

Original Article

The impact of storage time on HCV RNA quantitative detection in serum samples by TaqMan real-time PCR

Xianjun Lao, Dongmei Yang, Minyan Liu, Wenjun Tang, Xue Qin, Shan Li

Department of Clinical Laboratory, First Affiliated Hospital of Guangxi Medical University, Nanning, Guangxi, China

Received October 28, 2015; Accepted December 25, 2015; Epub February 1, 2016; Published February 15, 2016

Abstract: Information on the concentration of hepatitis C virus (HCV) RNA is useful for patient management, including assessing treatment initiation and treatment response, and monitoring follow-up. Previous studies have reported that poor sample processing and storage conditions might influence the stability of HCV RNA, and hence its detectability. We analyzed 30 patients known to be positive with high/intermediate/low HCV RNA load, and kept their serum samples in storage at 4 °C for 0, 1, 2, 3, 7, 14, 21 and 28 days. Quantitative detection of HCV RNA was performed by TaqMan real-time PCR. The results showed that no decrease in the concentration of HCV RNA was detected over the 14 days period at 4 °C ($P > 0.05$) for the high and intermediate HCV RNA load groups. However, after 21 and 28 days of storage, the mean viral loads in both groups differed significantly from those at Day 0 ($P > 0.005$). For the low HCV RNA load group, an important loss of HCV RNA concentration was observed when the serums were stored at 4 °C after 14 days, 21 days and 28 days ($P < 0.001$). These data demonstrated that serum samples with HCV RNA levels up to 10^4 IU/ml can be stored at 4 °C for 14 days, and that lower than 10^4 IU/ml can be performed for up to only 7 days when the samples are stored at 4 °C.

Keywords: HCV RNA stability, serum sample storage, TaqMan real-time PCR

Introduction

Hepatitis C virus (HCV) infection is a serious public health issue in both developing and developed countries. According to available estimates, approximately 130-170 million people are currently infected with HCV and 3-4 million people are newly infected each year throughout the world [1, 2]. In China, HCV infection is the fourth most common infectious disease after hepatitis B, and it has become an emerging burden for society and health care systems [3]. In a clinical laboratory, the routine methods for diagnosis of HCV infection are the detection of circulating HCV antibodies (anti-HCV) and polymerase chain reaction (PCR) of HCV RNA [4]. Since quantitative detection of HCV RNA with TaqMan real-time reverse transcription polymerase chain reaction (TaqMan real-time PCR) is considered to be more precise and sensitive than traditional anti-HCV testing, it has become the most valuable tool for diagnosing HCV infection. Furthermore, information on the concentration of HCV RNA has been found to be very important in guiding early ther-

apeutic intervention for patients during acute HCV infection and in monitoring patient follow-up [5].

In clinical practice, a serum or plasma specimen may be stored for a period of time before an analysis can be conducted. While RNA is labile and RNAases are ubiquitous, one major point relative to the success of analytic methods for HCV RNA detection is how the samples are stored after they have been collected. Suboptimal specimen processing and storage conditions may result in a significant decline in the concentration of HCV RNA or even in false-negative results, especially in low titer samples. To ensure the accuracy and reproducibility of the HCV viral load test results, it is necessary to define the optimal storage conditions of the clinical specimens when they are transported to the laboratory.

Many studies have examined the impact of storage time on HCV RNA stability in samples from different origins (plasma, blood, and serum) in the presence of different preservatives (citrate,

HCV storage condition in serum

Table 1. Mean HCV RNA viral loads of three categories in different time points

Groups	HCV RNA viral load (log ₁₀ IU/mL)							
	Day 0	Day 1	Day 2	Day 3	Day 7	Day 14	Day 21	Day 28
Group 1 (n = 10)	6.5067	6.4049	6.5106	6.5425	6.4206	6.3491	5.8065	5.5330
Group 2 (n = 10)	5.3542	5.0581	5.1885	5.1653	5.2175	4.9603	4.4826	4.6270
Group 3 (n = 10)	3.5207	3.2223	3.2391	3.2698	3.1140	2.9649	2.9798	2.8928

Group 1, high HCV-RNA viral loads; Group 2, intermediate HCV-RNA viral loads; Group 3, low HCV-RNA viral loads.

EDTA) at different storage temperatures (-80°C, -70°C, -20°C, 4°C, and room temperature) [6-15]. The results of these studies suggested that different storage conditions, such as short-term storage (≤ 5 days) at 4°C or long-term storage (up to 1 year) at -20°C or lower, had no effects on HCV viral load quantitation in serum or plasma samples. The effects that storage conditions of serum samples have been studied for the cDNA polymerase chain reaction (cDNA-PCR), HCV PCR, competitive reverse transcription PCR (cRT-PCR), and branched DNA signal amplification assay; However, the effects of those storage conditions on HCV RNA load in serum samples using TaqMan real-time PCR have not yet been studied [11-13, 15].

The study reported here was designed to assess the influence on HCV RNA quantitation by TaqMan real-time PCR in serum stored at 4°C for 0, 1, 2, 3, 7, 14, 21 and 28 days.

Materials and methods

Study population

Serum samples were obtained from 30 HCV RNA-positive patients at the Department of Clinical Laboratory, First Affiliated Hospital of Guangxi Medical University, Nanning, Guangxi China. The 30 patients were divided into three groups based on the HCV RNA loads: (i) 10 samples with high HCV RNA loads (10^6 - 10^7 IU/ml); (ii) 10 samples with intermediate HCV RNA loads (10^4 - 10^6 IU/ml), and (iii) 10 samples with low HCV RNA loads (10^3 - 10^4 IU/ml). All of the participants involved in the study signed a consent form and the study were approved by the Ethics Committee of the First Affiliated Hospital of Guangxi Medical University.

Serum sample processing

A whole blood sample was collected from each patient in 5.0 ml vacuum tubes (Shanghai Kehua Bio-Engineering, LLC, Shanghai, China) without using any anticoagulants. The serum

from the tubes was separated following a single centrifugation step (10 min at 3000 rpm), and then aliquoted in eight Eppendorf tubes. One of the sample tubes was immediately used for HCV RNA titer determination at Day 0. The seven remaining aliquots were stored at 4°C for 1, 2, 3, 7, 14, 21, and 28 days to evaluate the effect of storage time. All of the HCV RNA loads were investigated at the end of the aforementioned periods of time.

TaqMan real-time PCR assay

Nucleic acid extraction and TaqMan RT-PCR were performed using the Hepatitis C virus RNA Quantification Kit (Shanghai Kehua Bio-Engineering, LLC, Shanghai, China), according to the manufacturer's instructions. This method had a detection limit of 250 IU/mL and linearity to 10^3 - 10^7 IU/mL. Intra-assay variability was defined by testing 10 replicates of each quantitation standard (10^4 IU/mL and 10^6 IU/mL). The intra-assay coefficient of variability (CV) was $\leq 5\%$ for both 10^4 IU/mL and 10^6 IU/mL. The HCV RNA amplification and quantitative determination were carried out in a SLAN-96P real-time PCR system (Shanghai Hongshi Medical Technology Company, LTD, Shanghai, China). Cycling conditions were performed as follows: 25 min at 50°C, 2 min at 94°C, 10 sec at 94°C, 5 cycles of 10 sec at 94°C, 15 sec at 55°C, and then 15 sec at 72°C followed by 5 cycles of 10 sec at 94°C and 45 sec at 60°C. Each run contained a negative control, a positive and a blank control for the absence of PCR inhibition and to avoid false-positive results. Moreover, internal standard RNA was added to each test specimen. The internal standard RNA was constructed to have the same primer-binding site as the target viral RNA, which can serve as a non-competitive positive template [16].

Statistical analysis

The data were log₁₀ transformed prior for analysis and a pairwise comparison between each

HCV storage condition in serum

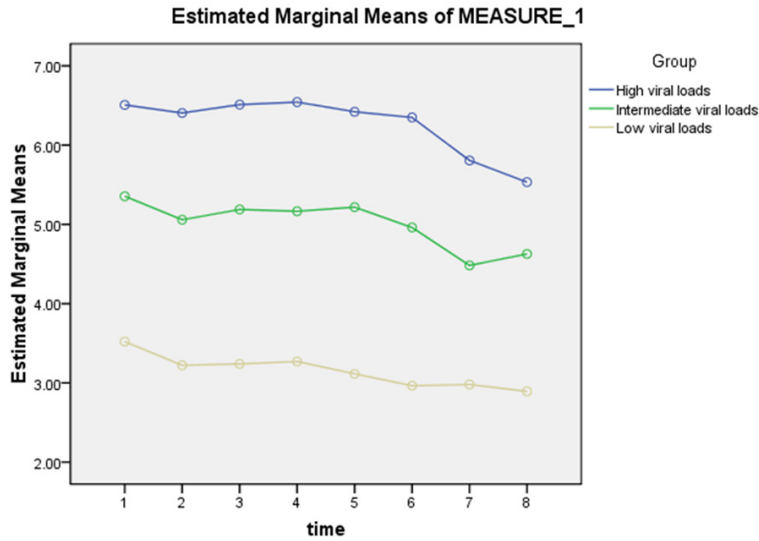


Figure 1. HCV RNA viral loads measurement in serum samples stored at 4 °C after 0, 1, 2, 3, 7, 14, 21, and 28 days. High: high HCV-RNA loads group; Intermediate: intermediate HCV-RNA loads group; Low: low HCV-RNA loads group.

time point in the same group was conducted using repeated measures analysis. Test results with *P* values less than 0.05 were considered to be statistically significant. The statistical analysis was performed with SPSS 16.0 software (SPSS Chinese, China).

Results

Thirty HCV RNA positive patients (16 males and 14 females; mean age, 45 years; range, 26-70 years) were enrolled in this study. The mean HCV RNA viral loads of all three sample categories at 0, 1, 2, 3, 7, 14, 21 and 28 days are shown in **Table 1** and **Figure 1**.

For the high HCV RNA group, the quantity of HCV RNA ranged from 6.18 log₁₀ IU/ml to 6.94 log₁₀ IU/ml, with a mean (\pm SD) of 6.50 \pm 0.29 log₁₀ IU/ml at Day 0. After 1, 2, 3, 7, and 14 days of storage at 4°C, the mean viral loads in the serum samples did not differ significantly from those taken at Day 0 (*P* > 0.05). However, when the serum samples stored at 4°C were checked after 21 days and 28 days of storage, the HCV RNA concentration was found to be significantly lower (0.70 log₁₀, *P* = 0.011 and 0.97 log₁₀, *P* = 0.005, respectively).

For the intermediate HCV RNA group, the quantity of HCV RNA ranged from 4.48 log₁₀ IU/ml to 5.95 log₁₀ IU/ml, with a mean (\pm SD) of 5.35

\pm 0.49 log₁₀ IU/ml at Day 0. Similarly, the mean viral loads in serum samples over the 14-day period were found to be stable (0.39 log₁₀, *P* = 0.116). However, the serum HCV RNA titer decreased significantly at 4°C after 21 days and 28 days of storage (0.87 log₁₀, *P* = 0.003 and 0.73 log₁₀, *P* = 0.005, respectively).

For the low HCV RNA group, the quantity of HCV RNA ranged from 3.22 log₁₀ IU/ml to 3.97 log₁₀ IU/ml, with a mean (\pm SD) of 3.52 \pm 0.25 log₁₀ IU/ml at Day 0. Notably, one serum sample on Day 3 and another serum sample on Day 7 became undetectable (data not shown). A statistical-

ly significant decline in the mean viral loads was found on Day 14, Day 21, and Day 28 (0.56 log₁₀, *P* < 0.001, 0.54 log₁₀, *P* < 0.001 and 0.63 log₁₀, *P* < 0.001, respectively).

Discussion

PCR detection of HCV RNA is a major technique that is used to screen and diagnose HCV infection. Quantitation of HCV RNA has been described as a key predictor for patient management, including assessing treatment initiation and treatment response, and monitoring follow-up. Previous studies have reported that different factors might have an effect on nucleic acid stability, such as shipping conditions, screening in blood banks, handling of samples, etc. To ensure optimal prognostic and therapeutic value in monitoring chronically infected patients, it is critical to evaluate the effects of handling clinical specimens in routine clinical laboratory practice.

In this study, 30 samples were included, covering an approximately 3 log₁₀ HCV RNA dynamic range, and these were divided based on HCV RNA loads, as follows: a high HCV RNA load group, an intermediate HCV RNA load group, and a low HCV RNA load group. The storage condition for all of the samples was 4°C for 0, 1, 2, 3, 7, 14, 21 and 28 days. The results showed high stability of HCV RNA over the

HCV storage condition in serum

14-day period at 4°C in the high- and intermediate HCV RNA load groups and no significant decrease in the HBV DNA titer was detected. However, for the samples stored for 21 and 28 days, the mean viral loads differed significantly from those at Day 0 for both of these days. For the low HCV RNA load group, an apparent decrease in the HCV RNA level was observed on Day 14. In our study, a 0.50 log₁₀ or greater loss of HCV RNA was defined as the probability of specimen failure. Because it is generally accepted that a difference of less than 0.5 log₁₀ IU/ml in viral loads detected using molecular assays should be considered the result of assay variance [17-19]. Furthermore, with the exception of two samples with low viral loads, all the samples were found to be positive for HCV RNA at 28 days at 4°C storage. This is possibly due to the fact that the particulate nature of the virus genome or the host's properties may affect the continuation of HCV RNA positivity [20]. Nevertheless, further studies are needed to more clearly address this issue.

Since Cuypers et al. [11] first examined the stability of HCV RNA in serum stored at 4°C using the cDNA-PCR method in 1992, this subject has been studied extensively over the past several years. In one study, seven serum samples taken from patients with histologically-proven chronic hepatitis C was stored at 4°C until HCV RNA determination. The results showed that the HCV RNA levels were not significantly different from the levels at time point 0 after five days at 4°C, but the levels were lost completely after 6 months at 4°C [12]. In another study, the impact of storage time on HCV RNA detection was analyzed in 11 serum samples subjected to 4°C storage for up to one month by using a Quantiplex branched-DNA assay. The authors reported that the estimated probability of specimen failure was 18%, which was lowest at 4°C [10]. Comparable results have also been presented in our studies. The probability of specimen failure was 14%, 14%, and 11% at Day 28 for the high, intermediate, and low HCV-RNA load groups, respectively. Moreover, Gessoni et al. [7] suggested that the standard procedures for storage schedules was 4°C for a maximum of 48 h without compromising the quality of the HCV RNA in whole blood samples. After examining the results of these studies, some differences were found in how the laboratories handled the specimens and in the standardizations they used for the nucleic acid test.

Therefore, evaluation of the optimal storage conditions for HCV RNA preservation should be conducted by individual laboratories based on their own technology.

Krajden et al. [10] the following elements are required to ensure an accurate assessment of the effect that storage at 4°C has on HCV RNA quantitative detection for clinical specimens: (1) sufficient samples to ensure statistical power; (2) a sensitive quantitative assay for viral load measurements; (3) an appropriate definition of a clinically relevant descriptive endpoint; and (4) reliable statistical analysis methods. In the present work, all of these criteria were applied. For clinical specimens, we included 30 HCV RNA-positive patients, and then these samples were equally divided into three groups in accordance with the HCV RNA loads. For the quantitative assay, a more direct and sensitive method, TaqMan real-time PCR, was used. This method had a detection limit of 250 IU/mL and linearity to 10³-10⁷ IU/mL. The intra-assay CV was ≤ 5% for both 10⁴ IU/mL and 10⁶ IU/mL. Moreover, in this assay, an internal standard RNA was added to each test specimen in order to avoid any false-negative results. For the descriptive endpoint, we defined the probability of specimen failure as a 0.50 log₁₀ or greater loss of HCV RNA. For statistical analysis, the repeated measures analysis method was performed with SPSS 16.0 software (SPSS Chinese, China). Test results with *P* values less than 0.05 were considered to be statistically significant.

In conclusion, our data demonstrated that serum samples with HCV RNA levels up to 10⁴ IU/ml can only be stored at 4°C for 14 days, and those lower than 10⁴ IU/ml can be performed for up to only 7 days when the samples are stored at 4°C.

Acknowledgements

We would like to thank Professor Zhiping Chen for the data analysis and Scribendi.com for its linguistic assistance during the preparation of this manuscript.

Disclosure of conflict of interest

None.

Address correspondence to: Dr. Shan Li, Department of Clinical Laboratory, First Affiliated Hospital

HCV storage condition in serum

of Guangxi Medical University, Nanning 530021, Guangxi, China. Tel: +86-771-5356052; Fax: +86-0771-865353342; E-mail: lis8858@126.com

References

- [1] Global Burden Of Hepatitis C Working Group. Global burden of disease (GBD) for hepatitis C. *J Clin Pharmacol* 2004; 44: 20-29.
- [2] Mohd Hanafiah K, Groeger J, Flaxman AD and Wiersma ST. Global epidemiology of hepatitis C virus infection: new estimates of age-specific antibody to HCV seroprevalence. *Hepatology* 2013; 57: 1333-1342.
- [3] Qin Q, Smith MK, Wang L, Su Y, Wang L, Guo W, Wang L, Cui Y and Wang N. Hepatitis C virus infection in China: an emerging public health issue. *J Viral Hepat* 2015; 22: 238-244.
- [4] Abe K and Konomi N. Hepatitis C virus RNA in dried serum spotted onto filter paper is stable at room temperature. *J Clin Microbiol* 1998; 36: 3070-3072.
- [5] Comert F, Aktas E, Terzi HA, Kulah C, Ustundag Y, Kokturk F and Aydemir S. Evaluation of hepatitis C virus RNA stability in room temperature and multiple freeze-thaw cycles by COBAS AmpliPrep/COBAS TaqMan HCV. *Diagn Microbiol Infect Dis* 2013; 75: 81-85.
- [6] Jose M, Curtu S, Gajardo R and Jorquera JI. The effect of storage at different temperatures on the stability of Hepatitis C virus RNA in plasma samples. *Biologicals* 2003; 31: 1-8.
- [7] Gessoni G, Barin P, Frigato A, Fezzi M, de Fusco G, Arreghini N, Galli P and Marchiori G. The stability of hepatitis C virus RNA after storage at +4 degrees C. *J Viral Hepat* 2000; 7: 283-286.
- [8] Damen M, Sillekens P, Sjerps M, Melsert R, Frantzen I, Reesink HW, Lelie PN and Cuypers HT. Stability of hepatitis C virus RNA during specimen handling and storage prior to NASBA amplification. *J Virol Methods* 1998; 72: 175-184.
- [9] de Moreau de Gerbehaye AI, Bodeus M, Robert A, Horsmans Y and Goubau P. Stable hepatitis C virus RNA detection by RT-PCR during four days storage. *BMC Infect Dis* 2002; 2: 22.
- [10] Krajden M, Minor JM, Zhao J, Rifkin O and Comanor L. Assessment of hepatitis C virus RNA stability in serum by the Quantiplex branched DNA assay. *J Clin Virol* 1999; 14: 137-143.
- [11] Cuypers HT, Bresters D, Winkel IN, Reesink HW, Weiner AJ, Houghton M, van der Poel CL and Lelie PN. Storage conditions of blood samples and primer selection affect the yield of cDNA polymerase chain reaction products of hepatitis C virus. *J Clin Microbiol* 1992; 30: 3220-3224.
- [12] Halfon P, Khiri H, Gerolami V, Bourliere M, Feryn JM, Reynier P, Gauthier A and Cartouzou G. Impact of various handling and storage conditions on quantitative detection of hepatitis C virus RNA. *J Hepatol* 1996; 25: 307-311.
- [13] Quan CM, Krajden M, Zhao J and Chan AW. High-performance liquid chromatography to assess the effect of serum storage conditions on the detection of hepatitis C virus by the polymerase chain reaction. *J Virol Methods* 1993; 43: 299-307.
- [14] Cardoso MS, Koerner K, Hinz W, Lenz C, Schwandt A and Kubanek B. Hepatitis C virus stability: the issue! *Vox. Sang* 1999; 76: 124-127.
- [15] Manzin A, Bagnarelli P, Menzo S, Giostra F, Brugia M, Francesconi R, Bianchi FB and Clementi M. Quantitation of hepatitis C virus genome molecules in plasma samples. *J Clin Microbiol* 1994; 32: 1939-1944.
- [16] Kim K, Park J, Chung Y, Cheon D, Lee IB, Lee S, Yoon J, Cho H, Song C and Lee KH. Use of internal standard RNA molecules for the RT-PCR amplification of the faeces-borne RNA viruses. *J Virol Methods* 2002; 104: 107-115.
- [17] Miskovsky EP, Carrella AV, Gutekunst K, Sun CA, Quinn TC and Thomas DL. Clinical characterization of a competitive PCR assay for quantitative testing of hepatitis C virus. *J Clin Microbiol* 1996; 34: 1975-1979.
- [18] Kessler HH, Santner B, Umlauf F, Urbaneck M, Kronawetter M, Pierer K, Stunzner D, Grunewald K and Marth E. Detection of hepatitis C viral sequences in serum by 'nested' polymerase chain reaction (PCR) and a commercial single-round PCR assay. *Clin Diagn Virol* 1995; 4: 239-250.
- [19] Farma E, Boeri E, Bettini P, Repetto CM, McDermott J, Lillo FB and Varnier OE. Single-step PCR in molecular diagnosis of hepatitis C virus infection. *J Clin Microbiol* 1996; 34: 3171-3174.
- [20] Sener K, Yapar M, Bedir O, Gul C, Coskun O and Kubar A. Stability of hepatitis C virus RNA in blood samples by TaqMan real-time PCR. *J Clin Lab Anal* 2010; 24: 134-138.