Preliminary study on the production of transgenic mice harboring hepatitis B virus X gene *

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Abstract

AIM To establish transgenic mice lineage harboring hepatitis B virus X gene and to provide an efficient animal model for studying the exact role of the HBx gene in the process of hepatocarcinogenesis.

METHODS The HBx transgenic mice were produced by microinjecting the construct with X gene of HBV (subtype adr) DNA fragment into fertilzed eggs derived from inbred C57BL/6 strain; transgenic mice were identified by using Nested PCR; expression and phenotype of HBx gene were analyzed in liver from transgenic mic at the age of 8 weeks by RT-PCR, pathologic examination and periodic acid-schiff staining (PAS), respectively.

RESULTS Five hundred and fourteen fertilized eggs of C57 BL/6 mice were microinjected with recombinant retroviral DNA fragment, and 368 survival eggs injected were transferred to the oviducts of 18 pseudopregnant recipient mice, 8 of them became pregnant and gave birth to 20 F1 offspring. Of 20 offsprings, four males and two females carried the hybrid gene (HBx gene). Four male mice were determined as founder, named X1, X5, X9 and X15. These founders were back crossed to set up F1 generations with other ibred C57BL/6 mice or transgenic littermates, respectively. Transmission of HBx gene in F1 offspring of X1, X5 and X9 except in X15

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followed Mendelian rules. The expression of HBx mRNA was detected in liver of F1 offspring from the founder mice (X1 and X9), which showed vacuolation lesion and glycogen positive foci. CONCLUSION Transgenic mice harboring HBx gene were preliminarily established.

INTRODUCTION

Despite overwhelming epidemiological evidence linking persistent Hepatitis B Virus (HBV) infection and the development of hepatocellular carcinoma (HCC)^[1], the pathogenetic mechanisms by which the virus contributes to liver cell transformation remain elusive. The case is mainly the limited host range and the lack of *in vitro* culture systems to propagate it. With the development of embryo microinjection technology, leading to the production of transgenic animals, it provided a powerful tool for studies of gene expression and creation of animal models of human diseases.

The HBx gene is one of the four genes in the HBV genome. It encodes a viral transactivator, the HBx antigen (HBxAg). This protein has been shown to convert immortalized mouse fetal hepatocytes into a fully malignant phenotype. Moreover, NIH 3T3 cells stably transfected with a HBx expression plasmid are carcinogenic in nude mice. Interestingly, recently it has been reported that the P53 and HBx proteins can be coimmunoprecipitated from HBV-related HCCs. Complexing of P53 protein by HBx protein should its DNA consensus binding inhibit and transcriptional transactivator function and provide a basis for the ways in which this interaction might contribute to malignant transformation^[2].

These observations suggest. that HBx may play a part in the molecular pathogenesis of HCC in humans. In supprt of this hypothesis, it has been shown that high level expression og the HBx gene can lead to HCC in transgenic mice^[3-5], although others have not observed the induction of HCC in independently derived X gene trasgenic mouse strains^[6-8]. Noticeably, the mice that develop HCC

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were produced and maintained on a CD1 backround that displays a high spontaneous rate of HCC. This noght suggest that the HBx protein functions are a cofactor in the process of hepatocarcinogenesis, and that it may not be sufficient to induce HCC by itself^[9].

To investigate the exact role of the HBx gene in the process of hepatocarcinogenesis. The HBx transgenic mice were produced by microinjecting the HBx gene of HBV (subtype adr) DNA fragment into fertilized eggs derived from inbred C57BL/6 strain mice that displays a lower spontaneous rate of HCC, and its phenotype were preliminarily analyzed by pathologic examination and PAS.

MATERIALS AND METHODS

Materials

Various restriction enzymes and other reagents. Modifying enzymes ECOR I, Hind III, BamHI, T4 DNA ligase, dNTP, Taq DNA polymerase, MMLV reverse transcriptase, etc, purchased from Promega, Sangon and Sigma Compant, respectively.

Mice. C57BL/6 mice (clean animal) were purchased from Animal Center. Shanghai.

Methods

Construction of recombinant plasmid. PSHDX42, a 6787bp recombinant retroviral vector plasmid containing the *X* gene of HBV (subtype adr) was constructed by DNA recombinant technique (Figure 1) (Plasmid constructs is to be published elsewhere).



Figure 1 Structure of recombinant plasmid PSHDX42 X: HBx gene; ↓: BamHI

Production of HBx transgenic mice. Plasmid PXSHD42 was linearized with Ecor I and purified with QIAGEN. It was microinjected into the male pronucleus of the fertilized eggs of C57BL/6 mice. The injected eggs were then transplanted into the oviduct of pseudopregnany mice. Operated mice got pregnant and given birth.

Detection of transgenic integration. Genomic DNA was extracted from transgenic mice tail (about 1cm -1.5cm) at age of 2-3 weeks. Total DNA was extracted by the method described by Sambrook^[10].

Nested-PCR. The primers were designed on human HBX (subtype adr) X gene sequence (synthesized by Cybersyn Biotechmology Company). The sequence of primer for the PCR were as follows:

Outer primer 1: 5'-CATGGCTGCTCGGGTGTGCT-3' 2: 5'-ATTAGGCAGACGTGAAAAG-3' Inter primer 3: 5'-CTTTGTCTACGTCCCGTCGGCGCTGAATC-3' 4: 5'-CAGTCTTTGAAGTATGCCTCAAGGTCGGT-3

The PCR reaction I contained 5µ1 of 10×PCR Buffer, $3\mu 1$ of 25mM Mg²⁺, $3\mu 1$ of 15mM dNTP, 1.5µ1 of 20pmol each of outer primer, 2U of Taq DNA polymerase, 1µg samle DNA. A 50µ1 total reaction volume was obtained by adding sterile water, the reaction procedure of PCR I was: beginning 5min at 95°C, then 30 cycles of 94°C denature for 50 sec, ammealing at 55°C for 1min and extension at 72°C for lmin. The PCR reaction II contained 5µ1 of 10; ÁPCR buffer, 3µ1 of 25mM Mg^{2+} , $3\mu 1$ of 15mM dNTP, $3\mu 1$ of 10pmol each of inter primer,2.5U of Taq DNA polymerase, 1µl-2µ1 PCR I products. The reaction procedure of PCR II was as follows: beginning 5 minat 94°C, 30 cycles of 94°C denature for 50 sec, ammealing at 60°C for 50 sec and extension at 72°C for 1 min, and final extension at 72°C for 10min. A 1.8% agarose gel was prepared and loaded with 10µ1 PCR II products, then electrophoresed at 45vol for 70min, and the result of electrophoresis was observed.

Detection of transgenic expreseeion. Tissue samples were removed from F1 transgenic mice liver at the age of 8 weeks and frozen in liquid nitrogen. Total RNA was extracted by the method described by Sambrook^[10].

RT-PCR. The sequence of primers for RT-PCR were previously described as outer primer 1,2. First stranded cDNA was synthesized as follows: The RT reaction contained 1µg RNA,5µ1 of $10 \times PCR$ buffer, 3µ1 of 15 mM dNTP, 1µ1 of RNasin (80U/µ1), 2µ1 10pmol outer primer 1 or 2, 1µ1 MMLV (200U). A50µ1 total reaction volume was obtained by adding sterile DEPC treated water and RT was fouowed at 37°C for DNA ploymerase, 3µ1 pf 10pmol outer primer 1,2 respectively. The reaction procedure of PCR was described as PCR I.

Histological procedures and cytohistochemical staining.

Liver tissues from F1 transgenic mice at the age of 8 weeks were divided into two portions: one portion fixed in 10% buffered formalin for Hematoxylin and Eosin as well as PAS stainin, the other quickly frozen in liquid nitrogen and used for RNA analysis.

RESULTS

Establishment of transgenic mice

Recombinant retrovial DNA fragment was microinjected into the male pronucleus of the 514 fertilized gees of C57-B1/6 mice, and 368 survived eggs were transferred to the oviducts of 18 pseudopregnant recipient mice, 8 of ther became pregnant and gave birth to 20 offspring mice. The zygotes survial rate and birth rate were 71% (368/514) and 5.4% (20/368). DNA extracted from tail of 20 offspring mice were screened by PCR amplification (Figure 2), and the results showed that four males and two frmales carried the hybrid gene (HBx gene). Four male mice were determined as founders, named X1, X5, X9 and X15. Total integration rate and efficiency of transgene was 30% and 1.1%, respectively.

Establishment of HBX transgenic mice lineage The founders (X1,X5,X9 and X15) were back crossed to set up F1 offspring with other inbred C57BL/6 mice or transgenic littermates. Of 20 F1 mice from X1, 9 F1 mice carried the hybrid gene; of 18 F1 mice from X5, 8 F1 mice carried the hybrid gene; of 17 F1 mice from X9, 8 F1 mice carried the hybrid gene. Surprisingly, no mice integrated from F1 generation of X15 were detected by PCR.

Expression of HBx gene in F1 mice

We performed RT^{*2}PCR to examine whether HBx transgenes experess in liver tissues of F1 mice of transgenic mice lineages (X1,X5 and X9). The PCR products (439bp) were detected in the liver of F1 mice of transgenic mouse lineage (X1,X9) (Figure 3). NO Band was detected in the liver of F1 offspring of transgenic mouse lineage X5.

Phenitypic characteristics of transgenic F1 mouse

The liver from the F1 offspring of lineage (X1,X9) showed the presence of areas of altered hepatocytes which were made up of cells with pooly stained cytoplasm and PAS positive foci (Figure 4), but not in thir normal littermates.





Figure 2 Electrophoresis analysis of Nested PCR products amplified from DNA obtained from tail tissue. lane 1, normal mouse (negative control; lane 2 and 3 (transgenic mice) showed positive bands (280bp). M, PCR marker (SABC).

Figure 3 Electrophoresis analysis of RT-PCR products. Lane 1, normal mouse liver tissue (negative control); lane 2 and 3 transgenic mice liver tissue showed positive bands (439bp). M, PCR standard marker (SABC).

Figure 4 Two-month old transgenic mouse liver showed glycogen positive foci PAS×200

DISCUSSION

Despite the increased frequency in the generation of transgenic animals, the methodologies for introducing exogenous DNA into mammalian early embryos have remained essentially unchanged since the pioneering work of Gordon in the early 1980s^[11]. To date, the most widely used method is microinjection of foreign DNA into the pronuclei of fertilized eggs. The method has enabled the production of transgenic animals from various mammalian species, but the success rate of generating a transgenic animal is extremely low. In this study, the zygotes survival rate and birth rate were 71% and 5.4% and efficiency of transgene was 1.1%. The lowas feeiciency of transgene may be influenced by many factors such as DNA structure, purity and concentration, gene vector and fertilized eggs as well as mouse strains, etc^[12]. The result of F1 generation mice integration demonstrated that foreign gene can stably transmit to next generation in a normal Mendelian fashion. As for F1 generation from founder X5 low integration efficiency, it was thought that the founder X5 was likely to mosaics. The mechanism of DNA integration into mouse chromosomes following microinjection is unknown. One possibility is that spontaneous breaks occur in chromosome, possibly exacerbated by the microinjection technique and DNA repair endonuclease, and these breaks are sites for integration of linear DNA^[12]

When the transgene results from random integration of a DNA fragment injected into fertilized eggs the pronucleus, its level of expression in the resulting transgenic animal varies and depends upon factors, such as the site, transgene cope number, genetic background of mouse, particularly promoter/enhancer of fusion gene. In this study, HBx gene under the control of retrovial promoter/ enhancer LTP and SV40 promoter, the expression of HBx gene was detected by RT-PCR in liver from partial transgenic mice lineage (X1, X9). The results demonstrated that the retrovial promoter/ enhancer LTR and SV40 promoter were probably functional, and HBX gene expression was influenced by different integrated site.

Above all initial histological examination of liver in HBX gene transgenic mice revealed the presence of patholoical change (vacuolation lesions) and glycogen positive foci. The results of phenotype observation were similar to KIM's^[3], and were consistent with their preneoplastic lesions, and further identified HBx transgenic mice model established. To sum up, we have produced transgenic mice harboring HBx gene. This work has laid a solid foundation for further study on the function of JBx gene in hepatocarcinogenesis, successively analysis on these transgenic mice are being undertaking.

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