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Original Article
A preclinical study on the combination therapy of everolimus and transarterial chemoembolization in hepatocellular carcinoma

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Abstract: Transarterial chemoembolization (TACE) is commonly used for the treatment of locally advanced hepatocellular carcinoma (HCC) by its dual effects of chemotherapy and ischemic hypoxia. However, one of the side effects of TACE is the introduction of hypoxic condition, which in turn activates hypoxia-induced survival pathways and enhances VEGF-induced neovascularization by stabilizing HIF-1α expression. Herein, the preclinical therapeutic efficacy of the combined treatment of everolimus, a novel mTOR inhibitor and TACE for the treatment of HCC was investigated. The MHCC-97L cells were used for the study of the effect of combined treatment on cell proliferation and cellular apoptosis. HUVEC cells were used for the study on tube formation under different treatments. Inhibitions on the Akt/mTOR pathways were also studied. Finally, the effect on tumor growth was further study using an in vivo orthotopic model. The results demonstrated that everolimus enhanced the therapeutic efficacy of TACE in inhibiting cell proliferation, promoting apoptosis and inhibiting tube formation of endothelial cells by blocking the Akt/mTOR signaling pathway in vitro and inhibiting tumor growth and neoangiogenesis in vivo. Based on this preclinical study, the potential of combining everolimus with TACE was guaranteed which suggested the use of the combination therapy in the clinical treatment of advanced HCC patients.

Keywords: Everolimus, hepatocellular carcinoma, hypoxia, mTOR signaling pathway, transarterial chemoembolization, tumor growth

Introduction

Hepatocellular carcinoma (HCC) is the fifth most common cancer worldwide [1]. Surgery for HCC including partial liver resection and liver transplantation remains the most effective treatment for HCC. However, only a minority of HCC patients are eligible to the surgical treatment and poor prognosis with high recurrence rate is always observed [2-4]. In addition, adjuvant systemic chemotherapy has not been shown to be beneficial in HCC patients due to the chemoresistance of HCC cells and poor tolerance and side effects of anti-tumor drugs to cirrhotic patients [2, 5]. Therefore, it is worthwhile to develop more efficient treatment for unresectable HCC patients.

Transarterial chemoembolization (TACE) is commonly used for the treatment of unresectable HCC. It is done by injecting an embolic particles coated with chemotherapeutic agents such as cisplatin into the hepatic artery. It then works by blocking the blood supply to the tumor as well as delivering chemotherapeutic drug directly to the tumor. TACE was found to prolong the overall survival of patients with unresectable HCC compared with conservative treatment in two randomized trials [6, 7]. However, some studies showed that TACE cannot completely eliminate the risk of a recurrence [8, 9]. In addition, blocking the blood supply induces hypoxic condition, which in turn stabilizes the hypoxia inducible factor 1α (HIF-1α) protein. HIF-1α activates the transcription of genes that are involved in angiogenesis, cell survival, and cell migration and invasion which are crucial to cancer growth and recurrence [10]. Therefore, there is still room for improvement to enhance the tumor eliminating effect of TACE in order to prevent recurrence in HCC.
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The mammalian target of rapamycin (mTOR) is a serine/threonine kinase and member of the phosphoinositide 3-kinase (PI3K)-related kinase family. It integrates and transmits signals from a diverse array of signaling pathways to regulate cell survival and growth [11]. mTOR exists in two complexes, mTORC1 and mTORC2, which controls cell growth and cell survival, respectively. Several reports have indicated that mTOR is a positive regulator of hypoxia inducible factor 1 (HIF-1) expression and activity [12], and the inhibition of HIF-mediated gene expression is considered to be related to the anti-tumor activity of mTOR inhibitors [13]. In addition, HIF-1 activates vascular endothelial growth factor (VEGF) gene expression [14]. Therefore, the Akt/mTOR signaling pathway is also important in regulating neovascularization which is a crucial factor for the success of TACE treatment [15].

Everolimus is an mTOR inhibitor which is currently used as an immunosuppressant [16-18] and as a molecular targeting drug in cancers such as renal cell carcinoma and breast cancer. Our previous study has demonstrated that everolimus suppressed HCC tumor growth and sensitized tumor cells to cisplatin-induced cytotoxicity [19]. Therefore, we hypothesized that everolimus enhances the therapeutic efficacy of TACE by inhibiting the mTOR mediated HIF-1 expression and sensitizing the cisplatin effect on HCC tumor cells. The combination effect of everolimus and cisplatin under hypoxic condition was first studied by in vitro experiments. We then mimicked the TACE treatment using a mouse orthotopic HCC model by hepatic artery ligation (HAL) and intraportal vein injection of cisplatin. The effect of everolimus on this in vivo model was then investigated. Therefore, with this study, the pre-clinical enhancing effect of everolimus on TACE treatment can be demonstrated which provides the basis for future clinical practice of everolimus on TACE treatment.

Materials and methods

Drugs and reagents

Everolimus was kindly provided by Novartis Pharmaceuticals Corporation. All reagents were purchased from Sigma-Aldrich (St. Louis, MO) unless specified below.

Cell culture and treatment

MHCC97L cells (Liver Cancer Institute, Fudan University) were maintained in DMEM (Life Technologies, Carlsbad, CA) supplemented with 10% heat-inactivated fetal bovine serum (Life Technologies) and 1% penicillin/streptomycin (Life Technologies) at 37°C in a humidified 5% CO₂ atmosphere. MHCC97L cells were stably transfected with luciferase expressing construct for the ease of detection in the in vivo study. Human umbilical vein endothelial cells (HUVEC; Life Technologies) were grown in a complete endothelial growth medium (Life Technologies) and were used between passages 4-6.

For the in vitro study, cells were treated as 1) Control group: in normoxic condition (20% O₂, 5% CO₂ and 75% N₂) without any drug treatment, 2) Hypoxia group: in hypoxic condition (1% O₂, 5% CO₂ and 94% N₂) maintained in an OxyCycler C42 hypoxia chamber (Biospherix, St. Lacona, NY), 3) Hypoxia + C group: in hypoxic condition and 6 μM cisplatin, 4) Hypoxia + E group: in hypoxic condition and 10 nM everolimus, and 5) Hypoxia + C + E group: in hypoxic condition, 6 μM cisplatin and 10 nM everolimus.

Cell proliferation assay

The treatment effect on MHCC97L cells was first examined by MTT assay as previously described [20]. Briefly, cells were plated in 96-well culture plates for 24 hours and media were replaced with the treatment media with increasing concentrations of everolimus (0 to 100 nM) under hypoxic condition was first studied by in vitro experiments. We then mimicked the TACE treatment using a mouse orthotopic HCC model by hepatic artery ligation (HAL) and intraportal vein injection of cisplatin. The effect of everolimus on this in vivo model was then investigated. Therefore, with this study, the pre-clinical enhancing effect of everolimus on TACE treatment can be demonstrated which provides the basis for future clinical practice of everolimus on TACE treatment.

Apoptosis assay

After MHCC97L cells were treated for 48 hours, cells were harvested, washed with ice-cold PBS, and stained with annexin V-FITC antibody (BD Biosciences Pharmingen). The percentage of apoptotic cells (annexin V positive cells) were determined by flow cytometry using the Cytomics FC500 Analyzer (Beckman-Coulter,
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Figure 1. Cell proliferation of MHCC-97L cells. Cells were treated with increasing concentrations of everolimus under normoxic, hypoxic or hypoxic condition with 6 μM cisplatin. Cell proliferation was analyzed by MTT assay. Data were presented as means ± SD from three independent experiments.

Figure 2. Cellular apoptosis of MHCC-97L cells. Cells were treated with Control: in normoxic condition (20% O$_2$, 5% CO$_2$, and 75% N$_2$) without any drug treatment, Hypoxia: in hypoxic condition (1% O$_2$, 5% CO$_2$, and 94% N$_2$), Hypoxia + C: in hypoxic condition and 6 μM cisplatin, Hypoxia + E: in hypoxic condition and 10 nM everolimus, and Hypoxia + C + E: in hypoxic condition, 6 μM cisplatin and 10 nM everolimus. The percentage of apoptotic cells were the percentage of annexin V positive cells analyzed by flow cytometry. Data were presented as means ± SD from three independent experiments and statistical analysis were performed by one-way ANOVA. *P<0.05 versus Control. #P<0.05 of Hypoxia + C + E versus Hypoxia, Hypoxia + C or Hypoxia + E.

CA). Unstained cells were used as a negative control.

Tube formation assay

The tube formation ability of the HUVEC cells was determined by the Endothelial Cell Tube Formation Assay (Life technologies) according to the manufacturer’s instructions. The bottoms of a 96-well plate were first coated with Geltrex™ Reduced Growth Factor Basement Membrane Matrix and 5 × 10^4 HUVEC cells were seeded on the surface of the gel and after treatment for 6 h, cells were stained with Calcein AM (Life technologies) and assessed under a fluorescence microscope (400 × magnification).

Multiplex bead assay

After MHCC97L cells were treated for 24 hours, cells were harvested and washed in ice-cold PBS. The expression levels of phosphorylated mTOR (Ser2448), p70S6K (Thr412), RPS6 (Ser235/Ser236), Akt (Ser473), TSC2 (Ser939), GSK3α (Ser21), GSK3β (Ser9), and p53 (Ser15) were quantified using a bead-based MILLIPEX® multiplex assay (Millipore, Billerica, MA) according to the manufacturer’s instructions. Briefly, cells were lysed in MILLIPEX MAP lysis buffer and diluted with an equal volume of MILLIPEX MAP cell assay buffer. Capture antibody beads were diluted in MILLIPEX MAP cell assay buffer and added to a Beadlike filter plate. 25 μl of the diluted cell lysate was then transferred to each well of the filter plate and incubated for 2 h at room temperature with shaking. After incubation, the beads were washed twice with cell assay buffer, and biotinylated detection antibodies diluted in cell assay buffer were added and incubated for 1 h at room temperature with shaking. After filtering, MILLIPEX MAP streptavidin-phycoerythrin solution was added and incubated for 30 min at room temperature with shaking. Finally, cell assay buffer was added after filtering, and signals were read using the Bio-Plex Luminex 200 system (Bio-Rad, Hercules, CA). The fluorescence signal of each phosphorylated protein was normalized with that of GAPDH.

Immunoblotting

After treatment, cells were lysed by ice-cold RIPA buffer containing 150 mM NaCl, 1 mM EDTA, 1% (v/v) NP-40, 0.25% (w/v) sodium deoxycholate, 1 mM PMSF, and protease inhibi-
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tor cocktail (Roche Diagnostics, Penzberg, Germany) in 50 mM Tris-HCl buffer, pH 7.4. Equal amount of protein was loaded onto a 10% SDS-polyacrylamide gel under reducing condition and transferred to PVDF membrane (Amersham Bioscience, Piscataway, NJ). Blots were probed with HIF-1α (Novus Biologicals, Littleton, CO), VEGF (Santa Cruz Biotechnology) and β actin (Sigma) overnight at 4°C. After washing, membranes were incubated with horseradish peroxidase-conjugated secondary antibodies (DAKO) and visualized using an enhanced chemiluminescence detection system (ECL Plus) (Amersham) according to the manufacturer’s instructions.

**In vivo orthotopic mouse HCC model**

All the animal experiments were approved by the Committee on the Use of Live Animals in
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Teaching and Research of The University of Hong Kong. Male CB17 severe combined immuno-deficient (SCID) mice of four to six weeks old were used and maintained in laminar flow cabinets under pathogen-free condition. Subcutaneous tumors were first grown by injecting MHCC97L cells (1 × 10^6) in a 0.2 ml culture medium subcutaneously into the right flank of mice. Once the subcutaneous tumors reached 0.8-1 cm in diameter, mice were sacrificed and tumors were removed and cut into 2 mm³ cubes, which were then implanted into the left liver lobes of another group of SCID mice. Six days after tumor implantation, the animals were randomly divided into five groups: 1) sham operation (Control; n=5), 2) hepatic artery ligation (HAL; n=5), 3) hepatic artery ligation combined with portal vein injection of cisplatin (3 mg/kg) (HAL + C; n=5), 4) hepatic artery ligation combined with oral feeding of everolimus daily (10 mg/kg) (HAL + E), and 5) HAL combined with portal vein injection of cisplatin (3 mg/kg) and oral feeding of everolimus daily (10 mg/kg) (HAL + C + E; n=5). Mice were sacrificed on day 25. Under anesthesia, D-luciferin (Life Technologies) was injected i.p. and the bioluminescence signal was detected by the IVIS imaging system 100 (Xenogen). Liver were disected, fixed in 10% formalin, and paraffin-embedded for further analysis. This animal model was repeated and overall survival was monitor for 80 days for the toxicity test of each treatment.

Microvessel density (MVD) measurement

After antigen retrieval of the liver sections, slides were incubated with anti-CD31 antibody (Thermo Scientific, Waltham, MA) overnight at 4°C and signal was detected by the LSAB + System-HRP kits (Dako) according to the manufacturer’s instruction. Sections were then counterstained with hematoxylin and dehydrated through a series of ethanol and xylene. Slides were then examined with light microscope (100 ×) and single endothelial cells or clusters of endothelial cells positive for CD31 staining were considered as a vessel. MVD was measured based on the Weidner’s method [21] by two independent investigators who were blinded to the study groups.

Statistical analysis

Data were presented as means ± SD from three independent experiments for the in vitro study and from n=5 for the in vivo study. Data were statistically analyzed with one-way ANOVA. Survival analysis was done by Mantel-Cox (Log-Rank) test. All analysis was performed by Prism 5.0 and data were considered statistically significant at P<0.05.

Results

Anti-proliferative effect of combined treatment with Hypoxia + C + E

The effect of different treatments on cell proliferation was first measured by the MTT cell proliferation assay (Figure 1). Cell proliferation decreased with increasing concentrations of everolimus under all the treatment condition. Without the addition of everolimus, treatment

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Figure 4. Expression of phosphorylated proteins in the Akt/mTOR signaling pathway. MHCC-97L cells were treated with Control: in normoxic condition (20% O2, 5% CO2, and 75% N2) without any drug treatment, Hypoxia: in hypoxic condition (1% O2, 5% CO2, and 94% N2), Hypoxia + C: in hypoxic condition and 6 μM cisplatin, Hypoxia + E: in hypoxic condition and 10 nM everolimus, and Hypoxia + C + E: in hypoxic condition, 6 μM cisplatin and 10 nM everolimus. The expressions of phosphorylated mTOR, p70S6, RPS6, Akt, TSC2, GSK3a and GSK3β were measured by the multiplex bead assay. The expression of GAPDH was used as internal control. Data were presented as means ± SD from three independent experiments and statistical analysis were performed by one-way ANOVA. *P<0.05 versus Control. #P<0.05 of Hypoxia + C + E versus Hypoxia, Hypoxia + C or Hypoxia + E.
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We then determined the apoptotic effect of different treatments with the annexin V assay (Figure 2). Treatment of cells with Hypoxia, Hypoxia + C and Hypoxia + E significantly increased the percentage of apoptotic cells by 3.01, 3.89 and 2.33 folds, respectively, when compared with the Control. For the combined treatment with Hypoxia + C + E, the percentage of apoptotic cells increased by 11.4 folds when compared with the Control. In addition, treatment with Hypoxia + C + E further increased the percentage of apoptotic cells by 3.78 folds when compared with the treatment with Hypoxia, by 2.93 folds when compared with the treatment with Hypoxia + C, and by 4.89 folds when compared with the treatment with Hypoxia + E treatment. This suggested that combined treatment with Hypoxia + C + E was significant in enhancing cellular apoptosis when compared with the single treatment.

Inhibiting effect of the combined treatment with Hypoxia + C + E on tube formation ability of HUVEC cells

Treatment with Hypoxia significantly enhanced the tube formation ability of HUVEC cells by 40.0% when compared with the Control (Figure 3). For the combined treatment with Hypoxia + C + E, the tube formation ability of HUVEC cells decreased by 52.0% when compared with the treatment with Hypoxia, and by 46.7% when compared with the treatment with Hypoxia + C. This suggested that combined treatment with Hypoxia + C + E was significant in reducing tube formation ability of HUVEC cells when compared with the single treatments.

of cells under hypoxic condition and under hypoxic condition with 6 μM cisplatin reduced the cell proliferation by 59.7% and 65.1%, respectively, when compared with that treated under normoxic condition. The IC50 under normoxic condition, hypoxia and hypoxia with cisplatin were 11.2 nM, 11.6 nM and 12.2 nM, respectively. With the addition of 10 nM everolimus, treatment of cell under hypoxic condition with 6 μM cisplatin significantly reduced the cell proliferation by 64.0% and 39.2% when compared with that treated under normoxic condition and hypoxic condition, respectively. This suggested that 10 nM everolimus is the optimized concentration in enhancing the anti-proliferating effect of single treatment.

Figure 5. Western blotting analysis of the expression of HIF-1α and VEGF. MHCC-97L cells were treated with Control: in normoxic condition (20% O2, 5% CO2 and 75% N2) without any drug treatment, Hypoxia: in hypoxic condition (1% O2, 5% CO2, and 94% N2), Hypoxia + C: in hypoxic condition and 6 μM cisplatin, Hypoxia + E: in hypoxic condition and 10 nM everolimus, and Hypoxia + C + E: in hypoxic condition, 6 μM cisplatin and 10 nM everolimus. Upper panel: Representative blots of the expression of HIF-1α and VEGF were shown. The expression level of β-actin was used as the loading control. Lower panel: Bar chart presenting the ratio of HIF-1α vs. β-actin and VEGF vs. β-actin was shown. Data were presented as means ± SD from three independent experiments and statistical analysis were performed by one-way ANOVA. *P<0.05 versus Control. #P<0.05 of Hypoxia + C + E versus Hypoxia or Hypoxia + C.
Inhibiting effect of the combined treatment with Hypoxia + C + E on Akt/mTOR signaling pathway

Phosphoprotein alteration was analyzed using the MILLIPLEX MAP 8-Plex Multi-Pathway Signaling Phosphoprotein Kit (Figure 4). Phosphorylation of different targets/effectors of the Akt/mTOR pathway including p70S6, RPS6, Akt, TSC2, GSK3α and GSK3β was studied and each single treatment caused a significant down-regulation of the Akt/mTOR signaling pathway when compared with the Control. In addition, the combined treatment with Hypoxia + C + E further reduced the levels of these phosphorylated proteins. These results
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Figure 7. Immunohistochemistry of CD-31 expression and the MVD analysis. The MVD measurement in the tumor of each liver section by CD31 expression. Upper panel: Representing images of the CD-31 staining (magnification: 100 ×) of tumor in the liver section from each group were shown. Lower panel: the average MVD measurements were plotted. Data were presented as means ± SD and statistical analysis were performed by one-way ANOVA. *P<0.05 versus Control. #P<0.05 of HAL + C + E versus HAL or HAL + C.

suggested a more potent inhibitory effect on the Akt/mTOR signaling pathway was observed in the combined treatment with Hypoxia + C + E than that of the single treatments.

Combined treatment with Hypoxia + C + E was effective in degrading hypoxia-induced HIF-1α expression

When compared with the Control, induced expression of HIF-1α was found in the treatment groups of Hypoxia by 2.34 folds and Hypoxia + C by 88.0% (Figure 5). Combined treatment with Hypoxia + C + E was found to reduce HIF-1α expression significantly by 2.66 folds when compared with the treatment with Hypoxia, and by 2.14 folds when compared with the treatment with Hypoxia + C. This change in expression pattern was also found in VEGF expression. When compared with the Control, induced expression of VEGF was found in the treatment groups of Hypoxia by 16.2% and Hypoxia + C by 19.5%. Combined treatment with Hypoxia + C + E was found to reduce
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HIF-1α expression significantly by 29.5% when compared with the treatment with Hypoxia, and by 13.9% when compared with the treatment with Hypoxia + C. This suggested that combined treatment with Hypoxia + C + E was effective in degrading HIF-1α, which in turn reduced VEGF expression.

Anti-tumor effect of the combined treatment with HAL + C + E

An in vivo orthotopic mouse model was used to demonstrate the anti-tumor effect of different treatments. Measurement of bioluminescence signal was done on day 6, day 10, day 15 and day 25 after orthotopic tumor implantation (Figure 6A-C). Longitudinal monitoring of tumor growth showed that significant inhibition of tumor growth and progression occurred in all the treatment groups starting on Day 15. In addition, a more pronounced anti-tumor effect was shown in the group with combined treatment with HAL + C + E when compared with the treatment group with HAL and HAL + C. The survival analysis was also performed to study the toxicity effect of different treatments (Figure 6D). Combined treatment with HAL + C + E significantly enhanced the overall survival from 40 days in the control group to 67.5 days in the HAL + C + E group.

To examine the effect of different treatments on microvascular density (MVD) in tumors, blood vessels expressing CD31 were counted (Figure 7). Treatment with HAL and HAL + C significantly enhanced angiogenesis with an increase in MVD by 2.63 folds and 2.78 folds, respectively, when compared with the Control group. For the group with combined treatment with HAL + C + E, MVD reduced by 8.27 folds when compared with the HAL group, and 8.73 folds when compared with the HAL + C group. These results suggested that combined treatment with HAL + C + E was more effective in inhibiting tumor growth and progression when compared with single treatments. In addition, treatment with HAL + C + E was effective in inhibiting hypoxia induced microvascular growth which makes the combined treatment a more potent treatment in reducing tumor size in this in vivo study.

Discussion

TACE is one of the effective treatments in advanced HCC with the simultaneous actions of blocking blood supply and administration of a high dose of chemotherapeutic agent to the tumor. However, one of the side effects of TACE is the introduction of hypoxic condition, which in turn activates hypoxia-induced survival pathways and enhances VEGF-induced neovascularization by stabilizing HIF-1α expression. A clinical study demonstrated that increases in neovascularization factors, VEGF and basic fibroblast growth factor (b-FGF), were found in non-responders of TACE treatment and VEGF was even singled out as an independent predictor of survival [22]. Another study on TACE also demonstrated that the residual surviving cancerous tissue in HCC after TACE was highly vascularized and TACE increased VEGF expression in the residual surviving cancerous tissue [23]. In addition, enhanced invasion of tumor and distant metastasis after hepatic artery ligation in orthotopic HCC mouse models were reported [24, 25]. These studies suggested that neovascularization is one of the most crucial obstacles affecting the response of TACE in HCC patients.

Everolimus is a potent mTOR inhibitor. Previous studies have provided evidence that the Akt/mTOR pathway is not only important for the survival of tumor cells but also for angiogenesis of tumor under hypoxia [26]. Studies demonstrated that mTOR is a positive regulator of HIF-1 dependent gene transcription in cells exposed to hypoxia [27]. HIF1-α plays an important role in the up-regulation of certain hypoxia-related proteins involved in angiogenesis, such as VEGF, and in the inhibition of apoptosis [28]. HIF1-α has also been reported to interact with Akt to mediate hypoxia-induced chemoresistance of cisplatin [29]. Therefore, blocking the Akt/mTOR/HIF-1 pathway by everolimus was expected to enhance the TACE efficacy in this study.

Our previous study showed that single treatment with everolimus inhibited HCC tumor growth and enhanced the sensitivity of HCC tumor cells to cisplatin-induced cytotoxicity [30]. In this study, we demonstrated that everolimus significantly enhanced the therapeutic efficacy of TACE using the in vitro and in vivo orthotopic mouse model. We showed that everolimus in combined with TACE significantly suppressed cell proliferation and enhanced cellular apoptosis in MHCC97L cells when compared with single TACE treatment or hypoxia treatment. In addition, the combined treatment was effective in inhibiting tube formation of
HUVEC cells. This is supported by the inactivation of the Akt/mTOR signaling pathway with the decrease in phosphorylated p70S6, RPS6, Akt, TSC2, GSK3α and GSK3β expressions and the reduction in HIF-1α and VEGF expression.

In conclusion, by blocking the Akt/mTOR signaling pathway, everolimus enhanced the therapeutic efficacy of TACE. Based on this preclinical study, the potential of combining everolimus with TACE was guaranteed which suggested the use of the combination therapy in the clinical treatment of advanced HCC patients.

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Disclosure of conflict of interest

None.

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