

5-azacytidine promotes terminal differentiation of hepatic progenitor cells

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Abstract. 5-azacytidine (5-azaC) is known to induce cardiomyocyte differentiation. However, its function in hepatocyte differentiation is unclear. The present study investigated the *in vitro* capability of 5-azaC to promote maturation and differentiation of mouse embryonic hepatic progenitor cells, with the aim of developing an approach for improving hepatic differentiation. Mouse embryonic hepatic progenitor cells (HP14.5 cells) were treated with 5-azaC at concentrations from 0 to 20 $\mu\text{mol/l}$, in addition to hepatocyte induction culture medium. Hepatocyte induction medium induces HP14.5 cell differentiation. 5-azaC may enhance the albumin promoter-driven Gaussia luciferase (ALB-GLuc) activity in induced HP14.5 cells. In the present study 2 $\mu\text{mol/l}$ was found to be the optimum concentration with which to achieve this. The expression of hepatocyte-associated factors was not significantly different between the group treated with 5-azaC alone and the control group. The mRNA levels of ALB; cytokeratin 18 (CK18); tyrosine aminotransferase (TAT); and cytochrome p450, family 1, member A1 (CYP1A1); in addition to the protein levels of ALB, CK18 and uridine diphosphate glucuronyltransferase 1A (UGT1A) in the induced group with 5-azaC, were higher than those in the induced group without 5-azaC, although no significant differences were detected in expression of the hepatic stem cell markers, DLK and α -fetoprotein, between the two groups. Treatment with 5-azaC alone did not affect glycogen synthesis or indocyanine green (ICG) metabolic function in HP14.5 cells, although it significantly increased ICG uptake and periodic acid-Schiff-positive cell numbers amongst HP14.5 cells. Therefore, the present study demonstrated that treatment with 5-azaC alone exerted no effects on the

maturation and differentiation of HP14.5 cells. However, 5-azaC exhibited a synergistic effect on the terminal differentiation of induced hepatic progenitor cells in association with a hepatic induction medium.

Introduction

Liver transplantation, a surgical procedure used to replace a diseased liver with a healthy liver allograft, is the most commonly used technique for the treatment of liver failure and end-stage liver disease (1,2). Due to the limitations of this form of treatment, including a shortage of donor organs, high technical difficulty and the requirement for lifelong immunosuppression, cell therapy-based treatment strategies have been developed (3,4). Terminally differentiated hepatocytes exhibit powerful liver function of detoxification, metabolism and synthesis. However, their availability and low expansion efficiency *in vitro* are significant obstacles to hepatocyte transplantation (5). Hepatic progenitor cells (HPCs) are bipotential stem cells, which arise in the liver and are capable of differentiation into either hepatocytes or cholangiocytes, under the appropriate conditions. Embryonic HPCs exhibit self-renewal and differentiation potential, in addition to low immunogenicity, indicating that they may be a useful alternative source of hepatocytes (6,7). Although a number of researchers have reported that hepatic progenitor cells are able to differentiate *in vitro* and *in vivo* into hepatic cells with certain function, the differentiation efficiency of these cells for use as a transplantation substitute remains unclear (8,9). Therefore, it is necessary to develop techniques to stably and efficiently obtain mature functional hepatocytes from hepatic progenitor cells.

5-azacytidine (5-azaC) is one of multiple DNA methylase inhibitors that is able to reverse the methylation status of a gene, and restore its expression (10), and is currently the only known effective chemical compound with which to induce the differentiation of mesenchymal stem cells (MSCs) into myocardial cells (11,12). Changes in DNA methylation status affect the differentiation of stem cells (13,14). The derivation of hepatic progenitor cells from embryonic fetal liver cells is of value in the study of early human liver organogenesis, as well as in the creation of an unlimited source of donor cells for hepatocyte transplantation therapy (15). In the present study, it was

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Table I. RT-PCR primers (5'-3').

Gene	Forward	Reverse
GAPDH	GGCTGCCCAAGAACATCAT	CGGACACATTGGGGGTTAG
DLK	GCTGGGACGGGAAATTCT	AACCCAGGTGTGCAGGAG
AFP	ACGAGGAAAGCCCCTCAG	GCCATTCCCTCACCACAG
ALB	CCAGACATTCCCAATGC	CAAGTTCGCCCTGTGCAT
CK18	CTGGGCTCTGTGCGAACT	ACAGAGCCACCCCAGACA
TAT	ACCTTCAATCCCATCCGA	TCCCGACTGGATAGGTAG
CYP7A1	GATTCTGATGCTGTCTTACTT	CAATATCATTAGTGGTGCC

RT-PCR, reverse transcription-polymerase chain reaction; AFP, α -fetoprotein; ALB, albumin; CK18, cytokeratin 18; TAT, tyrosine amino-transferase; CYP7A1, cytochrome p450, family 1, member A1.

demonstrated that 5-azaC significantly increased the hepatic differentiation of embryonic hepatic progenitor HP14.5 cells in the hepatocyte induction medium. The present study assists in the development of effective strategies to induce hepatic progenitor cells differentiation and lays a foundation for the use of progenitor cells as seed cells for liver transplantation in order to treat disease resulting from liver injury.

Materials and methods

Cell culture and chemicals. HP14.5 cells were isolated from the livers of embryonic mice at day 14.5 post coitus, and immortalized with SV40 large T antigen as described previously (16). HP14.5 cells were cultured in Dulbecco's modified Eagle's medium (DMEM; Gibco Life Technologies, Carlsbad, CA, USA) supplemented with 10% (v/v) fetal bovine serum (FBS, Gibco Life Technologies), 100 units/ml penicillin, and 100 μ g/ml streptomycin at 37°C in 5% CO₂. The hepatic differentiation induction medium was composed of 0.1 μ M Dexmethesone (Dex)/ 10ng/ml Hepatic growth factor (HGF)/20 ng/ml Fibroblast growth factor-4 (FGF4) and 2% horse serum (HS, Hyclone Laboratories, Inc., Logan UT, USA) in DMEM. Unless indicated otherwise, all chemicals were purchased from Sigma-Aldrich (St. Louis, MO, USA).

Transfection of albumin promoter-driven *Gaussia luciferase* (ALB-GLuc) reporter and *Gaussia luciferase* reporter assay. HP14.5 cells were transfected with the pSEB-ALB-GLuc reporter vector (17). To construct pSEB-ALB-GLuc vector, the mouse ALB promoter gene was amplified by polymerase chain reaction (PCR) and subcloned into pSEB-GLuc retroviral vector to drive the expression of *Gaussia luciferase*. Following 24 h of transfection, cells were replanted to 24-well plates and treated with 5-azaC at concentrations of 0, 0.1, 0.2, 0.5, 1, 2, 5, 10 or 20 μ mol/l, with or without hepatic differentiation induction. *Gaussia luciferase* possesses a natural secretory signal, which is secreted into the cell medium. Thus, at the indicated time points, the medium was collected in order to detect the activity of *Gaussia luciferase*, using a *Gaussia Luciferase* Assay kit (New England Biolabs, Ipswich, MA, USA). Each assay was performed in triplicate and three independent experiments were conducted.

Reverse transcription-quantitative (RT-q)PCR. As previously described (18), the total RNA from each of the HP14.5 cell groups was extracted using the TRIzol (Invitrogen Life Technologies, Carlsbad, CA, USA) method. In order to generate cDNA templates, 10 mcg of total RNA was reverse transcribed with random hexamer pairs using Superscript II reverse transcriptase (Invitrogen Life Technologies, Carlsbad, CA, USA). PCR primers were designed using the Primer3 program to amplify the gene of interest (Table I). qPCR reactions were conducted using a Bio-Rad protocol as follows: 94°C for 20 seconds, 55°C for 20 seconds, and 70°C for 20 seconds, for 40 cycles. Plates were read after each cycle. Data were reported as the fold-change with endogenous GAPDH normalization.

Western blot assay. Western blotting was performed as previously described (18,19). Total proteins were extracted from treated HP14.5 cells, which were lysed in radioimmunoprecipitation assay buffer with PMSF (Beyotime Institute of Biotechnology, Shanghai, China). Approximately 20 μ g of total protein per lane was electrophoretically separated on a 10% SDS-polyacrylamide gel (Beyotime Institute of Biotechnology) and then transferred to a polyvinylidene fluoride membrane (Millipore, Billerica, MA, USA). The membrane was blocked with 5% non-fat milk in Tris-buffered saline with Tween-20 (TBST; Beyotime Institute of Biotechnology) at room temperature for 1 h and incubated with rabbit anti-ALB polyclonal antibody (1:200; cat. no. sc-50536; Santa Cruz Biotechnology, Inc., Dallas, TX, USA) or mouse anti- β -actin monoclonal antibody (1:200; cat. no. sc-47778; Santa Cruz Biotechnology, Inc.) antibodies at 4°C overnight. Following washing with TBST, the membrane was probed with the appropriate secondary antibody, conjugated with horseradish peroxidase (Santa Cruz Biotechnology, Inc., Dallas, TX, USA), at room temperature for 1 h. Protein expression was visualized using enhanced Chemiluminescent substrate (Kaiji, Nanjing, China) and exposed under the Syngene GBox Imaging system (Syngene, Cambridge, UK).

Immunofluorescence Staining. Briefly, at 12 days following induction, cells were fixed in ice-cold methanol for 15 min, permeabilized with 1% NP-40 and blocked with 5% bovine serum albumin. The cells were then incubated with primary

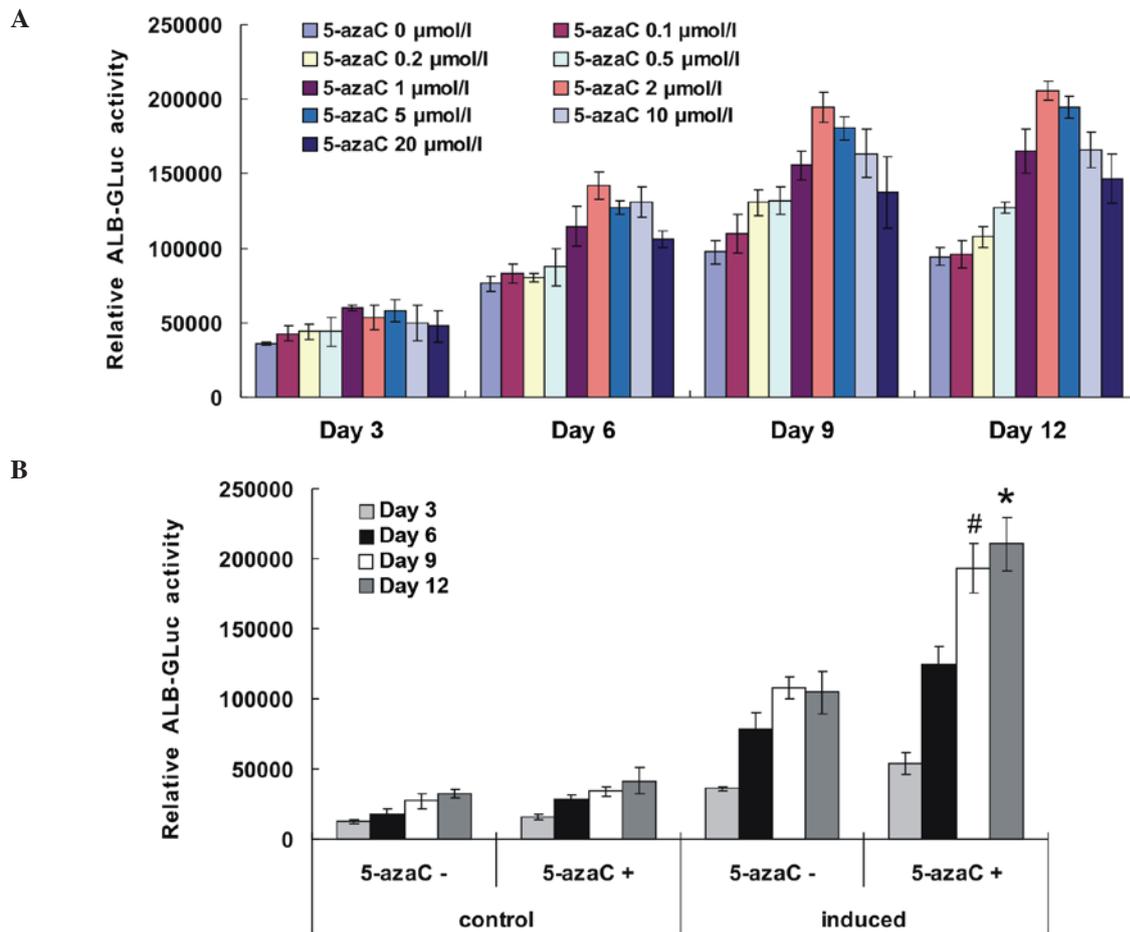


Figure 1. 5-azaC increased the ALB-GLuc activity of induced HP14.5 cells. (A) Effect of various concentrations of 5-azaC on ALB-GLuc activity in HP14.5 cells, cultured in a hepatocyte induction medium. Cells were transfected with pSEB-ALB-GLuc plasmid at 24 h prior to hepatic induction and treated with 5-azaC at concentrations of 0, 0.1, 0.2, 0.5, 1, 2, 5, 10 or 20 $\mu\text{mol/l}$, with hepatic differentiation induction. (B) HP14.5 cells were treated with 2 $\mu\text{mol/l}$ 5-azaC and hepatocyte induction medium. ALB-GLuc activity of HP14.5 cells was detected at 3, 6, 9 and 12 days following treatment. * $P < 0.05$, compared with the control group and # $P < 0.05$, compared with the group treated with induction medium alone. 5-azaC, 5-azacytidine; ALB-GLuc, albumin promoter-driven Gaussia luciferase.

goat anti-Cytokeratin 18 (CK18; 1:100; cat. no. sc-31700; Santa Cruz Biotechnology, Inc.) or rabbit anti-uridine diphosphate-glucuronosyltransferase 1A (UGT1A) polyclonal antibody (1:100; cat. no. sc-25847; Santa Cruz Biotechnology, Inc.) at 4°C overnight, followed by probing with DyLight 594- or 488-labelled secondary antibodies (Jackson ImmunoResearch Laboratories, West Grove, PA, USA) at room temperature for 30 min. Protein expression was examined under a fluorescence microscope (Nikon Intensilight C-HGF1; Nikon, Tokyo, Japan). Samples produced with control IgG were set up as negative controls.

ICG uptake and release. Cells were cultured in 24-well plates. At 12 days following treatment, cells were gently washed with PBS and incubated in 0.5 ml of complete DMEM medium, supplemented with 1mg/ml freshly-prepared cardiogreen at 37°C for 1 h. DMEM medium was then removed and the samples were gently washed several times with PBS. Green-stained cells were counted as ICG-positive cells under a microscope (Nikon Eclipse Ti-S; Nikon). Cells were then incubated in complete DMEM medium at 37°C for >6 h in order to assess ICG release, using a microscope (16,19). Ten nonoverlapping images were recorded.

Periodic acid-Schiff (PAS) staining. HP14.5 cells, cultured in 24-well plates, were treated for 12 days. Cells were fixed with 4% paraformaldehyde for 10 min and then incubated in 0.5% periodic acid solution for 5 min. Cells were then rinsed in ddH₂O for 3 min, incubated with Schiff's reagent for 15 min and counter-stained with hematoxylin solution for 2 min. Cells were subsequently thoroughly rinsed with tap water. All steps were performed at room temperature (16,19). Ten nonoverlapping visual fields were recorded using a microscope, and cells stained a purple-red color were counted as positive.

Statistical analysis. All data are presented as the mean \pm standard deviation, and were calculated using SPSS 15.0 software (SPSS, Inc., Chicago, IL USA). A two-tailed Student's t-test was used for statistical analysis. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

ALB-GLuc activity at various 5-azaC concentrations in induced mouse HP14.5 cells. 5-azaC, added at various concentrations to the hepatocyte induction culture medium, induced

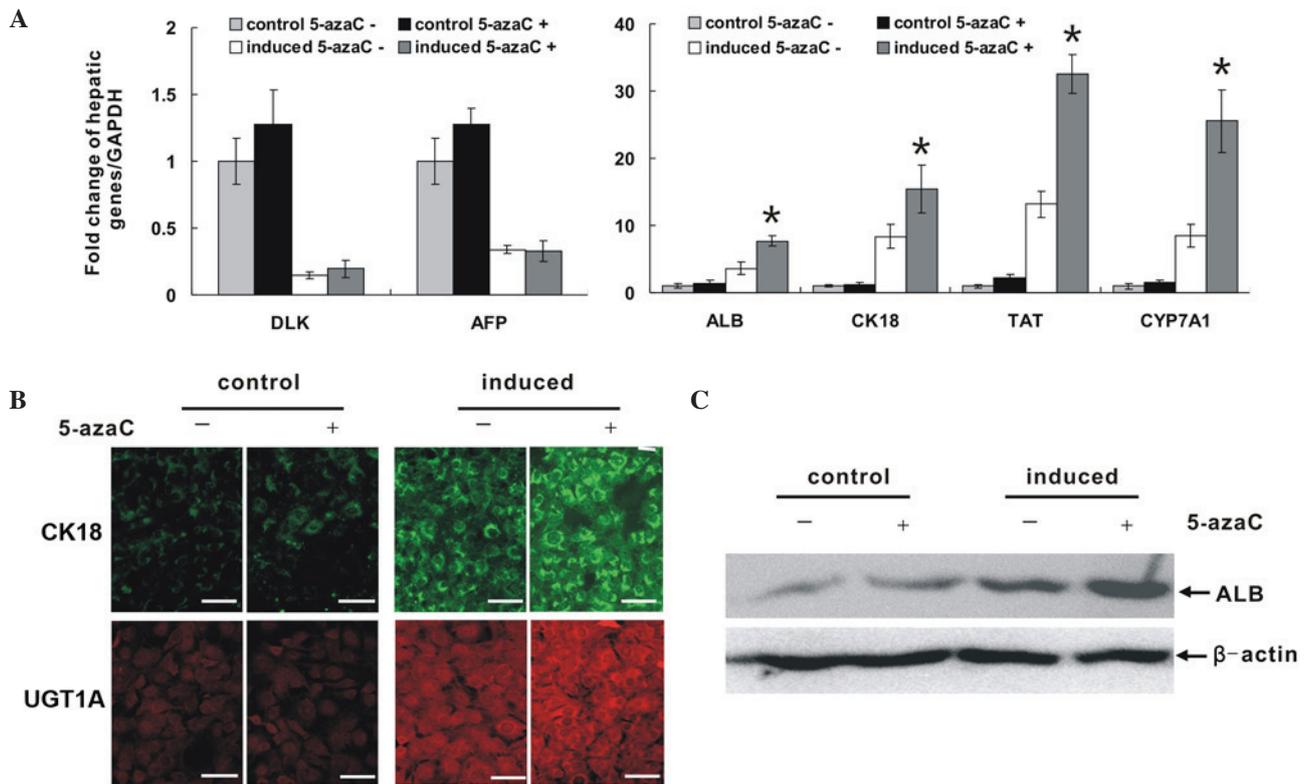


Figure 2. 5-azaC increased the expression of hepatic-associated markers of induced HP14.5 cells. HP14.5 cells were treated with 2 $\mu\text{mol/l}$ 5-azaC alone, hepatocyte induction medium alone, or a combination of the two, for 12 days. Untreated cells were used as a control. (A) mRNA expression of hepatic-associated marker genes, DLK, AFP, ALB, CK18, TAT and CYP7A1. Total RNA of cells in the different groups was extracted and reverse-transcribed into cDNA templates. The expression of genes was detected by qPCR and the fold of change was normalized to the expression GAPDH. qPCR results were confirmed in at least three independent experiments $^*P < 0.05$, compared with induction medium alone treated group. (B) Protein expression of CK18 and UGT1A was detected using immunofluorescence staining. Scale bar = 200 μm . (C) Expression of ALB was detected by western blot analysis. Equal loading of the samples was confirmed by β -actin. 5-azaC, 5-azacytidine; AFP, α -fetoprotein; ALB, albumin; CK18, cytokeratin 18; TAT, tyrosine aminotransferase; CYP7A1, cytochrome p450, family 1, member A1; qPCR, quantitative polymerase chain reaction; UGT1A, uridine diphosphate-glucuronyltransferase 1A.

the differentiation of HP14.5 cells. ALB-GLuc is transcribed from the ALB promoter and drives the luciferase reporter gene. Its activity indirectly reflects the level of ALB expression in cells, providing a useful means with which to detect hepatocyte maturation. At 6 days following induction, ALB-GLuc readings were higher in the induced group with 5-azaC than in the induced group without 5-azaC, and exhibited a progressive increase with increasing induction time, reaching a peak following induction for 9-12 days. It was shown that 2 $\mu\text{mol/l}$ 5-azaC was the optimal concentration for hepatic induction (Fig. 1A).

5-azaC enhances ALB-GLuc activity of HP14.5 cells treated with hepatocyte induction culture medium. The ALB-GLuc activity using 2 $\mu\text{mol/l}$ 5-azaC, with or without hepatocyte induction culture medium was subsequently detected. As shown in Fig. 1B, no significant difference in ALB-GLuc activity was observed between the group treated with 5-azaC alone and the control group. By contrast, 5-azaC significantly increased the ALB-GLuc activity of induced HP14.5 cells.

5-azaC increases the expression of hepatic-associated marker genes of HP14.5 cells in association with induction medium. qPCR results (Fig. 2A) showed that in the group treated with 5-azaC alone, the expression of various hepatic-associated factors increased slightly, although it was not significantly different compared with that in the control group. Hepatic

induction medium induced HP14.5 cells differentiation: The expression of DLK and α -fetoprotein (AFP), which are characteristic markers of hepatic stem cells, decreased significantly, while that of ALB, CK18, tyrosine aminotransferase (TAT) and cytochrome P450, family 7, subfamily A, polypeptide 1 (CYP7A1), which are mature hepatocyte markers, increased significantly. ALB, CK18, TAT and CYP7A1 expression in the induced group treated with 5-azaC, was higher than that of the induced group without 5-azaC. Western blotting results were consistent with these findings. 5-azaC treatment alone did not affect the expression of ALB protein, while it enhanced the expression of this protein in an induced environment (Fig. 2B). Immunofluorescence images demonstrated expression of the mature hepatocyte markers, CK18 and UGT1A, in cytoplasm, and no difference was detected between the group treated with 5-azaC alone and the control group. By contrast, the expression of these markers in the 5-azaC induced group was significantly higher than that in the induced group without 5-azaC ($P = 0.008597$; Fig. 2C). Thus, the results suggested that 5-azaC alone is insufficient to induce hepatic progenitor cell differentiation. However, it does stimulate hepatic maturation and differentiation within the appropriate induced environment.

5-azaC enhances the mature hepatic function of HP14.5 cells in association with induction. Mature hepatic cells are known to metabolize ICG, and ICG uptake may therefore be used for

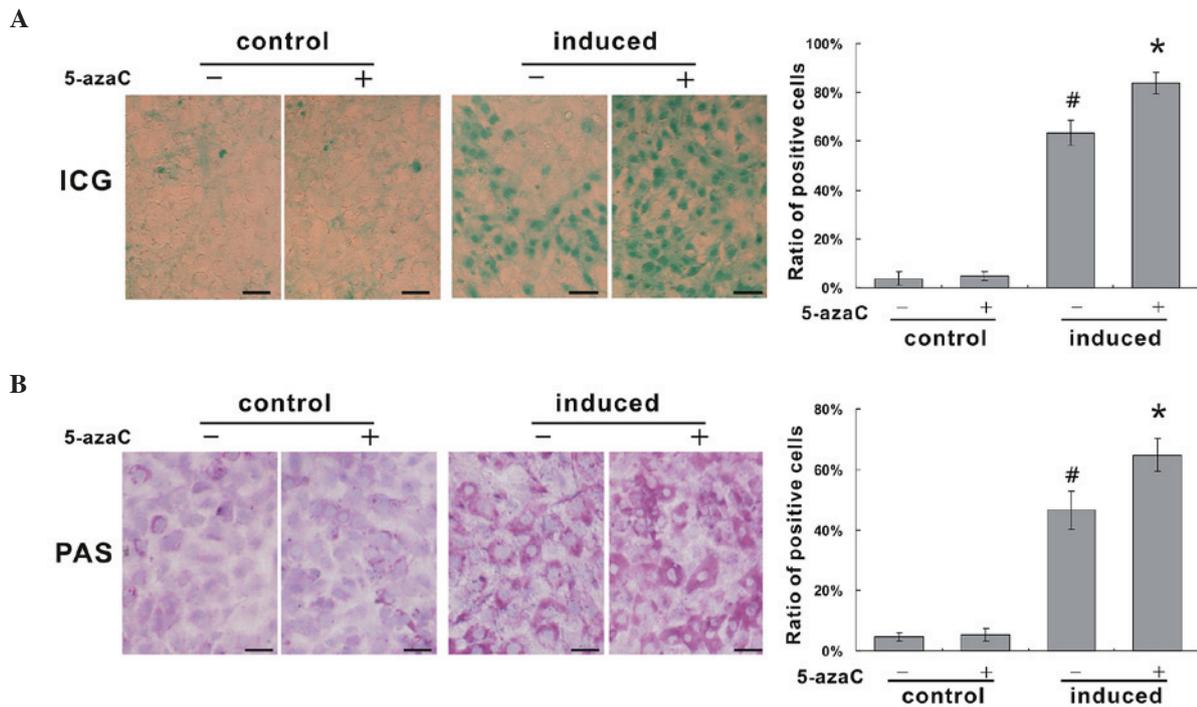


Figure 3. 5-azaC enhanced the mature hepatic function of HP14.5 cells in association with hepatic induction. Cells were treated as described in Figure 2. (A) Transport and metabolic function was evaluated by an ICG uptake assay. Positive cells exhibited a green-stained nucleus. (B) Glycogen storage and accumulation function in the induced HP14.5 cells was measured using a PAS staining method. Purple staining in the cell plasma indicated glycogen accumulation. Three independent experiments were performed. Scale bar = 200 μ m. Six nonoverlapping visual fields were randomly selected and cells were counted to calculate the ratio of positive to negative cells. #P<0.05, compared with the control group and *P<0.05, compared with the group treated with induction medium alone. 5-azaC, 5-Azacytidine; ICG, indocyanine green; PAS, periodic acid-Schiff.

the identification of differentiated hepatocytes *in vitro* (20,21). The ratios of ICG-positive cells to ICG-negative cells in the non-induced control group and in the group treated with 5-azaC alone, were $3.9 \pm 2.8\%$ and $4.8 \pm 1.9\%$, respectively, and no significant difference was detected between the two groups. Following 12 days of induction, the ratio of ICG-positive to ICG-negative cells was $63.4 \pm 5.1\%$, which was significantly higher than that of the control group, while it was significantly lower than the 5-azaC induced group ($83.9 \pm 4.5\%$; Fig. 3A).

Glycogen synthesis function is an important indicator in the evaluation of hepatic differentiation. Synthetic glycogen appears a purple-red color in the cytoplasm, upon PAS staining (22). The ability of HP14.5 cells to synthesize glycogen was weak, with a ratio of PAS-positive to PAS-negative cells of $4.6 \pm 1.4\%$. Treatment with 5-azaC alone, did not significantly increase this ratio ($5.3 \pm 2.1\%$). At 12 days following induction, the ratio of PAS-positive to PAS-negative HP14.5 cells increased to $46.6 \pm 6.3\%$, while the ratio in the induced group treated with 5-azaC was $64.7 \pm 5.4\%$, which was significantly higher than that in the induced group without 5-azaC. There was widespread purple-red throughout the cytoplasm, which was of a deeper color than that in the induced group (Fig. 3B). These results demonstrated that 5-azaC enhances the metabolic and synthetic function of HP14.5 cells in association with hepatocyte induction.

Discussion

Hepatic progenitor cells are a form of stem cell, which are able to self-proliferate; differentiate into hepatocytes and biliary

epithelial cells, and participate in liver repair and reconstruction. They may also be an important cell source for hepatic cell transplantation (7,23,24). Current research on hepatic stem cells remains at the theoretical and experimental stage. Extrahepatic sources of hepatic stem cells, including embryonic stem cells, hematopoietic stem cells and bone marrow MSCs are able to differentiate into hepatocyte-like cells that express hepatocyte-specific genes or exhibit partial hepatic cell function of regeneration *in vivo* (25,26). However, the differentiation efficiency and substitution function of these cells *in vivo* is far less than hepatic progenitor cells from embryos (27,28). The process of differentiation of stem cells into hepatocytes, includes bipotential hepatic progenitor cells. Embryonic hepatic progenitor cells are precursor cells, and may differentiate into hepatocytes or biliary epithelial cells. The intermediate bipotential state is therefore required during the differentiation of other types of stem cells into mature hepatocytes (26,29,30).

Hepatocyte differentiation is periodically regulated by different signals. In particular, the participation of associated cytokines is required to facilitate stem cell differentiation into hepatic endoderm (31-33). At the hepatic progenitor cell stage, in which cells exhibit bi-directional differentiation potential, the gene regulation system sends different instructions to the cells, in order to induce their differentiation into hepatocytes and biliary epithelial cells, thus producing a unique hepatic tissue morphology and function. It has been demonstrated that upregulating the expression of differential-related genes may improve the therapeutic efficacy of hepatic stem cells transplantation (34,35). Therefore, stable and efficient hepatocyte

resources obtained from hepatic progenitor cells may significantly improve the efficiency and biosafety profile of liver cell transplantation.

5-azaC is a type of cytosine chemical analogue, which is able to interfere with the physiological function of DNA by embedding in this molecule in order to exert cytotoxic and antitumor effects (10,36). 5-azaC has been used for the treatment of breast cancer, colon cancer, melanoma and acute myelogenous leukemia in clinical practice (37-39). 5-azaC also functions as a DNA methylation inhibitor. It combines with the methylation enzyme and inactivates it, which leads to hypomethylated DNA, thereby enhancing gene expression. In 1995, Wakitani *et al* (40) reported that 5-azaC induces the differentiation of bone marrow MSC into cardiomyocytes. 5-azaC has been widely used in a variety of directional stem cell induction. MSCs treated with 5-azaC *in vitro* express cardiac-specific structural protein (atrial natriuretic peptide, brain natriuretic peptide and α - β -myosin heavy chain) and cardiomyocyte-specific transcription factors (GATA4 and Nkx2.5/Csx), and partly improve myocardial systolic ventricular pressure and tension following transplantation into freezing damaged rat myocardium *in vivo* (41-43). Furthermore, brief exposure to 5-azaC induces pig dermal fibroblast reprogramming into insulin secreting cells (44). A high concentration (10 μ mol/l) of 5-azaC has been reported to enhance the induction of adipose-derived stem cell differentiation into myogenic cardiogenic cells (45).

5-azaC mediated-inhibition of DNA methylation is widespread rather than cell-specific. However, there are a number of studies that have investigated the effect of 5-azaC on hepatocyte differentiation and hepatic tumors. It has been reported that 5-azaC inhibits HepG2 and Hep3B liver tumor cell proliferation, induces apoptosis, and promotes their maturation and differentiation (46). The present study showed that 5-azaC exerts certain inducing differentiation effects, when administered at a suitable concentration. However, when treated with a high concentration condition of 5-azaC, the ALB-GLuc activity of induced HP14.5 cells decreased. This phenomenon may be due to higher doses of 5-azaC directly inhibiting cell proliferation, and mediating cell cytotoxicity by embedding into DNA and RNA (47,48). By contrast, lower doses of 5-azaC primarily inhibit DNA methylation, resulting in the recovery of gene normal expression (10,49,50). Concomitantly, it was found that hepatic-associated markers of HP14.5 cells treated with 5-azaC alone, did not increase significantly, suggesting that DNA hypomethylation is not the only factor that determines gene expression during the hepatic cell differentiation process; the appropriate induced culture conditions and the microenvironment also have an effect on transcriptional control. Similarly, MSCs may be induced to differentiate into cardiomyocytes by 5-azaC, although it is difficult to obtain functional beating myocardial cells (51). DLK and AFP proteins are markers of hepatic stem cells (52,53). If 5-azaC promotes the maturational differentiation of induced hepatic cells, the expression of these stem cell markers should decrease. However, as 5-azaC could inhibit DNA methylation, stem cell marker gene expression in the 5-azaC induced group remained at the same level as that in the induced group without 5-azaC treatment. The expression of CK18 and ALB, which are markers of mature hepatocytes, was higher in the induced group with 5-azaC than in the

induced group without 5-azaC. It was hypothesized that the different conformations of DNA cpG islands may determine the extent of the effect of 5-azaC, which functions primarily via modification of methylation (54,55). This proposal requires further investigation.

In conclusion, The present study demonstrated that 5-azaC synergistically promotes the hepatic differentiation of HP14.5 cells, significantly increases the expression of hepatic-associated marker genes, and enhances the ICG metabolism and glycogen synthesis function of these cells. The current study provides a basis for the clinical application of hepatic progenitor cells in liver disease. 5-azaC is known to induce differentiation into cardiomyocytes. The present study demonstrates that it is also involved in the terminal maturation and differentiation of induced hepatocytes, suggesting a wider role for this molecule as an inducing agent.

Acknowledgements

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