UCSF UC San Francisco Previously Published Works

Title

Design and Implementation of an External Quality Assessment Program for HIV Viral Load Measurements Using Dried Blood Spots

Permalink https://escholarship.org/uc/item/807257fw

Journal Journal of Clinical Microbiology, 53(3)

ISSN 0095-1137

Authors

Prach, Lisa M Puren, Adrian Lippman, Sheri A <u>et al.</u>

Publication Date

2015-03-01

DOI

10.1128/jcm.02698-14

Peer reviewed



Design and Implementation of an External Quality Assessment Program for HIV Viral Load Measurements Using Dried Blood Spots

Lisa M. Prach,^a Adrian Puren,^{b,c} Sheri A. Lippman,^a Sergio Carmona,^c Sophie Stephenson,^a Ewalde Cutler,^{b,c} Scott Barnhart,^d Teri Liegler^a

University of California San Francisco, San Francisco, California, USA^a; National Institute for Communicable Diseases/NHLS, Johannesburg, South Africa^b; University of Witwatersrand, Johannesburg, South Africa^c; University of Washington, Seattle, Washington, USA^d

An external quality assurance program was developed for HIV-1 RNA viral load measurements taken from dried blood spots using a reference panel and field-collected specimens. The program demonstrated that accurate and reproducible quantitation can be obtained from field-collected specimens. Residual proviral DNA may confound interpretation in virologically suppressed subjects.

IV-1 RNA viral load (VL) measurements from blood plasma are critical for assessing the response to antiretroviral treatment (ART) in individuals and populations (1–5). Sample preparation and storage, however, require laboratory and cold-chain infrastructure that may be limited when monitoring rural populations in resource-limited countries. These populations are often the most affected by the global pandemic. The collection of fingerprick blood on filter paper as dried blood spots (DBS) is an alternative strategy frequently used in global surveillance studies (6, 7). External quality assurance (EQA) programs for VL testing rely largely on blood plasma and do not effectively assess the unique qualities of the DBS sample matrix, including lower virion input copy number, inconsistent spot size, and the inclusion of proviral DNA, cellular RNA, and virion RNA.

To assess the quality of the VL measurements performed by a designated testing laboratory from DBS collected in a rural field setting for an HIV prevention program in North West Province, South Africa (8), an EQA program was developed and implemented that included two phases. First, to test overall concordance, 50 reference DBS cards were generated in duplicate at a 3rd-party laboratory under standardized conditions using venous EDTA blood (50 µl spots on Munktell filters; Ahlstrom Munktell, Helsinki, Finland) from HIV-seronegative and -seropositive donors with known plasma VL (Abbott RealTime HIV-1 viral load assay; Abbott Diagnostics, Des Plaines, IL, USA; lower limit of detection [LLOD], 40 copies/ml). The cards were dried at ambient temperature for 1 to 3 days and stored at -80° C in zip-lock bags with desiccant and humidity indicator cards for 3 months, followed by ambient shipment (3-day transit time) to the study testing and reference laboratories. Total HIV nucleic acid was measured at both laboratories from a single DBS using the Cobas AmpliPrep/Cobas TaqMan HIV-1 2.0 test (Roche Applied Sciences, Pleasanton, CA, USA; LLOD, 400 copies/ml). The HIV serostatuses and plasma VL of the donors that were used for the reference DBS (to which the testing and reference labs were blinded) were as follows: HIV seronegative, n = 10; HIV seropositive with plasma VL of <40 copies/ml (1.6 \log_{10}), n = 14; HIV seropositive with plasma VL of \geq 40 copies/ml, n = 25; and detectable plasma VL of unknown quantity, n = 1.

Second, following complete testing and evaluation of the reference cards, DBS cards (Munktell) were collected by finger-prick from the study participants, with informed consent, in three field sites. Blood was collected from 179 seropositive donors and confirmed by serial rapid testing or HIV DNA PCR. The DBS cards were transported in a cooler box to the field office, dried overnight, and packed in zip-lock bags with desiccant and humidity indicators. Within 6 days of collection, the cards were stored at ambient temperature until transport to the testing laboratory, where they were stored at -70° C until VL testing was performed. All seropositive samples from the first site and a random 15% of the DBS cards from sites 2 and 3 were transported at ambient temperature from the testing laboratory to the reference laboratory for VL testing.

Forty-nine of 50 (98%) reference cards produced valid results from both laboratories. All 10 HIV-negative samples had undetectable VL (<400 copies/ml) by both laboratories (100% specificity). Of the 26 reference DBS from subjects with detectable plasma viremia (>40 copies/ml; range, 67 to >10,000,000), 24 DBS samples yielded detectable values (≥400 copies/ml) in both labs. One sample (76 copies/ml in plasma) had undetectable VL by both laboratories, and another sample (>10,000,000 copies/ml in plasma) yielded repeated invalid results in one laboratory. Table 1 shows the mean, median, and range of the DBS VL results from the testing and reference laboratories. In a Bland-Altman analysis (Fig. 1A), 100% of the values were within the 95% limit of agreement. A pairwise comparison (Fig. 1C) yielded excellent agreement with a significant correlation (P = <0.0001; Pearson r =0.983; 95% confidence interval [CI], 0.961 to 0.993), with 24/26 (96%) having ≤ 0.5 -log₁₀ difference. The reference samples from HIV-seropositive subjects with undetectable plasma VL (<40

Received 26 September 2014 Returned for modification 17 November 2014 Accepted 10 December 2014

Accepted manuscript posted online 17 December 2014

Citation Prach LM, Puren A, Lippman SA, Carmona S, Stephenson S, Cutler E, Barnhart S, Liegler T. 2015. Design and implementation of an external quality assessment program for HIV viral load measurements using dried blood spots. J Clin Microbiol 53:964–966. doi:10.1128/JCM.02698-14.

Editor: Y.-W. Tang

Address correspondence to Teri Liegler, tliegler@sfgh.ucsf.edu.

Copyright © 2015, American Society for Microbiology. All Rights Reserved. doi:10.1128/JCM.02698-14

TABLE 1 Dried blood spot viral load results from the testing and	
reference laboratories	

Outcomes by sample type	HIV-1 DBS viral load (copies/ml) by laboratory		
	Testing	Reference	
Reference panel samples ^a	<i>n</i> = 25	n = 26	
Minimum	400	400	
Maximum	10,000,000	10,000,000	
Median	89,000	112,000	
Mean	85,000	79,500	
Field collected samples ^b	n = 27	n = 27	
Minimum	400	716	
Maximum	169,939	234,000	
Median	23,988	13,100	
Mean	12,988	12,365	

^{*a*} DBS VL results are shown for samples from HIV-seropositive donors with detectable plasma RNA (plasma VL > 40 copies/ml).

^b DBS VL results shown for samples from HIV-seropositive or indeterminate donors with unknown plasma RNA levels.

copies/ml) had poorer agreement: 3 of 14 (21%) samples had detectable VL by both laboratories, and 5/14 (36%) had undetectable VL by both. Six samples showed discordant results, with 3 each detected in either laboratory (range, 413 to 15,447 copies/ml).

Of the 179 samples collected at the three field sites that tested HIV seropositive or indeterminate, 27 (15%) randomly selected DBS samples were tested at both laboratories, of which 25 had a

VL in the quantifiable range by both laboratories. A Bland-Altman analysis showed that 92% (23/25) of the values were within the 95% limit of agreement (Fig. 1B). Pairwise comparisons of these showed a high correlation (Pearson r = 0.92; 95% CI, 0.788 to 0.956) (Fig. 1D), with 23 (92%) having a ≤ 0.5 -log₁₀ difference.

Overall, these results demonstrate the feasibility of a simple EQA program for DBS VL testing. We demonstrated highly correlative results between two laboratories when testing DBS samples from viremic subjects, despite the various collection conditions and spot sizes inherent in field-collected samples. Although the number of field specimens tested by both laboratories was limited, at 27/174 (15%), the agreement in the DBS VL results from the reference panel and the field-collected specimens lends confidence in the feasibility and accuracy of using DBS for a main study outcome.

However, we also found a high frequency (9/14 [64%]) of reference DBS samples from HIV-positive individuals with undetectable plasma viremia (<40 copies/ml) yielding detectable DBS VL (\geq 400 copies/ml) by either or both laboratories. These results are consistent with other reports and likely reflect the amplification of cell-associated HIV DNA and RNA in DBS samples in the absence of detectable blood plasma RNA (7, 9, 10). The magnitude of this discrepancy may be reduced using DBS preparation procedures that preferentially select for cell-free HIV RNA (11); however, the amplification of residual DNA in DBS to even a minimal extent may confound the interpretation of HIV RNA copy number in subjects with ART-induced virologic suppression. The significant discordance in DBS VL in samples from ART-suppressed

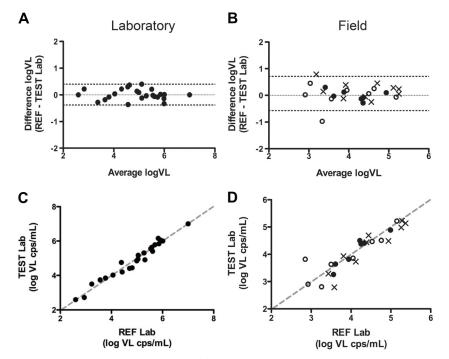


FIG 1 Agreement and correlation of DBS VL results from laboratory- and field-collected samples tested at two laboratories. (A and B) Bland-Altman analysis of \log_{10} HIV VL (copies/ml) results from the reference (REF) (National Institute for Communicable Diseases, Johannesburg, South Africa) and test laboratories (Clinical Laboratory Services, Johannesburg, South Africa) using DBS samples created in duplicate at an independent laboratory (AIDS Research Institute-University of California, San Francisco [ARI-UCSF] Laboratory of Clinical Virology, San Francisco, CA, USA) from infected subjects with detectable plasma viremia (\geq 40 copies/ml) (A), or from field-collected DBS from HIV-seropositive subjects with unknown plasma VL (B). The 95% limits of agreement for the results are -0.38 to 0.40 (A) and -0.57 and 0.71 (B). (C and D) Correlation plot of \log_{10} HIV VL (copies [cps]/ml) results from the reference and test laboratories analyzing DBS samples created in the laboratory (C) or collected from the field (D). The distinct symbols designate samples collected from 3 different study sites.

subjects underscores the limitations in interpreting the RNA copy number using this sample matrix and testing modality. However, in settings where few patients are virally suppressed, this bias may be minimal.

ACKNOWLEDGMENTS

This project was funded by the cooperative agreement 5U2GGH000324-02 from the Centers for Disease Control and Prevention. Its contents are solely the responsibility of the authors and do not necessarily represent the views of the CDC. Additional funding was provided through the University of California San Francisco-Gladstone Institute of Virology & Immunology Center for AIDS Research (CFAR), an NIH-funded program (P30 AI027763).

REFERENCES

- Das M, Chu PL, Santos GM, Scheer S, Vittinghoff E, McFarland W, Colfax GN. 2010. Decreases in community viral load are accompanied by reductions in new HIV infections in San Francisco. PLoS One 5:e11068. http://dx.doi.org/10.1371/journal.pone.0011068.
- Jain V, Liegler T, Kabami J, Chamie G, Clark TD, Black D, Geng EH, Kwarisiima D, Wong JK, Abdel-Mohsen M, Sonawane N, Aweeka FT, Thirumurthy H, Petersen ML, Charlebois ED, Kamya MR, Havlir DV, SEARCH Collaboration. 2013. Assessment of population-based HIV RNA levels in a rural east African setting using a fingerprick-based blood collection method. Clin Infect Dis 56:598–605. http://dx.doi.org/10.1093 /cid/cis881.
- Montaner JS, Lima VD, Barrios R, Yip B, Wood E, Kerr T, Shannon K, Harrigan PR, Hogg RS, Daly P, Kendall P. 2010. Association of highly active antiretroviral therapy coverage, population viral load, and yearly new HIV diagnoses in British Columbia, Canada: a population-based study. Lancet 376:532–539. http://dx.doi.org/10.1016/S0140-6736(10)60936-1.
- 4. Mermin J, Ekwaru JP, Were W, Degerman R, Bunnell R, Kaharuza F, Downing R, Coutinho A, Solberg P, Alexander LN, Tappero J, Campbell J, Moore DM. 2011. Utility of routine viral load, CD4 cell count, and

clinical monitoring among adults with HIV receiving antiretroviral therapy in Uganda: randomised trial. BMJ 343:d6792. http://dx.doi.org/10 .1136/bmj.d6792.

- Kantor R, Diero L, Delong A, Kamle L, Muyonga S, Mambo F, Walumbe E, Emonyi W, Chan P, Carter EJ, Hogan J, Buziba N. 2009. Misclassification of first-line antiretroviral treatment failure based on immunological monitoring of HIV infection in resource-limited settings. Clin Infect Dis 49:454–462. http://dx.doi.org/10.1086/600396.
- 6. Snijdewind IJ, van Kampen JJ, Fraaij PL, van der Ende ME, Osterhaus AD, Gruters RA. 2012. Current and future applications of dried blood spots in viral disease management. Antiviral Res 93:309–321. http://dx .doi.org/10.1016/j.antiviral.2011.12.011.
- Bertagnolio S, Parkin NT, Jordan M, Brooks J, Garcia-Lerma JG. 2010. Dried blood spots for HIV-1 drug resistance and viral load testing: a review of current knowledge and WHO efforts for global HIV drug resistance surveillance. AIDS Rev 12:195–208.
- Lippman SA, Treves-Kagan S, Gilvydis JM, Naidoo E, Khumalo-Sakutukwa G, Darbes L, Raphela E, Ntswane L, Barnhart S. 2014. Informing comprehensive HIV prevention: a situational analysis of the HIV prevention and care context, North West Province South Africa. PLoS One 9:e102904. http://dx.doi.org/10.1371/journal.pone.0102904.
- Monleau M, Montavon C, Laurent C, Segondy M, Montes B, Delaporte E, Boillot F, Peeters M. 2009. Evaluation of different RNA extraction methods and storage conditions of dried plasma or blood spots for human immunodeficiency virus type 1 RNA quantification and PCR amplification for drug resistance testing. J Clin Microbiol 47:1107–1118. http://dx .doi.org/10.1128/JCM.02255-08.
- Parkin NT. 2014. Measurement of HIV-1 viral load for drug resistance surveillance using dried blood spots: literature review and modeling of contribution of DNA and RNA. AIDS Rev 16:160–171.
- 11. Wu X, Baum P, Crask M, Do T, Honisch C, Will S. 2014. A simple method to elute cell-free HIV from dried blood spots improves the ascertainment of virologic suppression. Proceedings of the 20th International AIDS Conference, 20 to 25 July 2014, Melbourne, Australia.