recipients with high EBV loads showing low activity of specific CD8<sup>+</sup> T were at high risk for PTLD, using an enzyme-linked immunospot assay in pediatric solid-organ transplant recipients. Such a functional impairment may be present and influence the chronic high EBV load state.

Analyzing the expression profile of EBV-related genes is useful in clarifying the pathogenesis of EBV-associated diseases [16]. In each chronic high EBV load carrier, EBER1 and BARTs were abundantly detected, and LMP2 was found in half of the carriers. The mean expression level of LMP2 was low, compared with those of EBER1 and BARTs. No transcript specific for latency type 3 was detected in chronic high EBV load carriers. This restricted pattern is latency type 0, which is found in EBVinfected memory B cells [8, 29, 30], suggesting that the pattern of infected B cells in those with chronic high EBV loads may be the same as that in healthy EBV carriers. Qu et al [31] reported that peripheral blood lymphocytes in those with chronic high EBV loads after solid-organ transplant were divided into 2 types of infected cells: lymphocytes expressing LMP2 and carrying a low number of copies of EBV, and others coexpressing LMP1 and LMP2 and having a high number of copies of EBV. In the present study, expression of LMP1 was not observed in all chronic high EBV load carriers. Quantification in the blood sample may not have been sufficiently sensitive to detect a small population of infected cells expressing the LMP1 gene. Additionally, the expression pattern may vary with the time at which the blood sample is obtained. Transcripts specific for latency type 3 were detected in some samples from chronic high EBV load carriers obtained within 4 months after EBV infection (data not shown), and the pattern of EBV gene expression subsequently changed to latency type 0. Dynamic changes in the immune response, such as an increased precursor frequency of EBV-specific CD8+ T cells after reduced immune suppression, may influence expression patterns in chronic high EBV load carriers; however, this change was not evaluated in the present study.

After primary infection, EBV persists as a latent infection in memory B cells. Then, memory B cells occasionally differentiate into plasma cells that undergo lytic infection and produce new virus. Newly infected naive B cells become transformed but are controlled by CD8+ T cells specific for EBV. This cycle is necessary to maintain a latent infection in vivo [21, 32]. Thus, we suggest a novel mechanism for the maintenance of high viral loads in the blood of chronic high EBV load carriers. First, a large number of memory B cells latently infected with EBV survive after the primary or reactivated EBV infection occurring after transplant. Second, the latently infected memory B cells readily differentiate into plasma cells to produce the virus. Immunosuppression may influence these processes. Treatment with acyclovir or ganciclovir for a relatively prolonged period may modulate the chronic high EBV load state. However, our results did not provide mechanistic or therapeutic information.

## **Acknowledgments**

We thank the following people for providing samples from patients with infectious mononucleosis or posttransplant lymphoproliferative disorder: Naoko Nishimura and Takao Ozaki (Konan Kosei Hospital); Masayuki Hasegawa and Kazue Ito (Nagoya Ekisaikai Hospital); Masanori Yamada (Chubu Rosai Hospital); and Yasuko Abe (Shizuoka Children's Hospital). We also thank Fumiyo Ando for secretarial assistance.

## References

- Rickinson AB, Kieff E. Epstein-Barr virus. In: Knipe DM, Howley PM, eds. Fields virology. Fifth ed. Philadelphia, PA: Lippincott Williams & Wilkins, 2007:2655–2700.
- Cohen JI. Epstein-Barr virus infection. N Engl J Med 2000; 343:481– 492
- Kis LL, Takahara M, Nagy N, Klein G, Klein E. Cytokine mediated induction of the major Epstein-Barr virus (EBV)-encoded transforming protein, LMP-1. Immunol Lett 2006; 104:83–88.
- Williams H, Crawford DH. Epstein-Barr virus: the impact of scientific advances on clinical practice. Blood 2006; 107:862–869.
- 5. Wagner HJ, Wessel M, Jabs W, et al. Patients at risk for development of posttransplant lymphoproliferative disorder: plasma versus peripheral blood mononuclear cells as material for quantification of Epstein-Barr viral load by using real-time quantitative polymerase chain reaction. Transplantation 2001; 72:1012–1019.
- Lee TC, Savoldo B, Rooney CM, et al. Quantitative EBV viral loads and immunosuppression alterations can decrease PTLD incidence in pediatric liver transplant recipients. Am J Transplant 2005; 5:2222–2228.
- Kimura H, Morita M, Yabuta Y, et al. Quantitative analysis of Epstein-Barr virus load by using a real-time PCR assay. J Clin Microbiol 1999; 37:132–136.
- Hopwood PA, Brooks L, Parratt R, et al. Persistent Epstein-Barr virus infection: unrestricted latent and lytic viral gene expression in healthy immunosuppressed transplant recipients. Transplantation 2002; 74:194– 202
- Campe H, Jaeger G, Abou-Ajram C, et al. Serial detection of Epstein-Barr virus DNA in sera and peripheral blood leukocyte samples of pediatric renal allograft recipients with persistent mononucleosis-like symptoms defines patients at risk to develop post-transplant lymphoproliferative disease. Pediatr Transplant 2003;7:46–52.
- D'Antiga L, Del Rizzo M, Mengoli C, Cillo U, Guariso G, Zancan L. Sustained Epstein-Barr virus detection in paediatric liver transplantation: insights into the occurrence of late PTLD. Liver Transpl 2007;13: 343–348.
- Bingler MA, Feingold B, Miller SA, et al. Chronic high Epstein-Barr viral load state and risk for late-onset posttransplant lymphoproliferative disease/lymphoma in children. Am J Transplant 2008; 8:442–445.
- Green M, Soltys K, Rowe DT, Webber SA, Mazareigos G. Chronic high Epstein-Barr viral load carriage in pediatric liver transplant recipients. Pediatr Transplant 2009; 13:319–323.
- 13. Kamei H, Fujimoto Y, Nagai S, Suda R, Yamamoto H, Kiuchi T. Impact of non-congestive graft size in living donor liver transplantation: new indicator for additional vein reconstruction in right liver graft. Liver Transpl 2007; 13:1295–1301.
- Kimura H, Hoshino Y, Kanegane H, et al. Clinical and virologic characteristics of chronic active Epstein-Barr virus infection. Blood 2001; 98:280–286.
- Wada K, Kubota N, Ito Y, et al. Simultaneous quantification of Epstein-Barr virus, cytomegalovirus, and human herpesvirus 6 DNA in samples from transplant recipients by multiplex real-time PCR assay. J Clin Microbiol 2007; 45:1426–1432.
- Kubota N, Wada K, Ito Y, et al. One-step multiplex real-time PCR assay to analyse the latency patterns of Epstein-Barr virus infection. J Virol Methods 2008: 147:26–36.
- 17. Patel K, Whelan PJ, Prescott S, et al. The use of real-time reverse

- transcription-PCR for prostate-specific antigen mRNA to discriminate between blood samples from healthy volunteers and from patients with metastatic prostate cancer. Clin Cancer Res **2004**; 10:7511–7519.
- Muti G, Klersy C, Baldanti F, et al. Epstein-Barr virus (EBV) load and interleukin-10 in EBV-positive and EBV-negative post-transplant lymphoproliferative disorders. Br J Haematol 2003;122:927–933.
- Stevens SJ, Verschuuren EA, Pronk I, et al. Frequent monitoring of Epstein-Barr virus DNA load in unfractionated whole blood is essential for early detection of posttransplant lymphoproliferative disease in high-risk patients. Blood 2001; 97:1165–1171.
- Baldanti F, Grossi P, Furione M, et al. High levels of Epstein-Barr virus DNA in blood of solid-organ transplant recipients and their value in predicting posttransplant lymphoproliferative disorders. J Clin Microbiol 2000; 38:613–619.
- Kimura H, Ito Y, Suzuki R, Nishiyama Y. Measuring Epstein-Barr virus (EBV) load: the significance and application for each EBV-associated disease. Rev Med Virol 2008; 18:305–319.
- LaCasce AS. Post-transplant lymphoproliferative disorders. Oncologist 2006: 11:674

  –680.
- Bakker NA, van Imhoff GW, Verschuuren EA, van Son WJ. Presentation and early detection of post-transplant lymphoproliferative disorder after solid organ transplantation. Transpl Int 2007; 20:207–218.
- Hoshino Y, Kimura H, Tanaka N, et al. Prospective monitoring of the Epstein-Barr virus DNA by a real-time quantitative polymerase chain reaction after allogenic stem cell transplantation. Br J Haematol 2001; 115:105–111.
- 25. Rowe DT, Webber S, Schauer EM, Reyes J, Green M. Epstein-Barr virus

- load monitoring: its role in the prevention and management of post-transplant lymphoproliferative disease. Transpl Infect Dis 2001; 3:79–87.
- 26. Meij P, van Esser JW, Niesters HG, et al. Impaired recovery of Epstein-Barr virus (EBV)-specific CD8+ T lymphocytes after partially T-depleted allogeneic stem cell transplantation may identify patients at very high risk for progressive EBV reactivation and lymphoproliferative disease. Blood 2003; 101:4290-4297.
- Macedo C, Donnenberg A, Popescu I, et al. EBV-specific memory CD8+ T cell phenotype and function in stable solid organ transplant patients. Transpl Immunol 2005; 14:109–116.
- Smets F, Latinne D, Bazin H, et al. Ratio between Epstein-Barr viral load and anti-Epstein-Barr virus specific T-cell response as a predictive marker of posttransplant lymphoproliferative disease. Transplantation 2002; 73:1603–1610.
- Yang J, Tao Q, Flinn IW, et al. Characterization of Epstein-Barr virusinfected B cells in patients with posttransplantation lymphoproliferative disease: disappearance after rituximab therapy does not predict clinical response. Blood 2000; 96:4055–4063.
- Babcock GJ, Hochberg D, Thorley-Lawson AD. The expression pattern
  of Epstein-Barr virus latent genes in vivo is dependent upon the differentiation stage of the infected B cell. Immunity 2000; 13:497–506.
- Qu L, Green M, Webber S, Reyes J, Ellis D, Rowe D. Epstein-Barr virus gene expression in the peripheral blood of transplant recipients with persistent circulating virus loads. J Infect Dis 2000; 182:1013–1021.
- Thorley-Lawson DA, Gross A. Persistence of the Epstein-Barr virus and the origins of associated lymphomas. N Engl J Med 2004; 350: 1328–1337.

