

Fig. 3. Production of HBV in hydrodynamically-transfected mice. (A) Density of HBV. Sera from mice injected 3 days earlier with pHBV1.5 were treated with or without detergent then fractionated through sucrose density gradient. DNA was isolated from each fraction after treatment with DNase I and then the presence of the HBV DNA sequence was checked by PCR. The density (g/cm^3) of each fraction is indicated above each lane. (B) Viral titers in serum. The levels of HBV DNA were serially determined by real-time PCR.

template for HBV expression, in addition to the inoculated plasmid. To examine the presence of cccDNA in the liver, we used a PCR procedure which selectively detects cccDNA (Fig. 1). We also checked for the presence of inoculated plasmids by amplifying the ampicillin resistance gene by PCR. The authenticity of the cccDNA detection was confirmed by the detection of a specific signal from liver tissues of patients with chronic hepatitis B, but not from the serum of patients or pHBV1.5 (Fig. 5A). Viral cccDNA was clearly detected in wild-type pHBV1.5-injected livers at 3 days as well as 3 months after the injection (Fig. 5B). As expected, cccDNA was not detected in mutant pHBV1.5-injected livers. The levels of cccDNA were measured by real-time PCR ($n=5$ for each time point) and results were 2.4×10^7 and 6.0×10^5 copies per gram of liver tissue at 3 days and 2 months after the injection, respectively. Since the liver approximately contains 1.1×10^8 of hepatocytes, the average copy numbers of HBV cccDNA per core Ag-positive hepatocyte could be estimated to be 1 or 4. Ampicillin resistance gene was similarly amplified from both wild-type pHBV1.5- and mutant pHBV1.5-injected livers. The fact that HBV gene expression was terminated

within 2 months upon injection of mutant pHBV1.5 clearly indicates that the presence of residual plasmids in the livers at later time points is not sufficient for the expression of detectable levels of HBV genes; this is consistent with a previous report [20] demonstrating that transgene expression is rapidly terminated after hydrodynamic gene delivery despite the persistence of plasmid DNA in the livers. These results support the idea that viral cccDNA is critically involved in the long-term expression and carriage of HBV in this model.

3.4. Administration of IFN α gene transiently suppressed HBV DNA replication and failed to eradicate viral template

We next sought to examine the potential usefulness of this model for the assessment of anti-viral drugs. To examine the effect of IFN α in the phase of cccDNA-dependent HBV replication, we injected either pCMV-IFN α 1 or pCMV at 70 days after pHBV1.5 injection. Injection of pCMV-IFN α 1 led to substantial IFN α production at day 1 (Fig. 6A), although IFN α could not be detected in the mock-injected mice (data not shown). The levels of IFN α after pCMV-IFN α 1 injection rapidly declined at day 3 and could not be detected at day 28. Injection of pCMV-IFN α 1 significantly suppressed viral production at day 3 but did not affect HBs production (Fig. 6B and C); this is consistent with previous findings [15,21] that IFN α suppressed HBV replication at a step of reverse transcription. In spite of the substantial suppression of HBV production at day 3, the levels of viral titers of mice injected with pCMV-IFN α 1 increased to levels similar to those of pCMV-injected mice at day 14 and later. These results indicate that IFN treatment substantially suppressed viral replication, but could not eliminate the viral template from the infected host. This model should be useful for assessing anti-viral therapy aimed at eradication of the viral template.

4. Discussion

In the present study, we demonstrated that hydrodynamic injection of a plasmid encoding an overlength of HBV DNA into nude mice established long-term replication of HBV in the liver. Since hepatic damage was not observed, this model mimics the chronic carrier-like state of human HBV infections. This model reminds us of a 1988 report by Fiteelson et al., [22] in which they stated that intrahepatic injection of replication competent HBV DNA led to persistent HBs antigenemia as well as chronic liver injury in nude mice. They had no evidence of HBV replication such as production of Dane particles in the circulation. In a preliminary experiment, we intrahepatically injected pHBV1.5 into nude mice and monitored viral production in the serum. DNase I-resistant HBV DNA could not be detected in most mice tested; a small number of mice

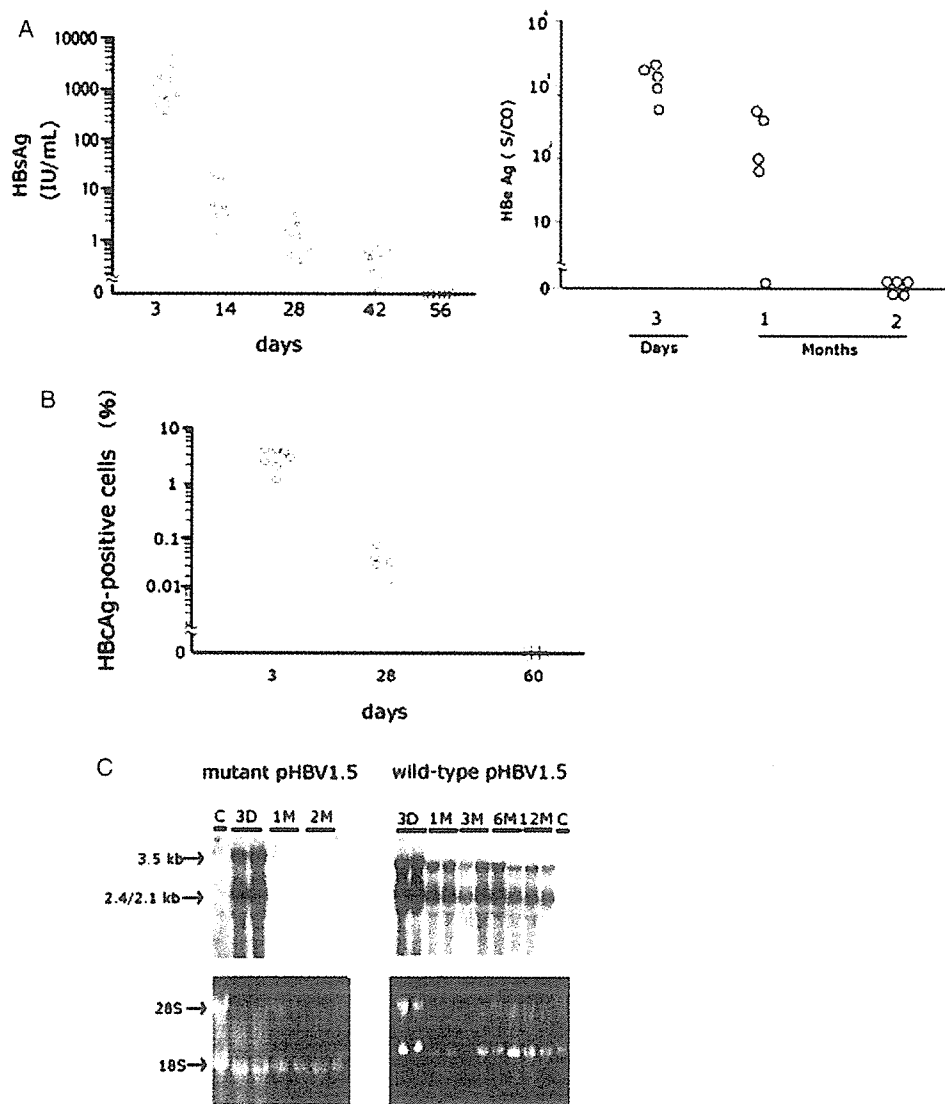


Fig. 4. Expression of HBV genes in nude mice injected with replication-incompetent pHBV1.5. (A) Serial detection of HBsAg and HBeAg in serum. (B) Frequency of HBcAg-positive hepatocytes in the livers. (C) Time-course of HBV RNA expression in the livers transfected with mutant pHBV1.5 or wild-type pHBV1.5 determined by Northern blot. C, control livers; 3D, 1M, 3M, 6M, and 12M, liver samples obtained at 3 days, 1 month, 3, 6, and 12 months after hydrodynamic injection, respectively.

produced low levels of virus at 3 days after injection but not at later time points (our unpublished data). Thus, the transfection efficiency of hydrodynamic injection of HBV DNA appeared to be higher than that of intrahepatic injection. Despite the difference in liver damage observed among these studies, we considered the absence of hepatitis in the present model reasonable, since the T-cell immune response towards HBV-related antigens could not occur. Furthermore, it should be noted that the antigenemia as well as HBV production achieved by the hydrodynamic procedure was very reproducible, which is critically important when applying this model to evaluate the efficacy of anti-viral drugs.

The duration of hydrodynamics-based gene expression varies among reports from days to months [19,23,24]. The plasmid-based gene expression of our model terminated

within 2 months, as demonstrated by the injection of replication-incompetent HBV DNA (mutant pHBV1.5). Replication-competent HBV DNA (wild-type pHBV1.5)-injected mice displayed a rapid decline of HBsAg production followed by relatively stable antigenemia for more than 1 year (Fig. 2B). Although the rapid decline observed in the first 2 weeks may reflect the plasmid-based gene expression, stable expression of HBsAg at later time points did not depend on residual plasmids in the livers, but required intracellular reproduction of HBV DNA. These results indicate that HBV replication in addition to immunological tolerance is critically important for long-term HBV expression in this system. Previous research on *in vivo* gene transfer [8] and transgenic mice [5] has indicated that HBV cccDNA, the template of HBV replication in natural infection, could not be detected in

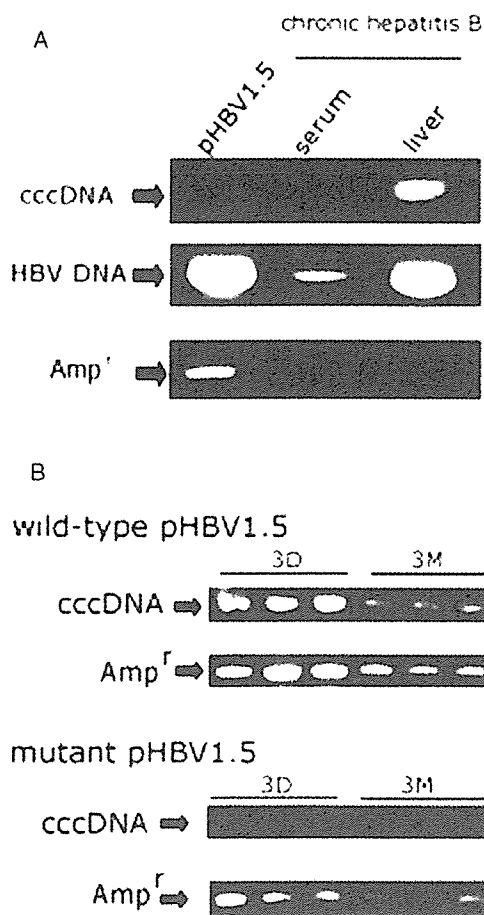


Fig. 5. Detection of HBV cccDNA and amplicin resistance gene by PCR. (A) Specificity of HBV cccDNA detection by PCR. DNA were isolated from the following samples and amplified for the detection of cccDNA, amplicin resistance gene, and HBV DNA sequence by corresponding PCRs. pHBV1.5, sample containing pHBV1.5; serum, serum from a patient with chronic hepatitis B; liver, a liver specimen from the same patient. (B) DNA was isolated from the liver samples 3 days (3D) or 3 months (3M) after hydrodynamic injection of either wild-type pHBV1.5 or mutant pHBV1.5 (*N*=3 for each group) and examined the presence of cccDNA and amplicin resistance gene by PCRs.

murine livers by Southern blot analysis. In the present study, we applied a highly sensitive PCR procedure and detected HBV cccDNA in pHBV1.5-injected livers. What is important is that the estimated numbers of HBV cccDNA per hepatocyte were 1 or 4, which should be sufficient for HBV gene expression. Taken together, the present study is the first demonstration of the production of viral cccDNA and its contribution to HBV replication in mice. Thus, the species restriction on the production of HBV cccDNA may not be as strict as has previously been believed.

Mutation of HBV DNA occurring during therapy with various nucleotide analogues leads to drug resistance and limits the success of these drugs for controlling HBV replication in humans [1,2]. Thus far, except for the in vitro recombinant HBV baculovirus system [25], there has been no useful model supporting reproduction of the HBV viral

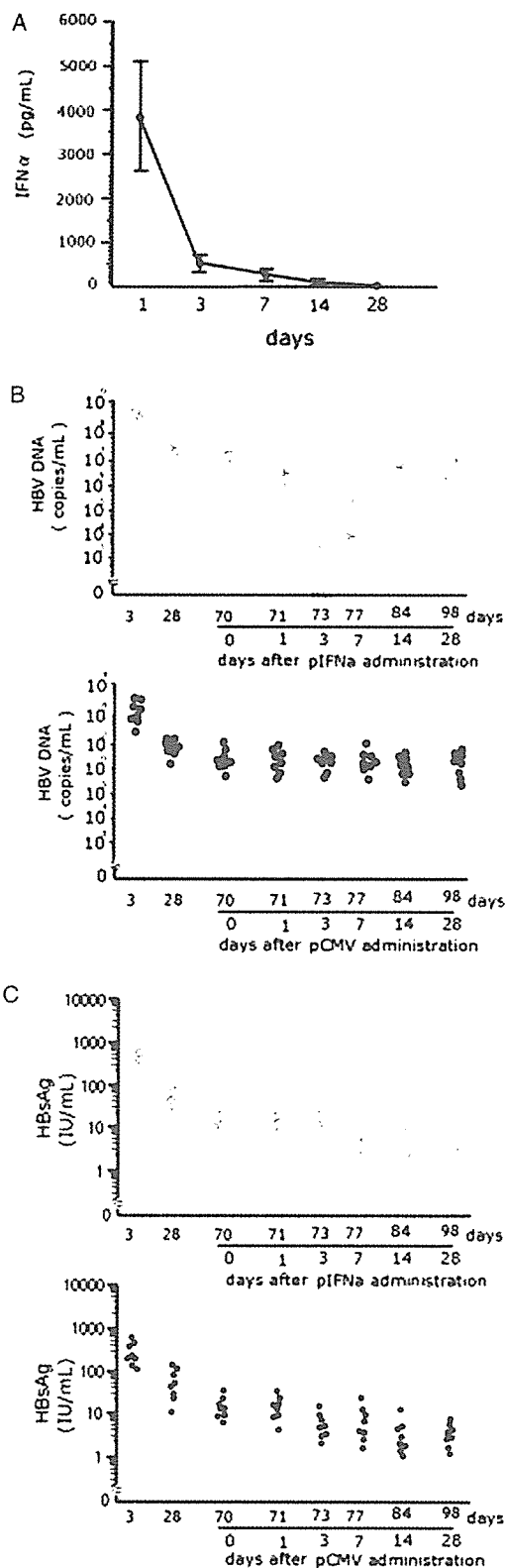


Fig. 6. Transient suppression of HBV production by IFN α gene therapy. (A) Serial determination of serum IFN α levels of nude mice after injection of pCMV-IFN α 1. Horizontal bars indicate SD. (B and C) Nude mice were transfected with pHBV1.5 and, at 70 days later, transfected with either pCMV-IFN α 1 (open circles) or pCMV (closed circles). Sera were serially obtained from the retro-orbital plexus, with HBV DNA (B) and HBsAg (C) titers being determined.

template as is the case of hepatitis C virus replicon systems [26,27]. Although HBV could not ‘infect’ murine hepatocytes, intracellular ‘reinfection’, namely recycling of HBV DNA occurs and leads to chronic viral production in the present model. Therefore, this model may provide a unique opportunity for analyzing possible mutations induced by long-term usage of various nucleotide analogs. Further study is needed to examine this possibility. Finally, intentional mutation could be easily introduced in inoculated DNA and a wide variety of mice with different genetic backgrounds can be used. The model presented here should enable analysis of viral as well as host factors that may regulate HBV replication.

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