Evaluation of miR-122-regulated suicide gene therapy for hepatocellular carcinoma in an orthotopic mouse model

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Objective: Intratumoral administration of adenoviral vector encoding herpes simplex virus (HSV) thymidine kinase (TK) gene (Ad-TK) followed by systemic ganciclovir (GCV) is an effective approach in treating experimental hepatocellular carcinoma (HCC). However, hepatotoxicity due to unwanted vector spread and suicide gene expression limited the application of this therapy. miR-122 is an abundant, liver-specific microRNA whose expression is decreased in human primary HCC and HCC-derived cell lines. These different expression profiles provide an opportunity to induce tumor-specific gene expression by miR-122 regulation. **Methods:** By inserting miR-122 target sequences (miR-122T) in the 3' untranslated region (UTR) of TK gene, we constructed adenovirus (Ad) vectors expressing miR-122-regulated TK (Ad-TK-122T) and report genes. After intratumoral administration of Ad vectors into an orthotopic miR-122-deficient HCC mouse model, we observed the miR-122-regulated transgene expression and assessed the antitumor activity and safety of Ad-TK-122T. **Results:** Insertion of miR-122T specifically down-regulated transgene expression *in vitro* and selectively protected the miR-122-positive cells from killing by TK/GCV treatment. Insertion of miR-122T led to significant reduction of tansgene expression in the liver without inhibition of its expression in tumors *in vivo*, resulting in an 11-fold improvement of tumor-specific transgene expression. Intratumoral injection of Ad vectors

mediated TK/GCV system led to a vector dosage-dependent regression influencies in the insertion of miR-122T does not influence the antitumor effects of suicide gene therapy. Whereas mice administrated with Ad-TK showed severe lethal hepatotoxicity at the effective therapeutic dose, no liver damage was found in Ad-TK-122T group. **Conclusions:** miR-122-regulated TK expression achieved effective anti-tumor effects and increased the safety of intratumoral delivery of adenovirus-mediated TK/GCV gene therapy for miR-122-deficient HCC.

Keywords: Suicide gene therapy; microRNA-122 (miR-122); hepatocellular carcinoma (HCC); adenovirus (Ad); thymidine kinase (TK)



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Introduction

Hepatocellular carcinoma (HCC) is one of the most common cancers worldwide and the third most common cause of death from cancers (1). Unresectable HCC lacks effective therapy and new therapeutic modalities are urgently needed (2,3). Suicide gene therapy is a rational, potential therapeutic approach for the treatment of HCC and liver metastases of digestive tumors (2,3). One specific suicide gene therapy strategy is the virusmediated transduction of herpes simplex virus (HSV) thymidine kinase (TK) gene followed by the systemic administration of ganciclovir (GCV). The TK gene product phosphorylates GCV to activate its potential to terminate DNA synthesis, leading to cell death (4,5). Not only were

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cells transduced by TK gene killed, but also TK-negative cells were killed as a result of a "bystander effect" (6,7). Among the commonly used gene transfer vectors, adenovirus (Ad) vectors are particularly attractive for *in vivo* gene therapy of cancers (8). Several studies have demonstrated the efficacy of intratumoral injection of Ad-mediated TK/GCV system for the treatment of HCC (9-13). In the model of HCC, tumors can be regressed and even eliminated in vivo by the TK/GCV system in an Ad vector dosage-dependent manner. One of the main obstacles limiting the application of this therapy is toxic side effects affecting normal liver tissue. Because of the high hepatic tropism of Ad vectors, an unwanted spread of vectors and TK expression in peritumoral and remote liver tissue lead to severe hepatic damage (12-15). The severe hepatoxicity occurred in TK/ GCV-treated animals especially at effective dosage and resulted in the failure of therapy. Stringent regulation of transgene expression is required to increase the safety and

efficacy of suicide gene therapy for HCC.

A vast post-transcriptional regulatory network mediated by microRNA (miRNA), a class of abundant small noncoding RNAs, provides improved control over gene expression (16). Mature miRNAs perfectly, or near perfectly, bind to complementary sequences in the 3'-untranslated regions (UTRs) of target mRNAs and cause either translational repression or mRNA degradation (17). Tissuespecific miRNA expression has been well clarified (18) and provides an opportunity to regulate transgene expression from therapeutic nucleic acids and viruses. miRNA target sequences have been inserted into the 3'-UTR of transgene to restrict its expression in a specific cell or tissue (16). MicroRNA-122 (miR-122) is the most abundant liver-specific miRNA (18,19). Studies show that miR-122 is significantly down-regulated in 50-70% of human primary HCC and in all HCC-derived cell lines (20-27). Since the decrease in hepatic miR-122 was a tumor-specific event, we hypothesized inserting miR-122 target sequences (miR-122T) into the 3'-UTR of the TK transcription cassette would improve tumorspecific expression by repressing TK expression in nonmalignant liver tissue, and as a result, reduce hepatotoxicity, while keeping the antitumor effects. In this study, we have constructed Ad vectors expressing miR-122-regulated TK (Ad-TK-122T) and report genes. By intratumoral delivery of these Ad vectors to a murine orthotopic HCC model based on the miR-122-deficient murine HCC cell line Hepa 1-6 (26), we have researched miR-122-regulated gene expression in HCC and liver, the anti-tumor effects and hepatotoxicity of miR-122-reglated suicide gene therapy for HCC.

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Materials and methods

Cell lines

HEK-293, HeLa, Huh7 and Hepa 1-6 cells were purchased from the American Type Culture Collection. All cells were grown in DMEM with 10% fetal bovine serum, 1% penicillin/streptomycin, and 1% L-glutamine, cultured in a humidified incubator at 37 °C with 5% CO_2 .

Animals

Five-week-old male C57BL/6 mice were purchased from HFK bioscience Co., Ltd. (Beijing, China) and housed in the Animal Care Center of the Chinese Center for Disease Control and Prevention under pathogen-free conditions. All procedures involving animals and their care were conducted in conformity with institutional guidelines, which are in compliance with national and international laws and guidelines.

Plasmids and Ad vectors

The AdMaxTM system was used to produce all Ad vectors for this research. Enhanced green fluorescent protein (EGFP), firefly luciferase (Fluc) and TK gene were cloned into the EcoR I and Bgl II sites of pDC316 which was a shuttle plasmid containing an expression cassette utilizing the CMV promoter. In this way, the EGFP, Fluc and TK expressing plasmids, named pDC316-EGFP, pDC316-Fluc and pDC316-TK, were constructed. Oligonucleotides containing four copies of miR-122 completely binding sites (forward: 5'-AGCTTCAAACACCATTGTCACACTCCA AGACCAAACACCATTGTCACACTCCAAGACCAAA CACCATTGTCACACTCCAAGACCAAACACCATGT CACACTCCAGAGCT-3', reverse: 5'-CTGGAGTGTG ACAATGGTGTTTGGTCTTGGAGTGTGACAATG GTGTTTGGTCTTGGAGTGTGACAATGGTGTTT GGTCTTGGAGTGTGACAATGGTGTTTGA-3') (the miR-122T sequences were underlined) were annealed at 95 °C, cooled and ligated into pDC316-EGFP, pDC316-Fluc or pDC316-TK cleaved with Hind III and Sac I. In this way, miR-122-regulated EGFP, Fluc, or TK expression plasmids, named pDC316-EGFP-122T, pDC316-Fluc-122T and pDC316-TK-122T, were constructed. All plasmids were cotransfected with an Ad genomic plasmid into HEK293 cells using the AdMax[™] system to produce Ad-EGFP, Ad-EGFP-122T, Ad-Fluc, Ad-Fluc-122T, Ad-TK and Ad-TK-122T (Figure 1), and further propagated in



Figure 1 Schematic diagram of Ad vectors used in this study. Ad vectors expressing miR-122-regulated TK and report genes were constructed by inserting miR-122 target sequences (miR-122T) into the 3'-UTR of the transcription cassettes. HSV-TK, herpes simplex virus thymidine kinase; EGFP, enhanced green fluorescent protein; Fluc, firefly luciferase; CMV, cytomegalovirus promoter; BGH pA, bovine growth hormone polyadenylation signal.

HEK293 cells. All Ad vectors were purified by two rounds of cesium chloride density gradient ultracentrifugation. Viral particle (vp) titers were determined by spectrophotometry at OD_{260} .

In vitro gene expression analysis

HeLa, Huh7 or Hepa 1-6 cells were infected with Ad-EGFP-122T or Ad-EGFP respectively. Forty-eight hours later, EGFP expression was observed in fluorescence microscope. Aforementioned cells were infected with Ad-Fluc-122T or Ad-Fluc respectively. Forty-eight hours later, relative light units (RLU) in cultured cells were measured using the commercial luciferase assay system (Promega), according to the manufacturer's instructions in a luminometer.

In vitro MTT assay

HeLa, Huh7 or Hepa 1-6 cells were infected with Ad-TK-122T or Ad-TK respectively and cultured in 96-well plates with 150 μ L complete medium in the presence of different concentrations of GCV for 5 d. On day 5, 20 μ L 3-(4,5)-dimethylthiahiazo(-z-y1)-3,5-diphenytetrazoliumromide (MTT) solution (5 mg/mL) was added to each well, and incubation at 37 °C in 5% CO₂ was continued for 3 h. Supernatants were removed and 100 µL dimethylsulfoxide (DMSO) was added to each well. After 30 min incubation, absorbances were measured using a microplate spectraphotometer equipped with a 570 nm filter.

Establishment an orthotopic transplant mouse model of HCC

Five-week-old male C57BL/6 mice were anesthetized with pentobarbital sodium (75 mg/kg). A transverse incision was made across the xyphoid process and extended approximately 2 cm. Hepa 1-6 cells (1×10^6 /mouse) in 10 µL were injected into the front of the left upper liver lobe, using a 30-gauge needle. The lobe was returned to the abdomen and the incision was closed with sutures.

Intratumoral injection of Ad vectors

Ten days after injection of Hepa 1-6 cells in C57BL/6 mice, when the tumor volume was 50-100 mm³, surgery was performed as described earlier and 10 μ L of Ad vector was injected slowly and carefully into the centre of the tumor, using a 30-gauge needle.

In vivo gene expression analysis

Two days after Ad-EGFP or Ad-EGFP-122T, Ad-Fluc or Ad-Fluc-122T injected into tumors in HCC models, as described earlier, tumor and liver tissues were isolated and frozen sections were made. The EGFP expression was observation in fluorescence microscope and hematoxylin & eosin (HE) stain was also made to differentiate liver and tumor tissues. The whole liver tissue or tumor tissue was grinded respectively and RLU was estimated using the commercial luciferase assay system, according to the manufacturer's instructions in a luminometer.

TK/GCV treatment and in vivo antitumor effect study

Twelve hours after Ad-TK or Ad-TK-122T injected into tumors in HCC models, as described earlier, the mice were received intraperitoneal injections of GCV (35 mg/kg) twice every day for 5 d. Ten days after TK/GCV treatment, the mice were scarified and tumor growth was monitored by calculating the tumor volume from caliper measurements. Tumor volumes were calculated according to the formula: (tumor volume; mm³) = (major axis; mm) × (minor axis; mm)²/2.

Measurement of serum alanine aminotransferase (ALT)

Ten days after TK/GCV was performed, blood samples were collected through tail vein bleeding and allowed to clot (15 min, room temperature) and spun at 1,200 g for 10 min to collect the serum. The serum ALT levels were determined using an ALT kit (Human GmbH) according to the manufacturer's instructions in an automatic chemistry analyzer.

Histology

The liver lobes from each mouse were immersed in 10% buffered formalin overnight at room temperature, embedded in wax and sectioned using a vibratome. Sections were stained with HE and analyzed by light microscopy at $200 \times$ magnification.

Statistical analysis

Data were expressed as $\overline{x}\pm s$. Significance was evaluated using *t*-test and denoted on the graphs as *P<0.05, **P<0.01.

Results

In vitro performance of miR-122-regulated report genes and TK/GCV system

To assess in vitro miR-122-regulated transgene expression mediated by Ad vectors, Ad-EGFP-122T or Ad-EGFP was infected into HeLa, Huh7 or Hepa 1-6 cells at doses of 5,000, 1,000 or 1,000 vp/cell respectively. EGFP expression was analyzed by fluorescence microscopy 48 h later. As shown in Figure 2A, in Huh7 cells which are known to be miR-122-positive (19,26), the expression of EGFP was down-regulated dramatically. In contrast, the presence of miR-122T did not affect the EGFP expression in miR-122-negative HeLa cells (26,28). Moreover, insertion of miR-122T did not reduce the EGFP expression in Hepa 1-6 cells, suggesting the murine HCC cell line Hepa 1-6 was lack of miR-122 expression. This result was in conformity with previous studies which reported miR-122 expressed at very low levels in Hepa 1-6 cells (26). We also evaluated the miR-122-regulated Fluc expression in Huh7, HeLa and Hepa 1-6 cells by using Ad-Fluc-122T and Ad-Fluc. Cells were harvested 48 h after Ad infection and luciferase activities in the cells were detected (Figure 2B). As compared to Ad-Fluc, 8.06-fold repression of luciferase activity was found in Huh7 cells infected with

Ad-Fluc-122T (P<0.05). However, there was no inhibition of luciferase activities in HeLa and Hepa 1-6 cells infected with Ad-Fluc-122T. These results suggested miR-122's negative regulation was effective and specific *in vitro* for genes containing miR-122T.

To evaluate if miR-122 regulation would protect miR-122-positive cells from conditional killing, we infected respectively Ad-TK-122T or Ad-TK into Huh7, HeLa or Hepa 1-6 cells in the presence of GCV, and tested the viable cell number by MTT assay. As shown in *Figure 2C*, Ad-TK-122T-infected Huh7 cells were much less injured in the presence of GCV at doses of 1-100 µg/mL, compared to Ad-TK-transfected cells (P<0.05, P<0.01). However, HeLa cells infected with Ad-TK-122T or Ad-TK were both effectively killed by TK/GCV system. For Hepa 1-6 cells, infection with Ad-TK-122T resulted in the same cell death as infection with Ad-TK. These data indicated that inhibiting expression of TK gene by miR-122 downregulation was a feasible way to protect cells endogenously expressing miR-122 from killing by GCV.

In vivo examining miR-122-regulated transgene expression

Next, we evaluated the potency of miR-122 regulation in vivo. Ad-EGFP-122T at the dose of 1×10^{10} vp or the control Ad-EGFP at the same amount was injected intratumorally in the orthotopic Hepa 1-6 tumor model. After 2 d, we found that mice treated with Ad-EGFP showed a high level of EGFP expression not only in tumor but also in peritumoral liver tissue (Figure 3A). In contrast, mice treated with Ad-EGFP-122T showed no EGFP expression in peritumoral liver tissue, compare to the high level of EGFP expression in tumor tissue (Figure 3B). We also evaluated the repression capabilities of miR-122 by using Fluc gene. Two days after intratumoral injection of 1×10^{10} vp of Ad-Fluc, luciferase activities had been detected not only in tumor tissue but also in peritumoral and remote liver tissue (Figure 3C). The average luciferase activity in liver tissue (586,889±325,833 RLU) was higher than that in tumor tissue (323,456±23,778 RLU). The luciferase activity ratio of tumor to liver tissue was 0.55:1.00 (Figure 3D). In contrast to Ad-Fluc, intratumoral injection of Ad-Fluc-122T resulted in significant reduction of Fluc expression in liver tissue (50,354±22,802 RLU) (P<0.05), but mediated nearly the same level of Fluc expression in tumor tissue (307,307±77,497 RLU) (Figure 3C) as that of Ad-Fluc. The luciferase activity ratio of tumor to liver tissue was 6.10:1.00



Figure 2 *In vitro* performance of miR-122-regulated report gene and TK/GCV system. Ad vectors encoding report genes were infected into HeLa, Huh7 or Hepa 1-6 cells, and the expression of report genes was analyzed 48 h after infection. Ad-TK-122T or Ad-TK was infected into the aforementioned cells in the presence of GCV, and the viable cell numbers were tested by MTT assay 5 d after infection. (A) Fluorescent microscopy analysis (100x) of EGFP expression in cells infected with Ad-EGFP or Ad-EGFP-122T; (B) Luciferase activities of cells infected with Ad-Fluc or Ad-Fluc-122T; (C) TK/GCV-mediated cytotoxicity in cells infected with Ad-TK or Ad-TK-122T. *, P<0.05; **, P<0.01.

(*Figure 3D*). These results suggested that the expression of transgene containing miR-122T was inhibited strongly in the liver, but the inserting of miR-122T did not influence the transgene expression in the Hepa 1-6 tumor. As a result, the insertion of miR-122T acquires an 11.09-fold improvement of tumor-specific transgene expression.

Antitumor effects of miR-122 regulated TK/GCV system

Next, we evaluated the antitumor effects of adenovirusmediated miR-122 regulated TK/GCV system. Ad-TK-122T, Ad-TK or Ad-Fluc at doses of 1×10^9 , 3×10^9 or 1×10^{10} vp and normal saline (NS) control was intratumorally injected into orthotopic Hepa 1-6 tumor models following GCV treatment for 5 d. Ten days after TK/GCV treatment, the mice were scarified and tumor growth was monitored (*Figure 4A*). The results showed that injection of Ad-TK-122T and Ad-TK at the doses of 1×10^9 and 3×10^9 vp did not lead to significant antitumor effects as compared to Ad-Fluc injected control (*Figure 4B*). When the dose of Ad vectors increased to 1×10^{10} vp, the tumors of mice injected with Ad-TK and Ad-TK-122T were significantly repressed, compared to the Ad-Fluc injected control (P<0.01) (*Figure 4B*). There were no significant differences in the antitumor activity between Ad-TK-122T and Ad-TK. These results suggested that intratumoral injection of Ad-TK-122T or

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Figure 3 *In vivo* examining the miR-122-regulated transgene expression in an orthotopic HCC mouse model. The orthotopic Hepa 1-6 tumor models were intratumorally injected with Ad vectors encoding report genes, and the tumors and livers were isolated and subjected to EGFP or luciferase expression analysis 48 h after injection. (A,B) Fluorescent microscopy analysis (left) and HE staining (right) (100×) of EGFP expression in tumor and liver of HCC mouse models injected with Ad-EGFP (A) or Ad-EGFP-122T (B); (C) Luciferase activities in tumor and liver of HCC mouse models injected with Ad-Fluc or Ad-Fluc-122T (n=6); (D) Luciferase activity ratio of tumor to liver in HCC mouse models injected with Ad-Fluc or Ad-Fluc-122T (n=6). **, P<0.01.



Figure 4 Antitumor effects of miR-122 regulated TK/GCV system. Ten days after injection of Hepa 1-6 cells in mice, the established tumors were intratumorally injected with $1 \times 10^{\circ}$, $3 \times 10^{\circ}$ or 1×10^{10} vp/mouse of Ad-Fluc, Ad-TK, Ad-TK-122T (n=7-10). Then the mice received intraperitoneal injections of GCV (35 mg/kg) twice every day for 5 d. Tumor growth was monitored by calculating tumor size 10 d after injection of Ad vectors. (A) Experimental time line; (B) Tumor growth. Intratumoral injection of NS was the control. *, P<0.05; **, P<0.01.



Figure 5 Assessment of hepatotoxicity in mice after TK/GCV treatment. (A) Measurement of serum ALT levels; (B) Measurement of relative body weight before and after treatments; (C) Assessment of liver histology (200×). *, P<0.05; **, P<0.01.

Ad-TK resulted in a vector dosage-dependent regression in HCC. The insertion of miR-122T does not influence the antitumor effects of suicide gene therapy for the Hepa 1-6 tumor model by intratumoral injection of TK-expressing Ad vectors.

Assessment of bepatotoxicity of miR-122 regulation TK/ GCV system

To examine the hepatotoxicity of the adenovirus-mediated miR-122-regulated TK/GCV system, examinations of ALT, relative body weight and histology were performed 10 d after TK/GCV treatment. Mice administered Ad-TK at doses of 3×10^9 and 1×10^{10} vp showed significantly increased ALT levels (1.2 and 4.0 times higher than control mice treated with Ad-Fluc at the same doses, P<0.05 and P<0.01, *Figure 5A*). Mice administered Ad-TK-122T at doses of 3×10^9 and 1×10^{10} vp showed no significant increase

of serum ALT as compared to the control mice treated with Ad-Fluc. Moreover, whereas mice that treated with Ad-TK at 1×10^{10} vp showed a significant reduction in body weight compared to the Ad-Fluc control (P<0.01), mice treated with Ad-TK-122T at the same dose did not show any significant loss in body weight (Figure 5B). Histological analysis showed a dramatic difference between mice administered Ad-TK and those administered Ad-TK-122T at the dose of 1×10^{10} vp (*Figure 5C*). Injection of Ad-TK resulted in vacuolation, hemorrhaging and abnormal nuclear morphology in mouse livers. However, livers from mice administered Ad-TK-122T showed very little pathological changes. In addition, while one mouse in the study which received 1×10^{10} vp of Ad-TK became hunched and immobile, and died after 9 d, all mice treated with Ad-TK-122T or Ad-Fluc were alive during the experiment. These results indicated that insertion of miR-122T effectively avoided the hepatotoxicity caused by

intratumoral delivery of adenovirus-mediated TK/GCV suicide gene therapy.

Discussion

miR-122 is highly and selectively expressed in hepatocytes and attracts much attention to be considered as elements for hepatic specific targeting viral vector (18,19,29,30). Several studies have confirmed that insertion of miR-122T is an effective strategy to specifically down-regulate gene expression in liver. By incorporating miR-122T in the 3'-UTR of the E1A gene, the recombinant Ad replicated normally in other cells but not in cells of hepatic origin (31,32). miR-122T could selectively down-regulate adenoviral luciferase expression in the liver and improve tumor-specific transgene expression in a human fibrosarcoma model (33). miR-122T incorporated in adeno-associated virus (AAV) vectors efficiently inhibited transgene expression in the liver (34). Adenovirus-mediated suicide gene therapy using the TK gene with miR-122T was reported to prevent hepatotoxicity when the Ad vector was injected locally into melanoma B16 tumor-bearing mice (35). miR-122 was significantly down-regulated in 50% of HCC compared to non-malignant liver tissue from the same individuals and was not detectable in HCC cell lines such as human (HepG2, Hep3B) or mouse (Hepa 1-6) cells and expressed at a significantly low level in human (Huh7) and rat (H4) cells compared to the hepatocytes (20-24). The different expression profiles in HCC as well as the specificity and abundance in liver make miR-122 attractive to be considered as elements for tumor-specific gene expression. In this study, we explored the efficiency and safety of suicide gene therapy for HCC in an orthotopic Hepa 1-6 tumor model based on intratumoral injection Ad vectors that mediated miR-122-regulated TK expression.

By inserting four copies of miR-122T sequences in the 3'-UTR of EGFP, Fluc and TK genes, the recombinant Ad expressing miR-122-regulated report gene or suicide gene have been constructed. We confirmed that miR-122's negative regulation was effective and specific *in vitro* for genes containing miR-122T in the 3'-UTR; miR-122 down-regulation could protect cells endogenously expressing miR-122 (Huh7) from killing and did not affect killing the miR-122-negitive cells (Hepa 1-6 and HeLa). We established the mouse HCC models by orthotopic injection of miR-122-deficient Hepa 1-6 cells and researched the performance of intratumoral injection of Ad vectors expressing miR-122-regulated report genes. We found intratumoral injection of Ad-EGFP or Ad-Fluc of less 1/5

of tumor volume would result in report gene expressing not only in tumor but also in peritumoral and remote liver tissue. The average report gene expression levels in liver tissue were even higher than those in tumor tissue. In contrast, when Ad-EGFP-122T or Ad-Fluc-122T was injected intratumorally, the report gene expression was not altered in tumor tissue but was repressed significantly in liver tissue. miR-122-regulation resulted in an 11.09-fold improvement of Hepa 1-6 tumor-specific transgene expression. It suggested that insertion of miR-122T is an effective strategy to realize the tumor-specific tansgene expression for intratumoral injection Ad vectors in the orthotopic miR-122-down-regulated HCC model. The further results showed that the tumor growth of mice after injection of Ad-TK-122T and Ad-TK were both inhibited in a vector dose-dependent manner. Treatment with 1×10¹⁰ vp of Ad-TK or Ad-TK-122T acquired significant antitumor effects. The insertion of miR-122T did not affect the antitumor effects in the Hepa 1-6 tumor model. While intratumoral injection of Ad-TK at the effective dose $(1 \times 10^{10} \text{ vp})$ resulted in fatal hepatotoxicity, no significant difference in hepatotoxicity was found at mice administrated with Ad-TK-122T at this dose as compared with those treated with a control Ad vector. These results indicated that suicide gene therapy based on intratumoral injection of Ad vectors encoding miR-122-regulated TK gene was safer and more efficient by reducing hepatotoxicity perfectly in the HCC model.

Several factors contribute to the effective antitumor response and safety of adenovirus-mediated miR-122regulated TK gene therapy in this study. Hepa 1-6 cells which highly express coxsakievirus and adenovirus receptor (CAR) can be effectively infected by Ad vectors (36). Due to the lack of miR-122 in Hepa 1-6 cells, TK gene expression has not be impaired by insertion of miR-122T. These features ensure the sufficient TK gene expression in Hepa 1-6 tumor. Although liver tropism of Ad vectors could induce high transduction of TK gene in hepatocytes, the abundance of miR-122 in the live induced strong miR-122's negative regulation which led to dramatic repression of TK expression, avoiding hepatotoxicity.

A phase I clinical trial of TK-based suicide gene therapy in HCC indicated that Ad-TK can be safely administered by intratumoral injection to patients with HCC up to 2×10^{12} vp per patient (37). And an ealier phase I clinical trial indicated that Ad-TK can be safely administered by intratumoral injection in patients with hepatic metastases at doses up to 1×10^{13} vp per patient (38). The doses of Ad vectors used in our study were higher than those in clinical trials. These indicate that the miR-122-regulated TK/GCV system could be used to increase the injected dose of TK-expressing Ad vectors and thereby result in greater antitumor effects in clinical trials for HCC.

It should be pointed out that the miR-122-regulated TK expression system is not fit for cases of HCC with high-level expression of miR-122 which would result in insufficient TK expression levels in tumor and then treatment failure. Although miR-122's down-regulation is a common feature in human and mouse HCC, different cases may vary greatly in miR-122's levels. However, due to the liver-specific expression of miR-122, the miR-122-regulated TK expression system could be used to eliminate hepatic metastatic tumors in which miR-122 is negative. Beside suicide gene therapy, miR-122 regulation could be also used to avoid liver toxicity when cytotoxic and proapoptotic genes such as FASL, TNF- α and TRAIL are used as therapeutic genes in gene therapy for HCC.

In conclusion, miR-122-regulated TK expression achieves effective antitumor effects and increases the safety of adenovirus-mediated TK/GCV gene therapy for miR-122-down-regulated HCC. Insertion of miR-122T makes it possible to increase the dose of adenovirus-mediated TK/GCV gene therapy for HCC to get more effective therapeutic effect. Intratumoral injection of adenovirusmediated miR-122-regulated TK may be useful in the treatment of miR-122-down-regulated HCC.

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