

Comparison of Real-time PCR Methods and pp65 Antigenemia Assay to Detect Cytomegalovirus Reactivation in Hematopoietic Stem Cell Transplantation

Seon-Young Lee¹, Byeong-Sun Choi¹, Sung Soon Kim¹

Su-Mi Choi, M.D.², Wan-Shik Shin, M.D.², and Joo-Shil Lee, Ph.D.¹

Center for Immunology and Pathology¹, National Institute of Health, Seoul, Korea

The Catholic Hematopoietic Stem Cell Transplantation Center², College of Medicine, The Catholic University of Korea, Seoul, Korea

Human cytomegalovirus (HCMV) is a common human pathogen that causes morbidity and mortality in hematopoietic stem cell transplantation (HSCT) recipients. Early diagnosis of HCMV infection or reactivation, and setting threshold values for effective pre-emptive therapies, are required for appropriate HCMV disease prevention in HSCT recipients. We compared the HCMV infections detected by the two methods, LightCycler-based PCR (LC PCR) and in-house immediate early protein PCR (in-house IE PCR) with the results of a pp65 antigenemia assay as the reference. The sensitivity and specificity for the in-house IE PCR were 79.3% and 72.7%, respectively, and 82.9% and 40.7%, respectively, for the LC PCR. The correlation between the HCMV viral load and pp65 antigenemia in HSCT recipients was $r=0.603$ with in-house IE PCR and $r=0.525$ with LC PCR. The discordant results between methods and relatively low (r) values suggest that we need more study to set threshold values according to the using methods with clinical outcome.

Key Words : HCMV, Real-time PCR, pp65 antigenemia, LC PCR, In-house IE PCR

Human cytomegalovirus (HCMV) is a common human pathogen that causes morbidity and mortality in hematopoietic stem cell transplantation (HSCT) recipients (1-3). Early diagnosis of HCMV infection or reactivation and effective pre-emptive therapies with setting cut-off values are required for appropriate HCMV disease prevention in HSCT recipients (4-6). Although the pp65 antigenemia assay is one of the most widely used methods to monitor CMV infection or reactivation, this assay has many limitations: it requires high-level skill in the experimenter; it is very labor intensive and the

specimens must be handled quickly (1, 3, 6-8). Recent reports suggest that HCMV infections with real-time PCR methods can be predicted more rapidly and sensitively than with the HCMV pp65 antigenemia assay (1, 7). Therefore, we compared the pp65 antigenemia assay and two kinds of quantitative HCMV real-time PCR methods in 67 patients who had undergone HSCT to apply the prediction of HCMV infections and reactivation.

Sixty-seven HSCT patients who had visited at a Korea Catholic Hematopoietic Stem Cell Transplantation Center between May 2000 and March 2002 were participated for this study. These patients were included in the study and gave their informed consent. There were 43 men and 24 women with the median age of 32 years (15-51 years). The median follow-up duration after organ transplant was 690 days (32-1,226 days) and the number of samples obtained from the 67 HSCT patients was 254.

Submitted 29 February 2008, Accepted 10 June 2008

This work was supported by the Korean National Institute of Health grant 2006-N00240-00. Seon-Young Lee and Byeong-Sun Choi contributed equally to this project.

Correspondence: Joo Shil Lee, Ph.D.

Director of Center for Immunology and Pathology, National Institute of Health, 194 Tongillo, Eunpyung-gu, Seoul 122-701, Korea

Tel : +82-2-380-2150, Fax : +82-2-382-6542

E-mail : jooshil@nih.go.kr

HCMV DNA was extracted from the plasma (0.2 mL) of HSCT recipients with a QIAamp Blood Mini Kit (Qiagen, Valencia, CA, USA) according to the manufacturer's instructions and the detection of HCMV DNA was determined by two distinct real-time PCR methods: Light Cycler-based PCR (LC PCR) and in-house immediate early protein-based PCR (in-house IE PCR). HCMV LC PCR was carried out using the commercial artus[®] CMV LC PCR kit (Qiagen, Hilden, Germany) and in-house IE PCR method was performed with the primers and probe reported by Ikewaki et al (7). The results for pp65 antigenemia assay were presented as the number of pp65-positive cells per 2×10^5 leukocytes (8).

HCMV infection from 254 plasma samples was detected in 82 (32.3%) by pp65 antigenemia assay, in 112 (44.1%) by in-house IE PCR, and in 170 (66.9%) by LC PCR.

Table 1. Results of HCMV Detection by pp65 Antigenemia Assay and Two Distinct Real-time PCRs in 254 Plasma Samples

Method and result type	No. of samples evaluated by pp65 antigenemia assay*		
	Positive	Negative	Total
Quantitative LC PCR			
Positive	68	102	170
Negative	14	70	84
Total	82	172	254
Quantitative in-house IE PCR			
Positive	65	47	112
Negative	17	125	142
Total	82	172	254

*Positive results of pp65 antigenemia assay were presented as number of pp65-positive cells per 2×10^5 leukocytes.

PCR. The sensitivity and specificity of the two distinct real-time PCR methods based on the results of pp65 antigenemia assay were 82.9% and 40.7% in the LC PCR and 79.3% and 72.9% in the in-house IE PCR, respectively (Table 1). The correlations between the HCMV viral load and pp65 antigenemia in HSCT patients were $r = 0.603$ with in-house IE PCR and $r = 0.525$ with LC PCR, respectively ($P < 0.0001$, Fig. 1). The above results showed that in-house IE PCR was higher specificity than the LC PCR.

To set the optimal cut-off values of the HCMV DNA load by the two distinct real-time PCR methods for detection of HCMV infection and reactivation, the receiver

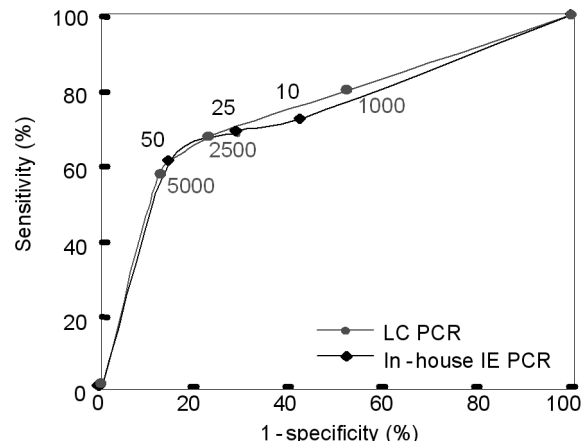


Fig. 2. Receiver-operating characteristic (ROC) curves constructed from the HCMV DNA loads measured by LC PCR and in-house IE PCR to evaluate the sensitivity and specificity of the other real-time PCR method at each cut-off values using eighty-two pp65 antigenemia positive samples. The numbers in the Fig. indicate cut-off values for detection of HCMV infection were set as 10, 25, and 50 copies/mL in the in-house IE PCR and as 1000, 2500, and 5000 copies/mL in LC PCR.

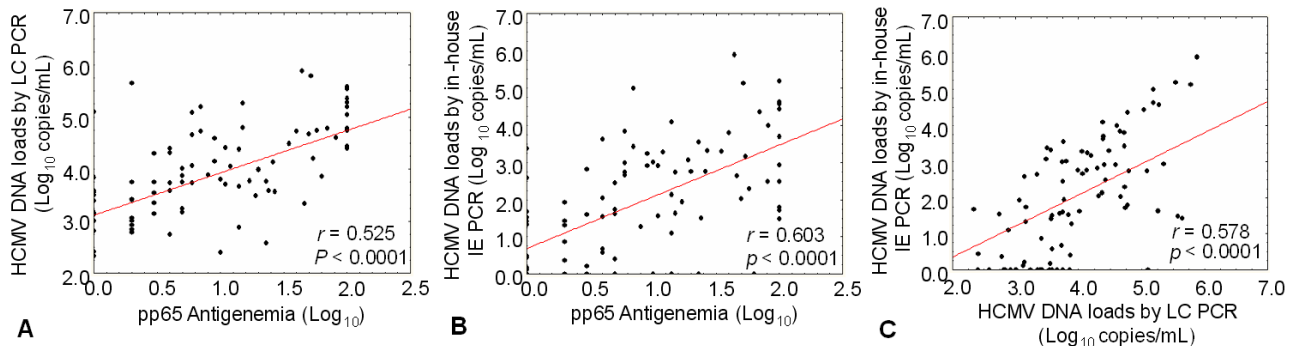


Fig. 1. The correlation between the HCMV DNA loads measured by real-time PCR methods and the numbers of pp65-positive cells obtained by pp65 antigenemia assay was analyzed using 254 plasma samples collected from HSCT patients. (A) Correlation between the HCMV DNA load measured by LC PCR and the pp65-positive cell numbers obtained by pp65 antigenemia assay; (B) Correlation between the HCMV DNA load measured by in-house IE PCR and the pp65-positive cell numbers obtained by pp65 antigenemia assay; (C) Correlation between the HCMV DNA loads measured by LC PCR and by in-house IE PCR.

operating characteristic (ROC) curves were constructed using the eighty-two pp65 antigenemia positive samples (Fig. 2). When cut-off values of HCMV DNA load in the in house IE PCR were set as 10, 25, and 50 copies/ mL, the sensitivity and specificity for detection of HCMV infection were 72.1% and 57.1% at 10 copies/mL; 69.1% and 71.4% at 25 copies/mL; 60.3% and 85.7% at 50 copies/mL, respectively. When cut-off values of HCMV DNA load in LC PCR were set as 1,000, 2,500, and 5,000 copies/mL, the sensitivity and specificity for detection of HCMV infection were 80.0% and 47.1% at 1,000 copies/mL; 67.7% and 76.5% at 2,500 copies/mL; 56.9% and 88.2% 5,000 copies/mL, respectively. From these data, the optimal cut-off values, 25 copies/mL for in-house IE PCR and 2,500 copies/mL for LC PCR, were applied to detect HCMV infection and reactivation for eighteen naïve HSCT patients and these cut-off values were useful (data not shown). Some studies suggested that the cut-off values of HCMV viral loads for preemptive therapy might differ such as 200 copies/ml and $\geq 10,000$ copies/mL depending on the kind of real-time PCR method and clinical samples (7, 8, 10, 11).

In conclusion, these results suggest that the optimal cut-off values of HCMV viral load for pre-emptive therapy in HSCT recipients should be differently considered by each individual method including detection protocols and specimen type as DNA sources (plasma or whole blood).

References

- 1) Alice T, Enrietto M, Pittaluga F, Varetto S, Franchello A, Marchiaro G, Ghisetti V: *Quantitation of cytomegalovirus DNA by real-time polymerase chain reaction in peripheral blood specimens of patients with solid organ transplants: comparison with end-point PCR and pp65 antigen test. J Med Virol* 78: 915-22, 2006
- 2) Boeckh M, Huang M, Ferrenberg J, Stevens-Ayers T, Stensland L, Nichols WG, Corey L: *Optimization of Quantitative Detection of Cytomegalovirus DNA in Plasma by Real-Time PCR. J Clin Microbiol* 42: 1142-8, 2004
- 3) Pumannova M, Roubalova K, Vitek A, Sajdova J: *Comparison of quantitative competitive polymerase chain reaction-enzyme-linked immunosorbent assay with LightCycler-based polymerase chain reaction for measuring cytomegalovirus DNA in patients after hematopoietic stem cell transplantation. Diagn Microbiol Infect Dis* 54:115-20, 2006
- 4) Boeckh M, Gallez-Hawkins GM, Myerson D, Zaia JA, Bowden RA: *Plasma polymerase chain reaction for cytomegalovirus DNA after allogeneic marrow transplantation: comparison with polymerase chain reaction using peripheral blood leukocytes, pp65 antigenemia, and viral culture. Transplantation* 64:108-13, 1997
- 5) Gor D, Sabin C, Prentice HG, Vyas N, Man S, Griffiths PD, Emery VC: *Longitudinal fluctuations in cytomegalovirus load in bone marrow transplant patients: relationship between peak virus load, donor/recipient serostatus, acute GVHD and CMV disease. Bone Marrow Transplant* 21:597-605, 1998
- 6) Ikewaki J, Ohtsuka E, Kawano R, Ogata M, Kikuchi H, Nasu M: *Real-time PCR assay compared to nested PCR and antigenemia assays for detecting cytomegalovirus reactivation in adult T-cell leukemia-lymphoma patients. J Clin Microbiol* 41:4382-7, 2003
- 7) Ikewaki J, Ohtsuka E, Satou T, Kawano R, Ogata M, Kikuchi H, Nasu M: *Real-time PCR assays based on distinct genomic regions for cytomegalovirus reactivation following hematopoietic stem cell transplantation. Bone Marrow Transplant* 35:403-10, 2005
- 8) Mori T, Okamoto S, Watanabe R, Yajima T, Iwao Y, Yamazaki R, Nakazato T, Sato N, Iguchi T, Nagayama H, Takayama N, Hibi T, Ikeda Y: *Dose-adjusted preemptive therapy for cytomegalovirus disease based on real-time polymerase chain reaction after allogeneic hematopoietic stem cell transplantation. Bone Marrow Transplant* 29:777-82, 2002
- 9) Choi SM, Lee DG, Choi JH, Yoo JH, Kim YJ, Park SH, Park SN, Min CK, Lee S, Kim HJ, Kim DW, Lee JW, Min WS, Shin WS, Kim CC: *Risk-adapted preemptive therapy for cytomegalovirus disease after allogeneic stem cell transplantation: a single-center experience in Korea. Int J Hematol* 81:69-74, 2005
- 10) Yakushiji K, Gondo H, Kamezaki K, Shigematsu K, Hayashi S, Kuroiwa M, Taniguchi S, Ohno Y, Takase K, Numata A, Aoki K, Kato K, Nagafuji K, Shimoda K, Okamura T, Kinukawa N, Kasuga N, Sata M, Harada M: *Monitoring of cytomegalovirus reactivation after allogeneic stem cell transplantation: comparison of an antigenemia assay and quantitative real-time polymerase chain reaction. Bone Marrow Transplant* 29:599-606, 2002
- 11) Li H, Dummer JS, Estes WR, Meng S, Wright PF, Tang YW: *Measurement of human cytomegalovirus loads by quantitative real-time PCR for monitoring clinical intervention in transplant recipients. J Clin Microbiol* 41:187-91, 2003