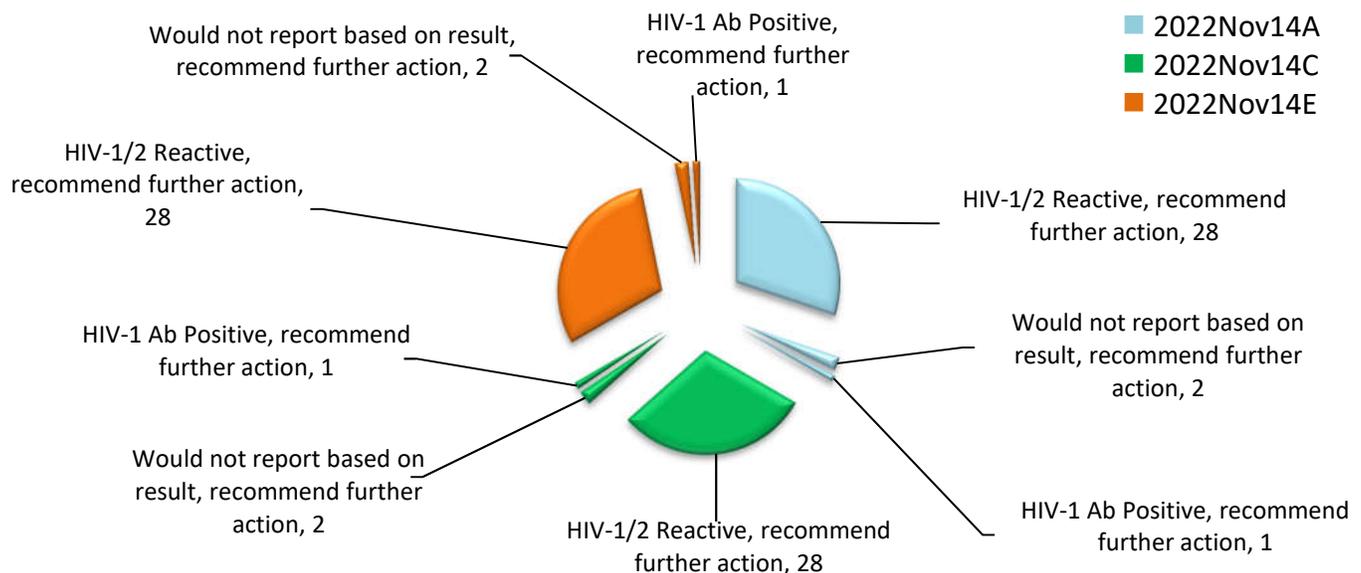


Figure 1: The final HIV serology status of the positive samples in the 2022Nov14 HIV serology panel submitted by participants using an HIV screening assay.

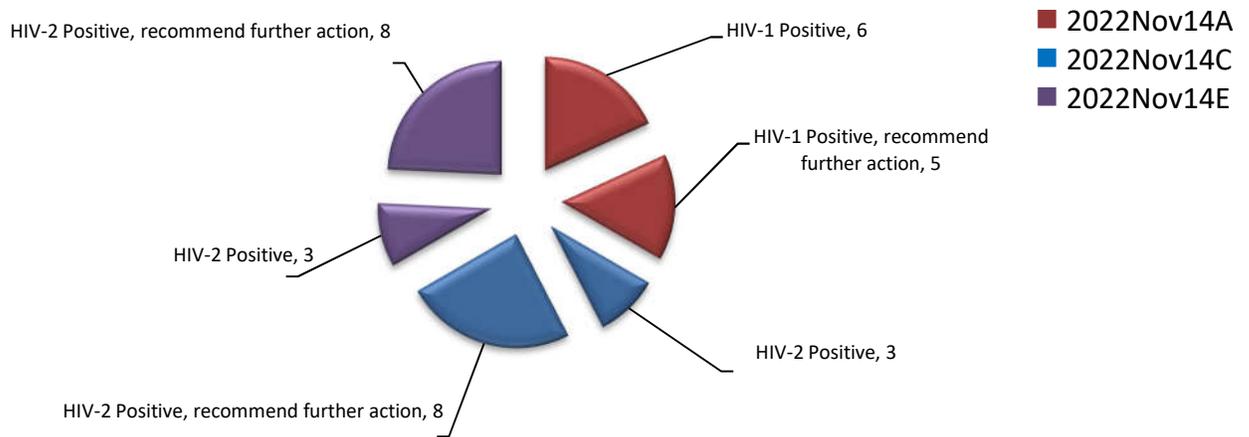


Figure 2: The final HIV serology status of the positive samples in the 2022Nov14 HIV serology panel submitted by participants (including NLHRS) using HIV screening and confirmatory assays.

Findings

The majority of participants correctly identified the serology status and/or provided an appropriate recommendation for the panel samples included in the 2022Nov14 test event.

One user of the bioLytical HIV-1/2 INSTI® Rapid Test made an error when selecting the final interpretation. This participant selected “HIV-1 Positive” for Sample A, C, and E. Because the bioLytical HIV-1/2 INSTI® Rapid Test is a screening test only, “HIV-1/2 Reactive” is the correct final interpretation to select when submitting a result for a sample that tested reactive. It is incorrect to select “HIV-1 Positive” as the final interpretation in this instance. One participant was not able to return results by the submission due date.

Since the 2021Apr19 test event, we have noticed several of the Abbott Architect users adopting the newer Abbott Alinity platform. In total, 11 users have switched over to the Abbott Alinity platform. Also, four other Abbott Architect users have switched over to other platforms: the Roche Elecsys (2), the Siemens Atellica (1), and the Ortho Clinical Diagnostics VITROS (1). In summary, of the 26 participants that have used the Abbott Architect, 15 had switched to other platform. We will continue to monitor if this trend continues in future events.

In closing, we value each laboratory’s participation in these QA test events and your suggestions for improvement. The NLHRS is committed to improve all aspects of the HIV serology proficiency-testing program in order to provide quality proficiency testing to our participants.

If you have any comments, suggestion or concerns, please contact us at:

nlhrs.qap-peq.insrv@phac-aspc.gc.ca

Thank you for your participation in the NLHRS HIV Serology QA Program



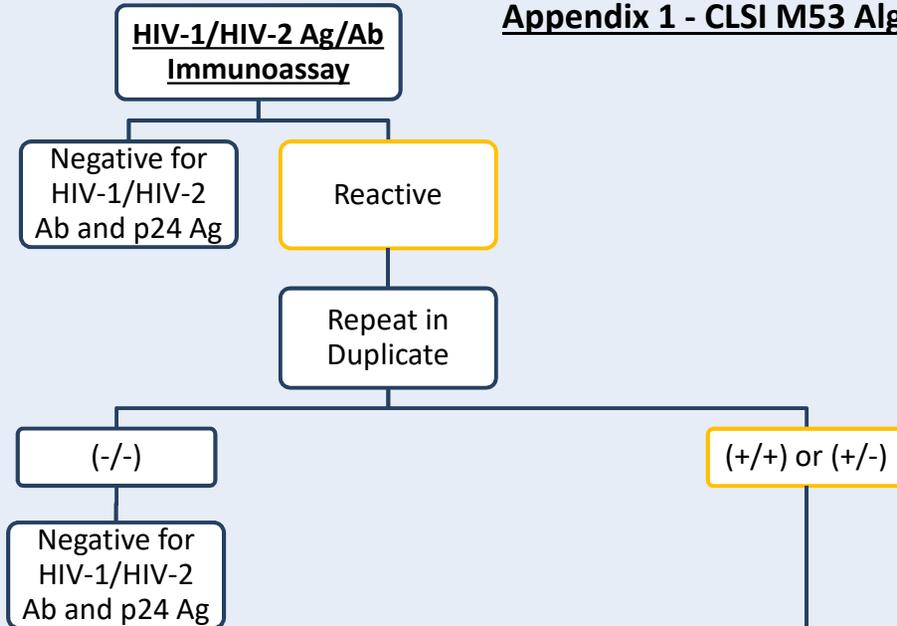
John Ho
Quality Assurance Program Coordinator
National Laboratory for HIV Reference Services
Public Health Agency of Canada
Tel: (204) 789-6518



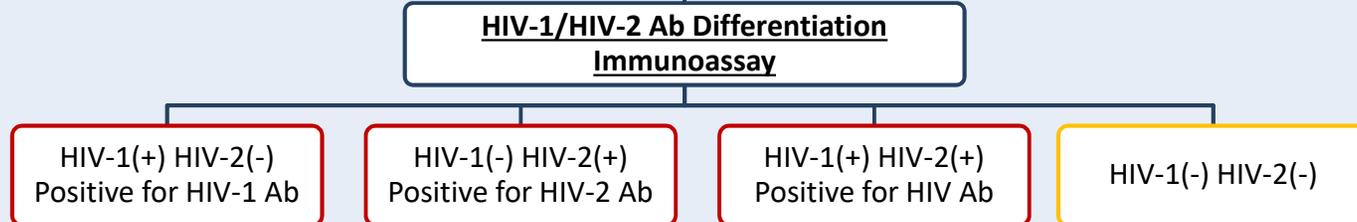
Dr. John Kim
Laboratory Chief
National Laboratory for HIV Reference Services
Public Health Agency of Canada
Tel: (204) 789-6527

Appendix 1 - CLSI M53 Algorithm I

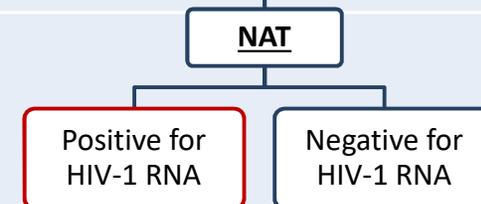
(i) HIV-1/HIV-2 Ag/Ab Immunoassay



(ii) HIV-1/HIV-2 Ab Differentiation Immunoassay



(iii) Nucleic Acid Testing

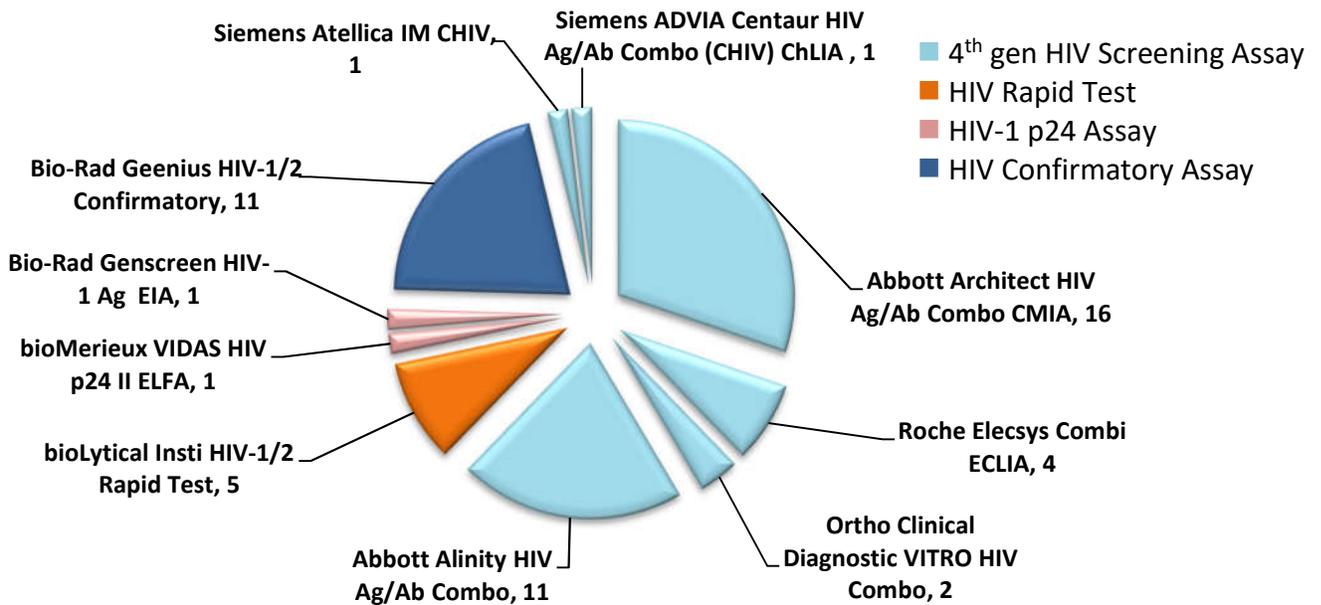


Appendix 1: Adaptation of the Clinical and Laboratory Standards Institute (CLSI) M53-*Criteria for Laboratory Testing and Diagnosis of Human Immunodeficiency Virus Infection: Approved Guideline* Algorithm I.

Appendix 2: Summary of NLHRS characterization of the 2022Nov14 HIV serology panel samples

Sample		B/D (Duplicate)	A	C	E
Final HIV Status		HIV-1/2 Ag/Ab Negative	HIV-1 Ab Positive	HIV-2 Ab Positive	HIV-2 Ab Positive
bioLytical INSTI® HIV-1/2 Rapid Test	Result	Non-Reactive	Reactive	Reactive	Reactive
Bio-Rad GS HIV p24	Result	Non-Reactive	Non-Reactive	Non-Reactive	Non-Reactive
Bio-Rad GS HIV p24 Confirmatory	Result	Not Tested	Not Tested	Not Tested	Not Tested
Fujirebio INNO-LIA HIV-I/II Score	Result	Negative	HIV-1	Not Tested	HIV-2
	sgp120	-	++	N/A	-
	gp41	-	+++	N/A	-
	p31	-	++	N/A	++
	p24	-	++	N/A	+
	p17	-	+	N/A	-
	sgp105	-	-	N/A	++
gp36	-	-	N/A	++	
Bio-Rad Geenius HIV-1/HIV-2 Supplemental Assay	Result	Negative	HIV-1	HIV-2	HIV-2
	gp36	-	-	+	+
	gp140	-	-	+	+
	p31	-	-	+	+
	gp160	-	+	-	-
	p24	-	-	-	-
	gp41	-	+	-	-
CTRL	+	+	+	+	

Appendix 3: Summary of assays used by the participants in the 2022Nov14 HIV serology test event



Appendix 4: Summary of bands detected for samples A, C, and E by the Bio-Rad Geenius HIV-1/2 confirmatory assay in the 2022Nov14 HIV serology test event (including NLHRS)

Bio-Rad Geenius	Frequency of Bands Detected						
	gp36	gp140	p31	gp160	p24	gp41	CTRL
2022Nov14A	-	-	-	11	4	11	11
2022Nov14C	11	11	11	-	-	-	11
2022Nov14E	11	11	11	-	-	-	11

Appendix 5: Troubleshooting

Troubleshooting; common causes of outlying and/or aberrant results in serology and molecular Laboratories.

Type of Error	Possible Cause(s)	Pre-Analytical	Analytical	Post- Analytical
Sample mix-up	Can occur during specimen reception or testing. May result in outlying/aberrant results for one or all samples mixed-up.	✓	✓	
Transcription	• Incorrect test ordering by physician	✓		
	• Incorrect shipment address	✓		
	• Selecting the wrong assay for data entry	✓		
	• Interchanging results for two or more specimens			✓
	• Entering incorrect results			✓
	• Entering values in the incorrect field (e.g., OD as S/Co)			✓
	• Entering values in the incorrect unit (e.g., IU/mL instead of log ₁₀ copies/mL)			✓
	• Using a comma instead of a dot to denote a decimal point			✓
	• Selecting the incorrect assay interpretation or analyte			✓
	• Failure to recommend follow-up testing where necessary			✓
	It is recommended all results that are manually transcribed or entered electronically be checked by a second individual to avoid transcription errors.			
Outlying and/or Aberrant Results (<u>random error</u>)	<u>Sporadic test results identified as outlying and/or aberrant can be classified as random events. Possible causes of random error include:</u>			
	• Incorrect sample storage/shipping conditions	✓	✓	
	• Incorrect test method	✓	✓	
	• Insufficient mixing of sample, especially following freezing		✓	
	• Poor pipetting		✓	
	• Ineffective or inconsistent washing		✓	
	• Transcription errors	✓		✓
	• Cross-contamination or carryover	✓	✓	
• Presence of inhibitors to PCR		✓		
Outlying and/or Aberrant Results (<u>systematic error</u>)	<u>A series of test results identified as outlying and/or aberrant may be due to a systematic problem. Systematic problems may be due to:</u>			
	• Reagents contaminated, expired, or subject to batch variation		✓	
	• Instrument error or malfunction		✓	
	• Insufficient washing		✓	
	• Incorrect wavelength used to read the assay result		✓	
	• Cycling times too long/short or temperature too high/low		✓	
	• Incubation time too long/short or temperature too high/low		✓	
	• Insufficient mixing/centrifuging before testing		✓	
	• Incorrect storage of test kits and/or reagents	✓		
	• Contamination of master-mix, extraction areas or equipment		✓	
	• Ineffective extraction process		✓	
	• Degradation of master-mix components		✓	
• Suboptimal primer design (in-house assays)		✓		

This table was modified from a report produced by the National Reference Laboratory (NRL), Melbourne, Australia.