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Novel therapeutic strategy targeting the Hedgehog signalling and mTOR pathways in biliary tract cancer

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Background: Activation of the PI3K/mTOR and Hedgehog (Hh) signalling pathways occurs frequently in biliary tract cancer (BTC). Crosstalk between these pathways occurs in other gastrointestinal cancers. The respective signalling inhibitors rapamycin and vismodegib may inhibit BTC synergistically and suppress cancer stem cells (CSCs).

Methods: Gene expression profiling for *p70S6k* and *Gli1* was performed with BTC cell lines. Tumour and pathway inhibitory effects of rapamycin and vismodegib were investigated in BTC preclinical models and CSCs.

Results: Rapamycin and vismodegib synergistically reduced BTC cell viability and proliferation. This drug combination arrested BTC Mz-ChA-1 cells in the G1 phase but had no significant effect on the cell cycle of BTC Sk-ChA-1 cells. Combined treatment inhibited the proliferation of CSCs and ALDH-positive cells. *Nanog* and *Oct-4* expression in CSCs was decreased by the combination treatment. Western blotting results showed the p-p70S6K, p-Gli1, p-mTOR, and p-AKT protein expression were inhibited by the combination treatment in BTC cells. In an Mz-ChA-1 xenograft model, combination treatment resulted in 80% inhibition of tumour growth and prolonged tumour doubling time. In 4 of 10 human BTC specimens, tumour p-p70S6K and Gli1 protein expression levels were decreased with the combination treatment.

Conclusions: Targeted inhibition of the PI3K/mTOR and Hhpathways indicates a new avenue for BTC treatment with combination therapy.

Biliary tract cancers (BTCs), which include cancer of the biliary duct, cholangiocarcinoma (CCA), and gallbladder cancer, are aggressive and have distinct epidemiologic patterns. A high incidence (85/100 000) of BTC has been reported for northeast Thailand, where CCA represents approximately 85% of primary liver cancers (Poomphakwaen *et al*, 2009). The incidence of intrahepatic CCA is rising rapidly particularly in the western world, including the United States, where an estimated 10 650 people will be diagnosed with BTC in 2014, with the majority presenting at an advanced and inoperable disease stage (Siegel *et al*, 2014). In Latin America, particularly in Chile, and in South Asia

and East Asia, gallbladder cancer is more common. Systemic treatment for BTC is commonly gencitabine plus cisplatin, but despite treatment, the median overall survival rate is <1 year. Novel therapeutics based on the underlying molecular profile of these tumors will be key to changing the treatment paradigm of BTCs.

Mammalian target of rapamycin (mTOR) is a 289-kDa serine/ threonine protein kinase of the PI3K/AKT/mTOR signalling pathway that regulates cell proliferation, survival, and angiogenesis. mTOR expression is common in gallbladder cancer and CCA and may have prognostic significance (Herberger *et al*, 2007; Leal *et al*, 2013;

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Moolthiya *et al*, 2014; Simbolo *et al*, 2014). Inhibition of mTOR has demonstrated anti-tumour effects in preclinical models of gallbladder cancer and CCA (Wu *et al*, 2007; Herberger *et al*, 2009; Zong *et al*, 2014). Furthermore, a clinical trial demonstrated a 5.1% objective response rate in BTC patients treated with the mTOR inhibitor everolimus (Buzzoni *et al*, 2014).

In contrast, the Hh pathway plays a fundamental role in stem cell maintenance, cell differentiation, tissue polarity, and cell proliferation. Dysregulation of the Hh signalling pathway has been associated with the presence of cancer stem cells (CSCs) and implicated in the initiation of pancreatic and other cancers (Rodova et al, 2012). Constitutive activation of the Hh pathway leading to tumourigenesis is seen in various cancers, including basal cell carcinoma and medulloblastoma. Cholangiocellular injury activates Hh ligands, which promotes dysfunctional repair and tumourigenesis. Inhibition of the Hh pathway attenuates the carcinogenesis of CCA cells and enhances necrosis. Expression analysis of human CCA samples has indicated Hh activation in almost 50% of cases (El Khatib et al, 2013). In these cases, noncanonical Hh pathway activation appears to play a key role and can be inhibited by cyclopamine (Kim et al, 2012; Razumilava et al, 2014; Riedlinger et al, 2014).

Our group recently reported crosstalk between the mTOR/ S6K1and Hh signalling pathways in oesophageal adenocarcinoma (Wang *et al*, 2012). mTOR inhibition is already being investigated in the clinical setting in BTCs and modest clinical efficacy has been observed (Buzzoni *et al*, 2014). We hypothesised that noncanonical Hh pathway activation via mTOR plays an oncogenic role in BTC and that simultaneous inhibition of these two pathways in BTC will provide incremental clinical benefit to patients.

MATERIALS AND METHODS

Cell lines and cell culture. Human BTC Mz-ChA-1and Sk-ChA-1cells were obtained from Dr. Reddy's lab (The University of Texas MD Anderson Cancer Center). Human BTC M214, M139, M156, M213, M055, M213LOH, and M213L5H cells were kindly provided by Dr. Lee's lab (The University of Texas Medical Branch). Cells were cultured in 5% CO₂ at 37 °C in DMEM containing 10% foetal calf serum, 100 units ml⁻¹ penicillin, and 100 mg ml⁻¹ streptomycin. Cancer stem cell culture medium consisting of DMEM/F12, glutaMAX, B27 supplement, epidermal growth factor and basic fibroblast growth factor were purchased from Life Technologies (Grand Island, NY, USA).

Reagents and antibodies. Rapamycin was purchased from LC laboratories (Woburn, MA, USA) and vismodegib was obtained from Genentech (South San Francisco, CA, USA). Antibodies recognising p70S6K, Phospho-p70 S6 Kinase (Thr389), mTOR, phospho-mTOR (Ser2448), AKT, and phospho-Akt (Ser473) were obtained from Cell Signaling Technology (Danvers, MA, USA) Antibodies against Gli1and phospho-Gli1(Ser84) were a kind gift of Dr. Mien-Chie Hung's lab (The University of Texas MD Anderson Cancer Center). Monoclonal anti- β -actin antibody was purchased from Sigma-Aldrich (St. Louis, MO, USA).

Cell viability assay. Biliary tract cancer Mz-ChA-1 and Sk-ChA-1 cells were seeded into white 96-well plates at 5×10^3 cells per well. After incubation overnight, the cells were treated with rapamycin, vismodegib, or both at serial concentrations from 0.25 to $50 \,\mu$ mol l⁻¹ for 72 h. The CellTiter-Glo luminescent cell viability assay kit (Promega, Madison, WI, USA) was used to determine the number of viable cells in culture. Following treatment, cellular proliferation was measured according to the manufacturer's instructions. All treatments were performed in triplicate and in three independent experiments.

Colony-formation assay. Colony-formation assay was performed as described previously. In short, 1×10^3 viable cells were subcultured in 6-well plates in triplicate. Cells were cultured overnight and then treated with $1 \mu \text{moll}^{-1}$ rapamycin, $10 \mu \text{moll}^{-1}$ vismodegib, or both for 72 h. Culture medium was aspirated, the cells were washed, and fresh complete medium was added. Cells were incubated for an additional 14 days. Thereafter, the colonies were fixed with 6.0% glutaraldehyde and stained with 0.5% crystal violet. Clonogenic cells were defined as those able to form a colony consisting of at least 50 cells.

Sphere-formation assay. To evaluate the formation of primary spheres, Mz-ChA-1 and Sk-ChA-1 cells were transferred to ultralow-adherence six-well plates (Corning, Inc., Corning, NY, USA) at a density of 5000 per well in 4 ml of CSC medium consisting of DMEM/F12 plus glutaMAX, B27 supplement, 20 ng ml⁻¹ epidermal growth factor and 20 ng l⁻¹ basic fibroblast growth factor. Tumour spheres were allowed to grow for 7 days. To establish spheres, cells were individualised by enzymatic dissociation, strained, and then plated at a density of 5×10^2 or 1×10^3 cells per well in 24-well plates (3 wells per treatment). Cells were treated with rapamycin1 μ moll⁻¹, vismodegib10 μ moll⁻¹, or both for 7 days, after which the culture medium was changed to fresh CSC medium. The resultant spheres were photographed after 14 days of culture.

Cell cycle assay. Mz-ChA-1 and Sk-ChA-1 cells were seeded in 25-cm² tissue culture flasks and then incubated with $1 \mu \text{moll}^{-1}$ rapamycin, $10 \mu \text{moll}^{-1}$ vismodegib, or both agents. After treatment for 48 h, cells (2 × 10⁶) were harvested by trypsinisation. Then, cells were fixed in 70% ethanol and then incubated with RNase A (100 μ g ml⁻¹) and propidium iodide (100 μ g ml⁻¹) for 30 min. The data from 10 000 cells for each sample were collected and assessed using a FACS scan (Becton Dickinson, Franklin Lakes, NJ, USA).

Quantitative reverse-transcriptase polymerase chain reaction (qRT-PCR). Total RNA was extracted from cells using the RNeasy mini kit (Qiagen, Valencia, CA, USA) according to the manufacturer's instructions. cDNA was synthesised from 1 μ g of total RNA using an iScript cDNA synthesis kit (Life Technologies Corp., Hercules, CA, USA). Real-time RT-PCR analysis was performed using a QuantiFast SyBR green PCR kit (Qiagen). The primer sequences were synthesised by Sigma (Supplementary Information 1). Gene expression levels were normalised against the average cycle threshold (Ct) values for the internal control gene *GAPDH*. Cycle threshold values were extracted using SDS version 2.3 software (Applied Biosystems, Foster City, CA, USA). Data analysis was performed using the $\Delta\Delta$ Ct method.

Flow cytometric analysis of aldehyde dehydrogenase (ALDH) activity. Mz-ChA-1 and Sk-ChA-1 cells (2.5×10^5) were treated with $1 \,\mu$ moll⁻¹ rapamycin, $10 \,\mu$ moll⁻¹ vismodegib, or both for 72 h. The cells were then harvested via trypsinisation and stained for 30 min at 37 °C using an Aldefluor kit (Stem Cell Technologies, Durham, NC, USA) following the manufacturer's instructions. Cells treated with diethylaminobenzaldehyde, a specific ALDH inhibitor, were used as a control. The ALDH-positive control population was detected using the FACS scan. ALDH-positive cells were identified with a dot plot.

Western blotting assay. Biliary tract cancer cells were treated with vehicle, $1 \mu moll^{-1}$ rapamycin, $10 \mu moll^{-1}$ vismodegib, or the combination of the two drugs for 48 h. Total cellular lysates were obtained by lysing cells in a buffer containing RIPA buffer (50 mm Tris-HCl, pH 7.5, 150 mm NaCl, 1% v/v Nonidet P-40, 0.5% v/v sodium deoxycholate, and 0.1% SDS) and a mixture of protease and phosphatase inhibitors. The protein concentration was measured using a bicinchoninic acid protein assay kit

(Pierce, Rockford, IL, USA). Electrophoresis was carried out through a 4%–20% Ready Gel Tris-HCl precast polyacrylamide gel (Life Science Technology, Hercules, CA, USA). Gels were transferred to Hybond–C extra nitrocellulose membrane (Amersham Bioscience, Freiburg, Germany). Proteins were detected with antibodies against Gli1, p-Gli1, mTOR, p-mTOR, p70S6K, p-p70S6K, AKT, and p-AKT(Ser⁴⁷³) Cell Signaling Technology, Danvers, MA, USA). Signals were detected by incubating the membrane with Pierce ECL western blotting substrate (Thermo Scientific, Rockford, IL, USA) and exposed to X-ray (BioExpress, Kaysville, UT, USA).

Mouse xenograft model. All experiments were performed at The University of Texas MD Anderson Cancer Center and were approved by the Institutional Animal Care and Use Committee. We examined the anti-tumour efficacy of rapamycin, vismodegib, or both in a subcutaneous cancer cell xenograft model. Nude (nu/nu) 6- 8-week-old mice were purchased from Charles River (Wilmington, MA, USA). Five million Mz-ChA-1 cells were subcutaneously injected into each flank in 50:50 PBS:Matrigel (BD Biosciences, Billerica, MA, USA). When tumours reached a mean volume of 100 mm³, animals were matched by size and distributed into four treatment groups consisting of eight mice per group. Mice were treated by oral gavage twice a day with vehicle, 1 mg kg^{-g} rapamycin, 100 mg kg^{-1} vismodegib, or both rapamycin and vismodegib. Tumour volume was calculated with the formula $1/2(L \times W^2)$, where L is length (longest dimension) and W is width (shortest dimension). To calculate tumour growth inhibition, the following formula was used: (TuG_{control}~-~TuG_{test})/TuG_{control}~\times 100%, where tumour growth (TuG) equals the final tumour size minus the pretreatment tumour size for individual treatment groups. 'T_uG_{control}' refers to the growth in mice treated with vehicle only and 'TuGtest' refers to the growth in mice treated with rapamycin, vismodegib, or both rapamycin and vismodegib. Each mouse was killed by CO₂ asphyxiation at the experimental end point or when its tumour was oversized, ulcerated, or necrotic.

Immunohistochemical analysis. Immunohistochemical was performed to examine the p-p70S6K and Gli1 protein expression levels in tumour xenografts and archival tumours from patients with resected gallbladder cancer. Xenograft tumour tissue was covered with OCT embedding compound and stored at -80 °C until ready for sectioning. Frozen sections were mounted on 3-aminopropyltriethoxysilane-coated slides. For xenograft tumour tissue slides, the tissue sections were immersed in pre-cooled acetone (-20 °C) for 10 min and the acetone was allowed to evaporate from the tissue sections. Human formalin-fixed, paraffin-embedded tissue sections were deparaffinised using xylene and ethanol. All sections were heated for epitope retrieval in citrate buffer (pH 6.0). Endogenous peroxidase was blocked by incubation in 3% hydrogen peroxide. Sections were incubated with the primary polyclonal rabbit anti-p-p70S6K antibody at a dilution of 1:400, rabbit anti-Gli1 antibody for 30 min at 37 °C, and then biotin-conjugated secondary goat anti-rabbit antibody (Vector Laboratories, Burlingame, CA, USA) for 30 min. Sections were stained with the avidin-biotin complex technique using reagents as described by the manufacturer (Vectastain kit, Vector Laboratories). Positive labelling was visualised by incubating the slides with horseradish peroxidase-conjugated streptavidin (DAB substrate kit, Vector Laboratories). Sections were counterstained with haematoxylin. Stained slides were dehydrated by sequential steps through a graded series of alcohol washes and xylene (Fisher, Suwanee, GA, USA) and were mounted using cover slips.

Statistical analysis. To determine whether the interaction between rapamycin and vismodegib was synergistic, additive, or antagonistic, we used the combination index (CI) method of Chou and Talalay (Chou and Talalay, 1984). Combination index = 1 indicates an additive effect, CI < 1 a synergistic effect, and CI > 1 an antagonistic effect. Results are expressed as the mean \pm s.d. of the indicated number of independent experiments. Student's *t*-test was calculated to compare the mean of each group with that of the control group. *P* values of <0.05 were considered significant. The statistical analysis of data in this study was performed using Student's *t*-test.

RESULTS

Hedgehog and mTOR pathway gene and protein expression in BTC cell lines. We first examined the mutation profile of nine known BTC cell lines for commonly mutated human oncogenes. We used a mass spectroscopy-based profiling for 159 point ('hot spot') mutations in 33 genes commonly involved in solid tumours. Our results indicated there were no significant hotspot mutations in these cell lines (Supplementary Information 2). We then conducted quantitative real-time RT-PCR assays to examine the relative expression levels of p70S6K and Gli1 in all 9 BTC cell lines, as indicated by the fold change relative to the gene with the lowest expression level (Figure 1). The real-time RT-PCR results showed a relatively high p70S6K and low Gli1 gene expression in Mz-ChA-1 cells and relatively low p70S6K and high Gli1 gene expression in Sk-ChA-1 cells. On western blotting, downregulation of p-p70S6K and p-Gli1 with vismodegib and rapamycin was noted in Mz-ChA-1 and Sk-ChA-1 cell lines (Supplementary Information 3). Therefore, we chose these two cell lines for further study. Prior phase I studies with the Hh antagonist, vismodegib, have successfully examined Gli1 expression using RT-PCR to measure the pharmacodynamic modulation of the Hh pathway in the

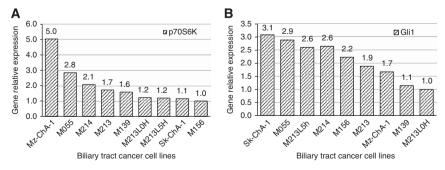


Figure 1. Relative expression levels of *p70S6K* and *Gli1* in BTC cell lines. Real-time RT-PCR analysis of *p70S6k* and *Gli1* relative expression in BTC cell lines. Values represent differences in normalised expression levels compared with the lowest gene expression (normalised against GAPDH mRNA levels). (A) *p70S6K* expression differences in BTC cell lines compared with M156 cell line. (B) *Gli1* expression differences in BTC cell lines compared with M213LOH cell line.

clinical setting (Von Hoff *et al*, 2009; Wong *et al*, 2011). As noted earlier, the crosstalk between the mTOR/S6K1 and Hh signalling pathway is through S6K1-mediated regulation of Gli1 in oesophageal cancer, thus p70 S6K1 was chosen for measurement of mTOR activity (Wang *et al*, 2012). We therefore used p70S6K and Gli1 as the surrogate biomarkers of mTOR and Hh pathway activation, respectively.

Effects of rapamycin, vismodegib, and both on BTC cell viability and proliferation. To explore the effects of rapamycin, vismo-

degib, and both on BTC cell proliferation, we used the CellTiter-Glo (Promega) luminescent cell viability assay to examine whether the combined treatment enhanced the inhibition of cell proliferation affected by either agent alone. Mz-ChA-1 and Sk-ChA-1 cells were treated at serial concentrations for 72 h. Our results showed that rapamycin and vismodegib inhibited proliferation in both cell lines in a concentration-dependent manner and that Mz-ChA-1 cells were more sensitive than Sk-ChA-1 cells to both drugs (Figure 2A and B). The results also suggested that combination therapy reduced cell viability more than either agent alone did.

To determine whether the combined treatment had a synergistic effect on cell viability, the CI values were calculated according to the method of Chou and Talalay (Table 1). The combination of vismodegib and rapamycin resulted in synergistic interaction in Mz-ChA-1 cells at IC₅₀, IC₇₅, and IC₉₀, and the CI values were

 0.40 ± 0.12 , 0.39 ± 0.16 , and 0.40 ± 0.16 , respectively. A modest additive effect also was observed for Sk-ChA-1 cells at IC₅₀, IC₇₅ and IC₉₀, and the CI values were 0.96 ± 0.18 , 1.08 ± 0.25 , and 0.78 ± 0.17 , respectively. In the colony-formation assays, compared with the control or vismodegib, rapamycin individually, the combination treatment significantly inhibited Mz-ChA-1 cell colony formation (*P*<0.01). For Sk-ChA-1 cell, the combination treatment also inhibited the colony formation (*P*<0.05) (Figure 2C and D).

To evaluate whether vismodegib, rapamycin, or their combination can inhibit cell proliferation via regulation of cell-cycle arrest, cell-cycle analysis was carried out by flow cytometry (Figure 3). Cells were treated with rapamycin, vismodegib, or both for 48 h. Vismodegib alone increased the accumulation of Sk-ChA-1cells in the S phase but had no notable effect on Mz-ChA-1 cells. Rapamycin alone arrested Mz-ChA-1 cells in the G1 phase and decreased the cell population in the S phase but had no notable effect on the Sk-ChA-1 cell population. For both cell lines, the combination of vismodegib and rapamycin arrested cells in the G1 phase and decreased the S phase cell population, although vismodegib alone increased the transition from G1 to S phase in the SK-ChA-1 cell population.

Effects of vismodegib, rapamycin, or both on BTC stem cell populations. Cancer stem cells have the exclusive ability to regenerate tumors. The capacity to generate a tumour sphere in nonadherent culture condition is a CSC characteristic. Mz-ChA-1

$IC_{50} \pm s.d.$					
Cell line	Rapamycin (µм)	Vismodegib (µм)	Combination treatment		
			Rapamycin (µм)	Vismodegib (µм)	Combination index
Mz-ChA-1	3.31±0.26	54.97 ± 3.45	0.82 ± 0.18	8.19 ± 1.02	0.40 ± 0.12
Sk-ChA-1	3.53±0.31	74.54 ± 2.58	2.28 ± 0.21	22.83 ± 1.35	0.96±0.18

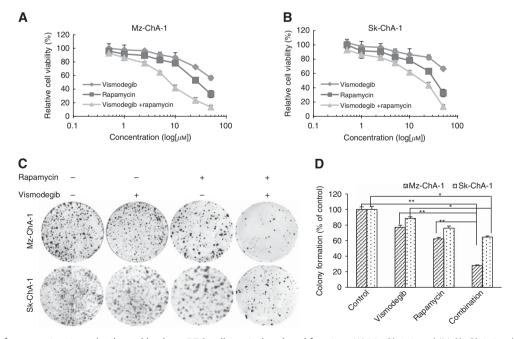


Figure 2. Effect of rapamycin, vismodegib, and both on BTC cell survival and proliferation. (A) Mz-ChA-1 and (B) Sk-ChA-1 cells were treated for 72 h at serial concentrations ($0.25-50 \mu moll^{-1}$). Rapamycin, vismodegib, and both caused a dose-dependent decrease in BTC cell proliferation, as measured by the CellTiter-Glo assay. (C) Clonogenic assay. Cells were treated with DMSO (control), $1 \mu moll^{-1}$ rapamycin, $10 \mu moll^{-1}$ vismodegib, or both for72 h, cells stained with 0.5% crystal violet. (D) Colony numbers as determined by clonogenic assay. The number of colonies with ≥ 50 cells was counted. Values are mean ± s.d. Results are representative of three independent experiments. *P<0.05, **P<0.01.

cells and Sk-ChA-1 cells were cultured in 24-well ultralowattachment plates and treated with vehicle (DMSO; control), $1 \,\mu$ moll⁻¹ rapamycin,10 μ moll⁻¹ vismodegib, or both rapamycin and vismodegib for 7 days. The combination of the two drugs dramatically reduced the total number and size of the tumour spheres compared with the control group, especially for Mz-ChA-1 cells (Figure 4A and B).

In order to further evaluate the effects of vismodegib, rapamycin, and both agents on CSCs, we used real-time RT-PCR to examine CSC-related gene expression in the tumour spheres.

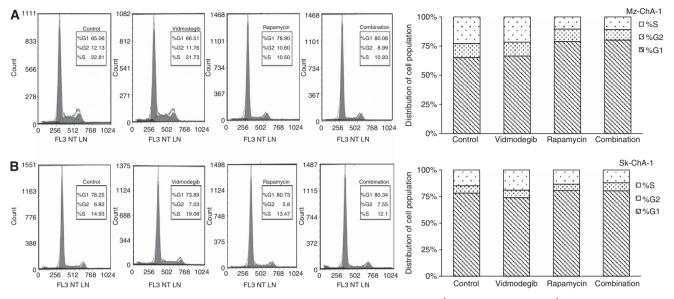


Figure 3. Cell-cycle analysis. (A) Mz-ChA-1 and (B) Sk-ChA-1 cells were treated with $1 \mu moll^{-1}$ rapamycin, $10 \mu moll^{-1}$ vismodegib, or both for 72 h. Cell-cycle distribution was analyzed by flow cytometry.

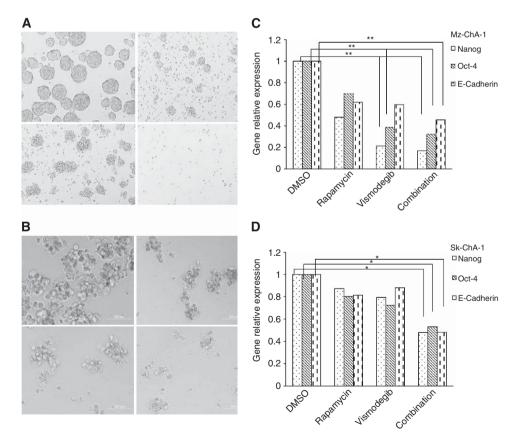


Figure 4. Effect of rapamycin, vismodegib, or both on stem cell sphere formation. Effects of vismodegib, rapamycin or both on tumoursphere formation and *Nanog*, *Oct-4*, and *E-Cadherin* gene expression. Mz-ChA-1 (**A**) and Sk-ChA-1 (**B**) tumour spheres. Third passage single cells were cultured for 72 h, and then treated with vehicle, rapamycin ($1 \mu moll^{-1}$), vismodegib ($10 \mu moll^{-1}$), or both for 7 days. Tumourspheres were imaged by Olympus microscope. Compared with control group, the combination treatment inhibited *Nanog*, *Oct-4*, and *E-Cadherin* expression in Mz-ChA-1 (**C**) (***P*<0.01) and Sk-ChA-1 (**D**) (**P*<0.05).

The results showed that combined treatment significantly decreased the Nanog, Oct-4, and E-cadherin gene expression in both Mz-ChA-1 and Sk-ChA-1 CSCs compared with the DMSO (control) group (Figure 4C and D).

Aldehyde dehydrogenase was also investigated as a possible marker for identifying CSCs, including BTC stem cells. We performed the ALDEFLUOR assay to explore the effect of the treatment agents on the Mz-ChA-1 and Sk-ChA-1 CSC populations (Figure 5A and D). The combined treatment significantly reduced the ALDH-positive population in Mz-ChA-1 cells but not in Sk-ChA-1 cells (Figure 5C and D).

Effects of vismodegib, rapamycin, and both on mTOR and Hh pathway signalling. We investigated the activity of rapamycin, vismodegib, and both on mTOR and Hh signalling in BTC cells by using western blots. The Mz-ChA-1 and Sk-ChA-1 cells were treated with rapamycin, vismodegib, or both for 48 h. We examined the expression levels of p70S6K, p-p70S6K, AKT, p-AKT(Ser⁴⁷³), Gli1, p-Gli1, mTOR, and p-mTOR. The initial ratios of pS6K/S6 in Mz-ChA-1 and Sk-ChA-1 cells were $51.62 \pm 9.07\%$ and $13.22 \pm 2.03\%$, respectively. The results showed that, for both cell lines, the expression of the phosphorylated proteins was significantly decreased in the combined treatment group (Figure 6). The western blotting results were repeated three times and with reproducible results.

Anti-tumour activity of the combination of rapamycin and vismodegib in a xenograft model. We next evaluated the antitumour activity of vismodegib, rapamycin, or both *in vivo* with a xenograft mouse model. Single-cell suspensions of 5×10^6 Mz-ChA-1 cells were subcutaneously injected into the right flank of 32 athymic nude mice. Once tumours grew to approximately 100 mm^3 , the mice were randomly allocated into four treatment arms (vehicle only, rapamycin, vismodegib, or both rapamycin and vismodegib) and treated twice daily through oral gavage. Compared with the control group, at day 27, tumour xenograft growth was $39.42 \pm 12.33\%$, $51.03 \pm 5.71\%$, and $80.39 \pm 11.18\%$ (P < 0.01) lower in the rapamycin, vismodegib, and combination groups, respectively (Figure 7A). The xenograft tumour doubling time was 7.11 ± 0.88 , 9.31 ± 1.29 , 12.40 ± 2.01 , and 20.04 ± 5.48 days in the control, rapamycin, vismodegib, and combined treatment groups. Nude mice were killed on day 27 because of the tumour size.

The mice tolerated the treatments without overt signs of toxicity. Body weight did not differ significantly between treatment groups (Figure 7B), and no adverse effects such as hunched posture, ruffled fur, and hypothermia were observed. Immunohistochemical assay results for the xenograft tumour tissues showed that the combined treatment significantly decreased p-p7086K and Gli1 protein expression levels as compared with the control (vehicle only) group *in vivo* (Figure 7C).

Immunohistochemical analysis of human samples of gallbladder cancer. In order to identify potential predictive biomarkers for vismodegib and mTOR inhibitors in human specimens, we investigated the protein expression levels of Gli1 and p-p70S6K in cases of resected gallbladder cancer. Our immunohistochemical results revealed a relatively high p-p70S6K protein level and low Gli1 protein expression level in 4 of 10 cases examined (Figure 8). This immunohistochemical pattern was similar to those we found in Mz-ChA-1 cell lines.

DISCUSSION

The combination of rapamycin and vismodegib significantly inhibited BTC cell viability and proliferation in our study; this effect was confirmed with our *in vivo* study. The protein expression levels of p-p7086K, p-mTOR, p-Gli1, and p-AKT in Mz-ChA-1 and Sk-ChA-1 cells were decreased by the combination regimen.

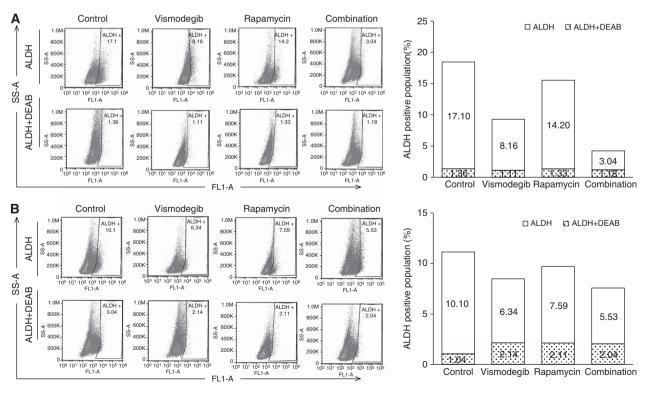


Figure 5. ALDH activity measured by FACS analysis. Aldefluor analysis of Mz-ChA-1 cells (**A**) and Sk-ChA-1 cells (**B**) treated with vehicle, rapamycin (1 μ moll⁻¹), vismodegib (10 μ moll⁻¹), or both for 72 h. ALDH⁺ Mz-ChA-1 cells were significantly decreased in the combined treatment group (compared with control and vismodegib, P<0.01 and P<0.05, respectively).

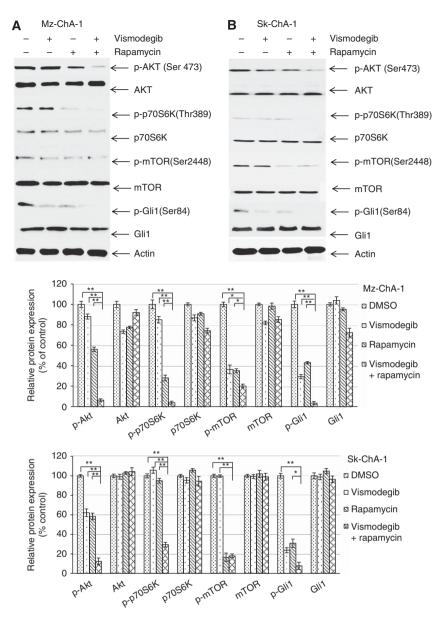


Figure 6. Effect of rapamycin, vismodegib, or both on protein expression. Mz-ChA-1 cells (A) and Sk-ChA-1 cells (B) were treated with vehicle, rapamycin (1 μ moll⁻¹), vismodegib (10 μ moll⁻¹), or both for 24 h. Actin was used as a loading control (*P<0.05, **P<0.01).

Decreased expression of p-p70S6K and Gli1 was noted in the BTC xenografts treated with this combination. High p-p70S6K expression along with low Gli1 expression was observed in Mz-ChA-1 cell lines, which were sensitive to the combination regimen. This expression pattern was also noted in 40% of the human BTC cases we examined.

Biliary tract cancer represents a diverse and genetically heterogeneous group of cancers, which makes the implementation of molecular therapy challenging. CCA and gallbladder cancer are often grouped together as BTCs in clinical trials despite their divergent clinical presentation and genetic background. Furthermore, there are very few BTC cell lines and preclinical *in vivo* models. All of these factors make the development of novel therapies in BTC challenging. Recently, there have been promising genomic and transcriptional profiling studies in intrahepatic CCA that suggest that this disease has distinct molecular subsets, some of which may be amenable to targeted therapy (Andersen *et al*, 2012; Borger *et al*, 2012; Sia *et al*, 2013; Voss *et al*, 2013).

The activation of Hh signalling is common in BTC and has prognostic implications. Tang *et al* (2013) recently demonstrated

that Gli1 expression was associated with a poor prognosis in CCA and demonstrated that inhibition of Hh signalling with cyclopamine led to apoptosis in CCA cells (Tang et al, 2013). Canonical activation of Hh signalling involves Hh ligand binding to its receptor, PTCH, which results in the activation of Smo and is followed by activation of Gli transcription factors. A critical role of Smo-independent regulation of Gli activity by several other signalling pathways, including the PI3K/AKT and RAS/RAF/ MEK/ERK pathways, has been suggested (Lauth and Toftgard, 2007; Jenkins, 2009). Canonical Hh signalling is attenuated in CCA, and non-canonical Hh signalling contributes to disease progression, thereby supporting a role for Hh and other targeted inhibitors in this disease (Razumilava et al, 2014). In our study, combined mTOR- and Hh-directed therapy had the optimal antitumour effect in BTC cell lines, particularly Mz-ChA-1. Although combined therapy was superior, inhibition of p70S6K also occurred with vismodegib and p-Gli1 with rapamycin, which supports the idea of bi-directional cross talk between the pathways. Our group described the proposed mechanism of this cross talk in a previous report: activated mTOR pathway promotes Gli1

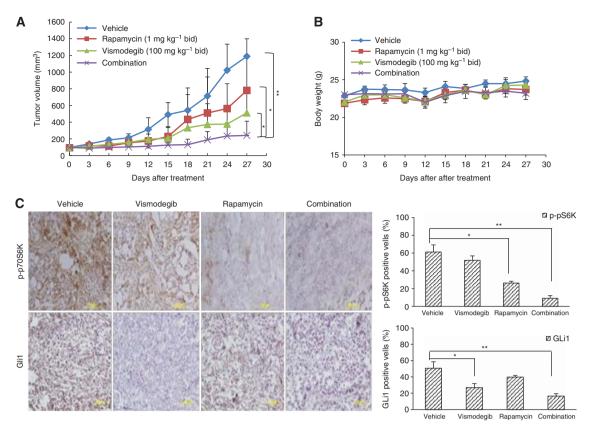


Figure 7. Effect of rapamycin, vismodegib, or both on Mz-ChA-1 cell xenograft tumors. (A) Effects on xenograft growth. Mice treated with vehicle only, rapamycin (1 mg kg⁻¹, b.i.d.), vismodegib (100 mg kg⁻¹, b.i.d.), or both when tumour volume reached 100 mm³. Values are expressed as mean \pm s.d. (n = 8 mice per treatment group). (B) Nude mouse weight. The mice were weighed every 3 days and there was no significant change in nude mice weight. Values are expressed as mean \pm s.d. (C) Immunohistochemical analysis of xenograft tumour tissue. Tumour tissues were subjected to immunostaining for p-p70S6K and Gli1 protein expression. Quantified p-p70S6K and Gli1 protein expression levels in a representative xenograft tumour from each treatment group are shown at right. *P < 0.05, **P < 0.01.

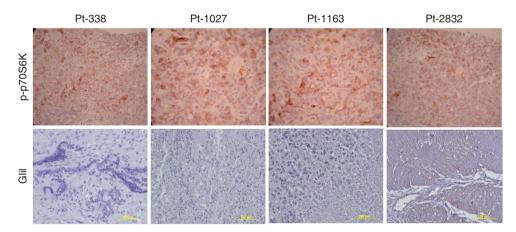


Figure 8. Immunohistochemical analysis of p-p70S6K and Gli1 protein expression. Ten gallbladder cancer patient tumours were examined and four patient tumours with high p-p70 S6K and low Gli1 protein expression.

transcriptional activity and oncogenic function through p70S6K1mediated Gli1 phosphorylation at Ser⁸⁴, which releases Gli1 from its endogenous inhibitor, SuFu (Wang *et al*, 2012).

For studies of the combined inhibition of Hh signalling and mTOR in BTC cells, it would be useful to identify a potential clinical phenotype for investigational therapy. For our study, we first characterised the BTC cell lines for p70S6K and Gli1 gene expression. We chose the Mz-ChA-1 and Sk-ChA-1 cell lines for further investigation, as these represented relative extremes of

p70S6K and *Gli1* gene expression levels, with Mz-ChA-1 cells having a high *p70S6K* gene expression level and Sk-ChA-1 having a high *Gli1* gene expression level. Furthermore, these cell lines have been widely used in prior BTC preclinical studies. The cell line Mz-ChA-1, which is derived from gallbladder cancer, was significantly inhibited by the combination of vismodegib and rapamycin. The dose of rapamycin and vismodegib used *in vivo*, when converted to human equivalent doses are within clinically acceptable dose ranges (Supplementary Information 4) (LoRusso *et al*, 2011;

Oct-4, and E-cadherin gene expression level. Nanog and Oct-4 are essential for maintaining the pluripotency of embryonic stem cells, whereas aberrant expression of *E-cadherin* has been associated with the development of metastases. Although the prevailing belief is that loss of E-cadherin is a prerequisite for progression to metastatic disease and tumourigenesis, tumours are actually remarkably heterogeneous and E-cadherin loss does not always correlate with invasion. E-cadherin is also highly overexpressed in prostate cancer, inflammatory breast cancer, and early-stage ovarian carcinoma. E-cadherin enhances proliferation and survival by inducing ligand-independent activation of the receptor tyrosine kinase epidermal growth factor receptor (Putzke et al, 2011; David and Rajasekaran, 2012). Loss of E-cadherin regulation and expression levels in stem cells is likely to reflect the different culture conditions for these cell lines (Mohamet et al, 2011). High ALDH activity is a characteristic of CSCs in several malignancies, including BTC (Douville et al, 2009). In our study, ALDH activity was notably decreased in Mz-ChA-1 cells with the combined treatment but not in Sk-ChA-1 cells, which supports a role for combination therapy against CSCs in a subset of BTCs.

In summary, the addition of vismodegib to an mTOR inhibitor may lead to an incremental anti-tumour effect in BTCs. In particular, a high p70S6K/Gli1gene expression ratio may predict responsiveness to the combination in the clinical setting.

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CONFLICT OF INTEREST

The authors declare no conflict of interest.

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