Quantitation of Human Immunodeficiency Virus Type 1 RNA in Plasma by Using Blood Dried on Filter Paper

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We compared whole blood dried on filter paper to the standard assay with frozen cell-free plasma for use in the quantitation of the human immunodeficiency virus RNA load in blood. RNA values from filter paper, corrected for the hematocrit, gave results comparable to those of the standard assay in terms of sensitivity and reproducibility.

The measurement of human immunodeficiency virus (HIV) RNA levels in plasma is rapidly becoming the most important laboratory tool for staging HIV infection and for the management of therapy (6, 7). However, quantitation of HIV levels in plasma requires access to certain laboratory equipment which may not be readily available in some settings. By the year 2000 the World Health Organization predicts that 30 million to 40 million new HIV infections will have occurred, with 90% of these occurring in developing countries (5). Even the simple act of centrifuging a tube of blood, aliquoting and freezing the plasma, and subsequently shipping the plasma to a centralized laboratory might prove difficult in some field situations, where many clinical trials of new antiretroviral drugs and potential HIV vaccines as well as transmission intervention studies will be conducted in the future.

The purpose of this research was to develop a simple method for collecting specimens for analysis of the HIV RNA level in plasma that would yield accurate and reproducible results without the need for electricity at the collection site. Our ultimate goal is to obtain blood from a fingerstick or heelstick, dry it on filter paper, and airmail it to a central laboratory, thus avoiding the need for a skilled phlebotomist and laboratory technician on site and also avoiding the need for centrifuges, freezers, and dry ice for shipping. Dried blood spots have been used for many years to screen for several metabolic disorders such as phenylketonuria and sickle cell disease. For HIV they have been used for the anonymous screening of newborns to assess the seroprevalence of HIV among childbearing women (4) and for the diagnosis of perinatal HIV infections by using HIV DNA (2, 3). More recently, it has been reported that DNA obtained from dried blood spots is suitable for gene sequence analysis (1). Consequently, it has been reported that DNA obtained from dried blood spots is suitable for gene sequence analysis (1). Consequently, we decided to evaluate the possibility of using whole blood dried onto filter paper for the quantitation of the HIV RNA level in plasma.

One hundred four HIV-positive patients seen in the University of North Carolina’s adult and pediatric infectious disease clinics provided 4 ml of blood collected in EDTA for the purposes of viral load testing. HIV RNA was quantitated from 100 μl of whole blood from each of 76 patients spotted in quadruplicate onto Schleicher & Schuell no. 903 filter paper. A total of 50 μl of whole blood from an additional 28 patients was spotted in eight replicates onto Schleicher & Schuell Isocode filter paper. The blood spots were dried overnight at room temperature in a biohazard hood. We also used the whole blood to determine the hematocrit (mean of two determinations). Cell-free plasma obtained from the same tube of blood was stored at −70°C. HIV type 1 (HIV-1) RNA levels were determined by a commercially available nucleic acid sequence-based amplification assay (NASBA; NASBA HIV-1 QT; Organon Teknika, Durham, N.C.). Two 30-μl aliquots of dried blood spots were placed in 9 ml of NASBA lysis buffer, and the mixture was rocked at room temperature for 2 h to isolate the RNA, after which the filter papers were removed. The HIV-1 RNA from cell-free plasma was isolated by placing 100 μl of plasma in 0.9 ml of NASBA lysis buffer. For the rest of the procedure, the manufacturer’s instructions were followed, including the use of diluted calibrators. The numbers of RNA copies per milliliter of plasma from the spots were calculated as follows: correct number of spot RNA copies per milliliter of plasma = (number of spot RNA copies per milliliter of blood)/(100 − hematocrit)/100. Blood from the first 76 patients was tested singly by using the no. 903 filter paper. Blood from the remaining 28 patients was tested in duplicate by using the Isocode filter paper. Some dried spots were stored at room temperature for various lengths of time before testing.

We initially used the no. 903 paper to measure HIV-1 RNA levels from dried blood spots, with encouraging results (Fig. 1A). However, a second lot of paper gave uninterpretable results and we switched to Isocode paper, which is pretreated with guanidinium isothiocyanate. The log_{10} hematocrit-corrected RNA concentration from the two types of paper is compared with the log_{10} RNA concentration from plasma in Fig. 1A (no. 903 filter paper) and Fig. 1B (Isocode filter paper). The diagonal line on each graph indicates where the points would fall if the estimates from the dots and the plasma were the same. Correlation coefficients were 0.88 (no. 903 filter paper versus plasma) and 0.90 (Isocode filter paper versus plasma). There is little evidence of a systematic difference between the values from no. 903 filter paper and those from plasma. The mean difference, 0.0478, was not statistically significantly different from zero (P = 0.33). Most (91%) of the differences in RNA values between the no. 903 filter paper dots and plasma were between −0.5 and 0.5 log_{10}. However, the
mean difference between values from plasma and Isocode filter paper was significantly different from zero (mean = 0.22; \( P < 0.01 \)). On average, then, the values from plasma were 1.66 times the values from the Isocode filter paper. The difference did not vary systematically over the range of the data (Fig. 1B), so the use of a correction factor for the Isocode paper might be feasible.

We tried to correct RNA values from filter paper for the hematocrit in three ways. Initially, we simply adjusted the RNA value obtained from the filter paper for each patient’s hematocrit, producing the data shown in Fig. 1. Then we tried the use of two constant correction factors. The first, 1.82, was derived from the mean hematocrit of 45 for this cohort. The second, 2.0, was based on the assumption that the mean hematocrit was close to 50. Use of a constant factor could be appropriate when hematocrits are not available. The two constants produced very similar results (data not shown).

The sensitivities of filter dot RNA and plasma RNA were compared by classifying each RNA determination as above or below the limit of detection (1,000 HIV RNA copies/ml). For

![Figure 1A](image1.png)

**FIG. 1.** (A) Log_{10} hematocrit-corrected HIV RNA concentrations for 55 patients tested with no. 903 filter paper plotted against the log_{10} RNA concentration for the matching plasma sample. Samples for which either estimate was below the detection limits were excluded (n = 21). The diagonal line indicates where points would fall if the estimates from the dots and the plasma were the same (correlation coefficient = 0.88). The points in the plot are roughly scattered around this line, indicating that there is little evidence of a systematic difference between the two approaches. For 91% of the hematocrit-corrected dot values, the values were within threefold (0.5 log_{10}) of the value for plasma. (B) Mean log_{10} hematocrit-corrected HIV RNA concentrations for the Isocode filter paper plotted against the mean log_{10} RNA concentration for the matching plasma sample (correlation coefficient = 0.90). The points are shifted to the right, indicating that higher values were obtained in the assay with plasma. The mean difference was 0.22 (\( P = 0.0016 \)), meaning that, on average, the estimates from the plasma were 1.66 times the estimates from the Isocode filter paper. For 95% of the hematocrit-corrected dot values, the values were within threefold (0.5 log_{10}) of the value for plasma.
The preliminary results reported here are extremely encouraging. Using the Isocode filter paper and correcting for hematocrit and even without correcting for the use of the Isocode filter paper, we were able to quantify the amount of HIV RNA in plasma fairly accurately (Fig. 1B). No difference in sensitivity was observed between the assays with Isocode filter paper and the plasma. The reproducibility of the results of assays with Isocode filter paper were comparable to the reproducibility of the results of the standard assay with plasma. However, problems with stability exist. We are not sure whether the RNA is being degraded with time or whether we are simply having increased difficulty in removing the RNA from the filter paper with prolonged storage.

We believe that modifications and refinements of this method can provide an accurate measurement of the plasma HIV RNA level suitable for primitive field conditions. In addition, the nucleic acid recovered from the spots should be amenable to amplification and sequencing and should thus be of value for drug sensitivity and HIV clade testing (1).

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REFERENCES


