RESEARCH ARTICLE

Increased activation of PI3K/AKT signaling pathway is associated with cholangiocarcinoma metastasis and PI3K/ mTOR inhibition presents a possible therapeutic strategy

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Abstract Phosphatidylinositol 3-kinase (PI3K) signaling plays a critical role in cholangiocarcinoma (CCA), as well as anti-cancer drug resistance and autophagy, the type II program cell death regulation. In this work, we aimed to: (1) determine the expression levels of several key components of PI3K signaling and (2) evaluate whether NVP-BEZ235, a novel dual PI3K/mTOR inhibitor, could inhibit CCA cell growth. Immunohistochemistry for p85 α , p110 α , AKT, p-AKT (T308), mTOR, p-mTOR (S2448), GSK-3 β , p-GSK-3 β (S9), PTEN, and p-PTEN (S380, T382/383) was performed in 30 CCA patients. Western blotting was used to analyze PTEN and p-PTEN expression in the cell lines (KKU-OCA17, KKU-100, KKU-M055, KKU-M139, KKU-M156, KKU-M213, and KKU-M214). The effects of

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Department of Neurosurgery, School of Medicine, Johns Hopkins University, Baltimore, MD, USA NVP-BEZ235 on CCA cells were evaluated using a growth inhibition assay, flow cytometer and migration assay. Increased activation of PI3K/AKT signaling was reproducibly observed in the CCA tissues. The expression of $p85\alpha$. mTOR, and GSK-3 b was significantly correlated with metastasis. Interestingly, PTEN suppression by loss of expression or inactivation by phosphorylation was observed in the majority of patients. Furthermore, NVP-BEZ235 effectivelyinhibited CCA cell growth and migration through reduced AKT and mTOR phosphorylation and significantly induced G1 arrest without apoptosis induction, although increase autophagy response was observed. In conclusion, the constitutive activation of PI3K/AKT pathway in CCA is mainly due to PTEN inactivation by either loss of expression or phosphorylation along with an increased expression in its pathway components heralding a poor prognosis for CCA patients. This work also indicates that inhibition of PI3K and mTOR activity by the inhibitor NVP-BEZ235 has anticancer activity against CCA cells which might be further tested for CCA treatment.

Keywords PI3K/AKT \cdot mTOR \cdot PTEN \cdot NVP-BEZ325 \cdot Cholangiocarcinoma

Abbreviations

CCA	Cholangiocarcinoma
PI3K	Phosphoinositide 3-kinase
mTOR	Mammalian target of rapamycin
MAPK	Mitogen-activated protein kinase
JAK/STAT	Janus kinase/signal transducer and activator
	of transcription

Introduction

Cholangiocarcinoma (CCA) is an aggressive cancer originating from the bile duct epithelium. CCA is a major public health problem in the northeastern part of Thailand where the highest incidence in the world has been reported [1, 2]. Chronic inflammation caused by liver fluke (Opisthorchis viverrini, Ov) infection is documented as the major etiology of CCA development in this area [1, 3, 4]. CCA is recognized as a highly metastatic cancer [5-7]. CCA patients are mostly clinically silent and difficult to diagnose until the disease is advanced and often in the metastatic stage resulting in a poor prognosis. Surgical treatment of early stage disease is the only curative therapy, but unfortunately, the majority of CCA patients present late with advanced stage disease, which is not surgically curable [8]. The addition of current chemotherapy regimens is helpful but does not appreciably improve the long-term survival rate [9]. In order to develop new molecular therapies for CCA, a better understanding of the molecular pathogenesis of CCA is needed. So far, the majority of molecular strategies target anti-apoptotic or growth-stimulating pathways, in order to interfere with cancer-specific mechanisms and reduce adverse effects on normal cells have been extensively explored [10].

We have previously demonstrated that PRKAR1A, a regulatory subunit 1 alpha of protein kinase A type I, is involved in CCA carcinogenesis [11]. Furthermore, inhibition of PRKAR1A had anti-cancer activity against CCA cells and enhanced apoptosis induction with an associated decrease in several protein kinases pathways including MAPK, PI3K/AKT, JAK/STAT, and Wnt/β-catenin signaling. This indicates that PRKAR1A might be a target for CCA treatment [12]. Therefore, these data lead us to focus more on the roles of other protein kinases in CCA.

We have recently identified that multiple protein kinases are activated in both CCA cell lines and CCA tissues. Among those activated protein kinases in CCA, PI3K/AKT signaling pathway is prominently activated [13]. In addition, a previous study has shown upregulation of a downstream of AKT in CCA tissues, mTOR, which was analyzed by a genomic DNA copy number arrays [14]. These results convince us to pay more attention on the roles of PI3K/AKT signaling pathway in CCA development.

PI3K/AKT signaling pathway plays a role as a regulator of many cellular processes including cell division, growth, and programmed cell death. On the other hand, activation of this particular pathway is central to the growth of many human cancers, including colon cancer [15], glioblastoma [16, 17], and CCA [18–24]. PI3K/AKT activation leads to increased resistance to radiation therapy and chemotherapy, and inhibition of this pathway can sensitize CCA cells to these therapies [23, 25, 26]. In addition, Menakongka and Suthiphongchai reported hepatocyte growth factor (HGF)induced invasiveness in CCA cell lines was seen via PI3K/AKT signaling [27]. Conversely, inhibition of this pathway markedly suppressed HGF-stimulated CCA cell invasion.

In the present study, we demonstrated that the aberrant activation of the PI3K signaling pathway in human CCA correlated with tumor metastasis. We also evaluated the in vitro anti-tumorigenic effect of NVP-BEZ235, a dual PI3K/mTOR inhibitor which is an imidazo[4,5-c]quinoline derivative that inhibits activity by binding to the kinase ATP-binding cleft. This compound equally targets wild-type, mutated p110 and mTOR [28] in CCA.

Materials and methods

Human CCA specimens

The 30 paraffin-embedded tissues collected from primary tumors of CCA patients were obtained from the specimen bank of the Liver Fluke and Cholangiocarcinoma Research Center, Faculty of Medicine, Khon Kaen University, Thailand. Informed consent was obtained from each patient prior to surgery, and the Ethics Committee for Human Research at Khon Kaen University approved the research protocols (#HE561035).

Cell lines and cell culture

Human CCA cell lines including KKU-OCA17, KKU-100, KKU-M055, KKU-M139, KKU-M156, KKU-M213, and KKU-M214 were obtained from CCA patient, which were established at Khon Kaen University Liver Fluke and Cholangiocarcinoma Research Center. All cell lines were cultured in Ham's F-12 medium (Gibco/BRL, Grand Island, NY, USA) supplemented with 44 mM NaHCO₃, penicillin (100 units/ml), streptomycin (100 mg/ml), and 10 % fetal bovine serum in a humidified atmosphere containing 5 % CO₂.

Antibodies and inhibitor

Antibodies used for this study were as follows: anti-PI3K (p85 α), PI3K (p110 α), AKT, p-AKT(S473), p-AKT (T308), mTOR, p-mTOR (S2448), GSK-3 β , p-GSK-3 β (S9), PTEN were purchased from Cell Signaling Technology (Cell Signaling Technology, Danvers, MA, USA.), p-PTEN (S380, T382/383) and LC3B were purchased from Abcam (Abcam, Cambridge, UK), and anti- β -actin antibody was purchased from Sigma (Sigma-Aldrich, St. Louis, MO, USA). The dual PI3K/mTOR inhibitor, NVP-BEZ235, was kindly supplied by Novartis Pharma AG (Basel, Switzerland).

The inhibitor was dissolved in dimethylsulfoxide (DMSO; Amresco, Solon, OH, USA) at a stock concentration of 10 mM and stored at -20 °C until used.

Immunohistochemistry staining and scoring

Paraffin-embedded tissues were sectioned for immunohistochemistry (IHC) staining according to standard methods. Briefly, the sections of CCA tissues were deparaffinized and rehydrated through a graded series of aqueous ethanol solutions. Next, microwave antigen retrieval was performed with sodium citrate buffer for 10 min. Then, endogenous peroxidase activity was blocked with 0.3 % hydrogen peroxide in phosphate-buffered saline (PBS) for 30 min, and non-specific binding was blocked by 10 % skim milk in PBS for 1 h. The tissue sections were incubated overnight with the primary antibody against designed target proteins at 4 °C. After that, sections were incubated with peroxidaseconjugated Envision[™] secondary antibody (DAKO, Denmark) for 1 h. After washing, peroxidase-labeled polymer, 0.1 % diaminobenzidinetetrahydrochloride solution was used for the signal development and then counterstained with Mayer's hematoxylin. The stained sections were observed under a light microscope by using the high magnification power ×200 and ×400 (Axioscope A1, Carl Zeiss, Jena, Germany).

The IHC scoring system was used for quantitation of results. Grading of staining depended upon the intensity and frequency of staining in tumor area. Intensity of protein expression was classified into four groups: 0, negative; +1, weak expression; +2, moderate expression; +3, strong expression, respectively. The frequency of staining was divided into four groups: 0, negative; +1, 1–25 %; +2, 26–50 %; +3, >50 %, respectively. Staining scores were calculated by multiplying intensities and frequencies in each case which are classified into two groups: low expression levels; scores are <4 and high expression levels; sores are \geq 4 [6, 29].

Growth inhibition assay

CCA cells (5×10^3) in 100 µl media were seeded in 96-well plates (Costar, Corning, NY, USA) and incubated overnight. Then, the cells were treated with different concentrations of NVP-BEZ235 ranging from 1 to 10^5 nM. After 48 h of incubation, cell numbers were estimated using the sulforhodamine B(SRB) assay (Sigma-Aldrich, St. Louis, MO, USA). The optical densities were read using a microtiter plate reader (Sunrise, TECAN Trading, Switzerland) at 540 nm. The experiments were performed in triplicate. The percentage of growth inhibition (%GI) was calculated by using the equation: %GI=(1-(Nt/Nc)×100, where Nt and Nc represent the absorbance in treated and control cultures, respectively. IC₅₀, the drug concentration causing cells GI of 50 %, was determined by interpolation from dose–response curves, as previously described [30].

Flow cytometry analysis

CCA cells were seeded at 1.5×10^5 cells in six-well plates, cultured overnight, and then cells were treated with 10, 100, and 1,000 nM of NVP-BEZ235 or control medium for 48 h. After that, the cell cycle distribution was detected by staining DNA with propidium iodide (Invitrogen, Paisley, UK), while apoptosis and necrosis were detected by using Annexin-V-FLUOS staining kit (Roche, Penzberg, Germany), respectively. The cell cycle distribution and apoptosis were determined by flow cytometry (BD FACSCantoTM II, BD Biosciences, CA, USA) and analyzed by BDFACSDivaTM software (BD Biosciences, CA, USA).

Western blot analysis

Cells were treated with NVP-BEZ235 at a concentration as 10. 100, and 1,000 nM or control medium for 48 h, respectively. Cell lysates were electrophoresed and transferred to polyvinylidenedifluoride membranes (Millipore, Bedford, USA). Membranes were blocked with 5 % skim milk in Trisbuffered saline (TBS) at room temperature for 1 h and incubated with primary antibody at 4 °C overnight. After rinsing with TBS containing 0.1 % polyoxyethylenesorbitanmonolaurate (Tween-20; TBST), membranes were incubated with horseradish peroxidase-conjugated secondary antibody (Santa Cruz Biotechnology, CA, USA) at room temperature for 1 h. After rinsing with TBST, membranes were exposed to the ECL Prime Western Blotting Detection System (GE Healthcare Bio-Science, UK). The immunoblot and intensity were analyzed by the ImageQuant[™] analysis system (GE Healthcare Bio-Science, UK). Human β -actin was used as a loading control.

Wound healing assay

CCA cells were seeded and grown to confluence in the fibronectin-coated 12-well plates (Costar, Corning, NY, USA). The in vitro wound healing assay was made by scraping the middle of the cell monolayer using a sterile micropipette tip. Floating cells were removed by extensive washing with sterile PBS. After that, cells were cultured with control medium (1 % of DMSO) and 10 nM of NVP-BEZ235 in culture medium. The experiments were performed in duplicate well for both conditions. Photographs were taken using a phase-contrast microscope with a digital camera (Axiovert 40, Carl Zeiss, Germany) at 0, 12, and 24 h after wounding. Degrees of cell spreading between groups were observed.

Statistical analysis

Statistical analyses were performed by SPSS software version 17 (IBM Corporation, NY, USA). The correlation between IHC scores of each protein in CCA tissues was calculated using the Pearson's correlation coefficient. The association of protein expression in CCA tissues and patients' clinico-pathological factors were assessed by Fisher's exact test. Results of cell growth inhibition were presented as mean \pm SEM of a representative experiment. The cell cycle analysis was represented as mean \pm SD, and the significance of differences was addressed by Student's *t* test. A *P* value <0.05 was considered statistically significant.

Results

Altered expression and association of PI3K/AKT signaling pathway molecules with CCA clinicopathological characteristics

A total of 30 CCA tissues were studied, 70 % were male and 30 % were female. The age of patients ranged from 37 to 71 years old (median=58 years old). The patients were at an advanced stage with 50 % presenting with metastasis. In this study, the histological types were classified as papillary type CCA (47 %) and non-papillary type CCA (53 %). The results of immunohistochemical staining revealed no or very weak positive staining of all proteins investigated in normal bile duct epithelia, while the increased expression was observed in pre-cancerous and cancer cells, respectively. The percent expression of p85α, p110α, AKT, p-AKT (T308), mTOR, pmTOR (S2448), GSK-3β, and p-GSK-3β (S9) in CCA tissues is 27, 90, 50, 57, 67, 20, 40, and 27 %, respectively (Fig. 1a, Table 1). Moreover, expression levels of the $p110\alpha$, catalytic subunit of PI3K, have a significant correlation with its regulatory subunit; $p85\alpha$ and its downstream targets as shown in Table 2. Interestingly, the expression of $p85\alpha$, mTOR, and GSK-3ß was significantly correlated with metastasis (P=0.035, P=0.008, and P=0.025, respectively; Tables 3 and 4). Additionally, elevated expression of AKT, p-AKT, and GSK-3ß was significantly correlated with nonpapillary type of CCA (P=0.028, P=0.030, and P=0.007, respectively; Tables 3 and 4). Furthermore, there was a trend towards significance between p-AKT expression and metastasis status of patients (P=0.065). In addition, the expression levels of p110 α also showed a trend associated with nonpapillary (P=0.073), with 67 % of these cases are metastatic CCA. Interestingly, we found loss of PTEN expression in the majority of CCA patients (70 %) as shown in Table 1. Furthermore, we observed that within 89 % of the PTEN positive cases, their PTEN were inactivated by phosphorylation. In addition, Western blot analysis revealed the PTEN inactivation by phosphorylation in all CCA cell lines studied (Fig. 1b). Therefore, loss of PTEN expression as well as inactivation by phosphorylation seems to be the major cause of PI3K/AKT overactivation in this series of CCA (Table 5).

NVP-BEZ235 inhibits CCA cell growth through blocking the PI3K/mTOR activity

We examined further if NVP-BEZ235, a dual PI3K/mTOR inhibitor, could inhibit CCA cell proliferation. Seven CCA cell lines (OCA17, KKU100, M055, M139, M156, M213, and M214) were treated with a range of concentration of the small molecule, and cell proliferation was assessed by SRB assay. The results showed that NVP-BEZ235 effectively suppressed CCA cell growth at one nanomolar and in a dose-dependent manner (Fig. 2a). The IC₅₀ values (mean±SD) of NVP-BEZ235 in seven CCA cell lines, OCA17, KKU100, M055, M139, M156, M213, and M214, were 48 ± 19 , 15 ± 4 , 5 ± 2 , 135 ± 48 , 2 ± 1 , 30 ± 5 , and 45 ± 5 , respectively. Additionally, Western blot analysis revealed that NVP-BEZ235 was able to inhibit the phosphorylation of S473 AKT and S2448 mTOR in two representative cell lines in a dose-dependent response (Fig. 2b), suggesting that NVP-BEZ235 blocks the PI3K/AKT pathway resulting in inhibition of CCA cell proliferation. Moreover, flow cytometry analysis demonstrated that NVP-BEZ235 induced significantly G1 arrest as well as reduced S phase progression of CCA cell lines, for M139 at 1,000 nM and M214 at 100 and 1,000 nM (Fig. 2c).

NVP-BEZ235 induces autophagy without a remarkable effect on apoptosis in CCA cells

To test if NVP-BEZ235 inhibited cell growth, at least in part by inducing apoptosis, we performed flow cytometry using Annexin-V-FLUOS staining. The results indicated that NVP-BEZ235 has no remarkable effect on apoptosis induction (Fig. 3a), although NVP-BEZ235 decreased phosphorylation of AKT and mTOR which function as a key negative regulator for autophagy, the type II programmed cell death. We therefore examined the effect of NVP-BEZ235 on autophagy induction. The results revealed that NVP-BEZ235 induces LC3B (LC3-II) protein, a hallmark of cells undergoing autophagy in a dose- and time-dependent manner in treated CCA cell lines (Fig. 3b).

NVP-BEZ235 inhibited migration of CCA cells

The above IHC data indicated the association of several key components of PI3K pathway with the metastasis status of the CCA patients studied. Therefore, we sought to determine if NVP-BEZ235 could inhibit CCA cell migration, an important step in the metastatic process. We performed an Fig. 1 Expression of several key components of PI3K/AKT signaling pathway. a The expression of $p85\alpha$, $p110\alpha$, AKT, p-AKT (T308), mTOR, p-mTOR (S2448), GSK-3 β , p-GSK-3 β (S9), PTEN, and p-PTEN (S380, T382/383) in normal adjacent bile duct and tumor area by immunohistochemistry staining. Original magnification is ×400. b Western blot analysis of the expression of PTEN and p-PTEN (S380, T382/383) in a panel of CCA cell lines



 Table 1
 Summary of the PI3K/

 AKT signaling pathway components expression in human CCA tissues

Molecules	Total cases	Results								
		Positive	Negative (%)							
		High	Low	Total (%)						
1. p85α	30	3	5	8 (27 %)	22 (73 %)					
2. p110α	30	18	9	27 (90 %)	3 (10 %)					
3. AKT	30	6	9	15 (50 %)	15 (50 %)					
4. p-AKT (T308)	30	5	12	17 (57 %)	13 (43 %)					
5. mTOR	30	8	12	20 (67 %)	10 (33 %)					
6. p-mTOR (S2448)	30	1	5	6 (20 %)	24 (80 %)					
7. GSK3	30	6	6	12 (40 %)	18 (60 %)					
8. p-GSK3 (S9)	30	3	5	8 (27 %)	22 (73 %)					
9. PTEN	30	4	5	9 (30 %)	21 (70 %)					
10. p-PTEN (S380, T382/383)	9	5	3	8 (89 %)	1 (11 %)					

in vitro wound healing assay in conditioned culture medium with or without NVP-BEZ235. The results showed that the migration of the M139 and M214 cells was reduced by treating cells with 10 nM of NVP-BEZ235, a concentration which has no effect on cell proliferation (Fig. 4).

Discussion

Although an increased activation of PI3K/AKT pathway along with PTEN loss [20, 22] has been reported in CCA,

until now there has been no comprehensive study of PI3K, AKT, p-AKT, mTOR, p-mTOR GSK-3 β , p-GSK-3 β together with PTEN and its inactive form, p-PTEN, in human CCA tissues. A previous study revealed that altered expression of p110 α , the catalytic subunit of class IA PI3K, by amplification, overexpression, or somatic mutation has been found in various human tumors [31]. In this current work, we report the overexpression of p110 α in human CCA tissues. Moreover, our results demonstrated the significant positive correlation between expression levels of p110 α with its regulatory subunit, p85 α , and its downstream targets.

 Table 2
 Correlation coefficients between immunohistochemistry scores of the PI3K/AKT signaling pathway components in human CCA tissues

		p110a	AKT	p-AKT (T308)	mTOR	p-mTOR (S2448)	GSK-3β	p-GSK-3β (S9)	PTEN	p-PTEN
Ρ85α	Correlation coefficient	0.535	0.593	0.619	0.619	0.667	0.597	0.564	0.621	0.283
	Р	0.002*	0.001*	0.000*	0.000*	0.000*	0.000*	0.001*	0.000*	0.460
Ρ110α	Correlation coefficient		0.465	0.528	0.464	0.351	0.452	0.282	0.332	0.473
	Р		0.010*	0.003*	0.010*	0.057	0.012*	0.132	0.073	0.199
AKT	Correlation coefficient			0.580	0.681	0.270	0.835	0.681	0.678	0.200
	Р			0.001*	0.000*	0.148	0.000*	0.000*	0.000*	0.606
p-AKT (T308)	Correlation coefficient				0.715	0.323	0.564	0.770	0.728	0.228
	Р				0.000*	0.082	0.001*	0.000*	0.000*	0.555
mTOR	Correlation coefficient					0.478	0.686	0.664	0.828	0.807
	Р					0.008*	0.000*	0.000*	0.000*	0.009*
p-mTOR (S2448)	Correlation coefficient						0.083	0.234	0.439	0.452
	Р						0.665	0.213	0.015*	0.222
GSK-3β	Correlation coefficient							0.637	0.625	0.326
	Р							0.000*	0.000*	0.392
p-GSK-3β (S9)	Correlation coefficient								0.751	0.369
	Р								0.000*	0.328
PTEN	Correlation coefficient									0.754
	Р									0.019*

P=0.05, significant correlation at this level (two-tailed)

 Table 3
 Correlation between clinicopathological findings and expression of PI3K subunits, AKT and p-AKT (T308) in human CCA tissues

Clinicopathologic factors	No. of patients	p110a			p85a			AKT			p-Al	p-AKT (T308)		
		-/Low	High	Р	_	+	Р	_	+	Р	_	+	Р	
Age (years)														
≤58	14	6	8	0.765	10	4	1.000	5	9	0.143	7	7	0.491	
>58	16	6	10		12	4		10	6		6	10		
Gender														
Female	9	4	5	1.000	7	2	1.000	5	4	1.000	3	6	0.691	
Male	21	8	13		15	6		10	11		10	11		
Histological types														
Non-paillary	16	4	12	0.073	10	12	0.266	5	11	0.028*	4	12	0.030*	
Papillary	14	8	6		6	2		10	4		9	5		
Metastasis														
Non-metastasis	15	8	7	0.136	14	1	0.035*	9	6	0.273	9	6	0.065	
Metastasis	15	4	11		8	7		6	9		4	11		

*P≤0.05, statistically significant

However, expression of $p85\alpha$ showed fewer positive cases than $p110\alpha$, although $p110\alpha$ could be directly activated by RAS proteins [32], which are usually altered in CCA. A high frequency of KRAS proto-oncogene mutation was found in 4–60% among Japanese and Thai CCA patients [33–36] and 100% of English patients [37]. Moreover, there are studies that indicate elevated expression of mutated KRAS has been correlated with a more aggressive phenotype of CCA [38–40]. Furthermore, our data revealed that the expression of p85 α was significantly associated with metastasis. Phosphorylation at T308 by PDK1 and S473 by mTORC2 is essential for maximal activation of AKT, although phosphorylation of T308 in AKT is essential and necessary for the cell transforming activity and for signaling [41], possibly as a result of changing in conformation that allows substrate binding and phosphorylation [42]. However, several studies focused on S473 rather than T308 phosphorylation, probably caused by relatively effective S473 antibodies, particularly in immunohistochemistry resulting in analyzing p-AKT (T308) in few studies. Thus, this study aimed to reveal the expression of p-AKT (T308) in CCA. Previous studies demonstrated an activation of the PI3K/AKT pathway in human CCA tissues which more than 50 % of cases expressed p-AKT and correlated with poor prognosis

Table 4 Correlation between clinicopathological findings and expression of mTOR, p-mTOR (2448), GSK-3β and p-GSK-3β (S9) in human CCA tissues

Clinicopathologic factors	No. of patients	mTOR			p-mTOR (S2448)			GSK-3β			p-GSK-3β (S9)		
		_	+	Р	_	+	Р	_	+	Р	_	+	Р
Age (years)													
≤58	14	3	8	0.105	12	2	0.657	7	7	0.296	11	11	0.689
>58	16	11	8		12	4		11	5		3	5	
Gender													
Female	9	3	8	0.419	7	2	1.000	5	4	1.000	5	4	0.195
Male	21	11	8		17	4		13	8		17	4	
Histological types													
Non-papillary	16	4	12	0.156	13	3	1.000	6	10	0.007*	10	6	0.226
Papillary	14	7	7		11	3		12	2		12	2	
Metastasis													
Non-metastasis	15	9	6	0.008*	13	2	0.651	12	3	0.025*	12	3	0.682
Metastasis	15	2	13		11	4		6	9		10	5	

*P≤0.05, statistically significant

Clinicopathologic factors	No. of patients	PTEN		
			+	Р
Age (years)				
≤58	14	9	5	0.694
>58	16	12	4	
Gender				
Female	9	6	3	1.000
Male	21	15	6	
Histological types				
Non-papillary	16	8	8	0.017*
Papillary	14	13	1	
Metastasis				
Non-metastasis	15	12	3	0.427
Metastasis	15	9	6	

Table 5 Correlation between clinicopathological findings and PTEN expression in human CCA tissues

*P≤0.05, statistically significant

[18–23]. These data are consistent with our results as shown by 57 % of the cases positive for p-AKT (T308) and correlated with poor prognosis. Our results showed a few cases, 20 % positive for p-mTOR with no correlation to any clinical data. Chung and coworkers reported that p-mTOR expression was elevated in 83.7 % in extrahepatic CCA tissues [20]. Additionally, there was an altered expression of GSK- 3β , which was previously reported to be a tumor suppressor in various types of cancer [43-45]. There is a single study that demonstrated the increased levels of p-GSK-3 (S9), an inactive form in a mouse model of CCA carcinogenesis [22]. In this study, we report the expression of GSK-3 β and its inactive form, p-GSK-3ß (S9), in human CCA tissues. Our IHC results revealed a strong positive staining intensity in tumors when compared with adjacent normal bile ducts. We found that about 40 % of cases expressed GSK-3β, which correlated with metastasis and histological type of the poor prognosis. Our finding was supported by growing evidence which indicated that GSK-3 β can positively regulate cell survival and proliferation [46], as well as promote tumor development and progression. Moreover, inhibition of GSK-3 β can sensitize tumor cells to chemotherapeutic drug as shown in pancreatic cancer [47]. In addition, we found that only 8 of 30 cases were positive for p-GSK-3 β , an inactive form of GSK-3ß. Furthermore, the expression of PTEN, which was defined as a tumor suppressor and a major negative regulator of PI3K/AKT signaling, was investigated. We found loss of PTEN expression in the majority of CCA patients (70 %) and consistent with the study of Chung and coworkers, which showed the absence of PTEN expression in extrahepatic cholangiocarcinoma patients [20]. Moreover, we found that among PTEN positive cases, their PTEN was inactivated through phosphorylation. The results are consistent with the study of Xu and coworkers [22].

We analyzed for the first time in CCA the therapeutic potential of a PI3K/mTOR inhibitor, NVP-BEZ235, an imidazo[4,5-c]quinoline derivative that inhibits activity by binding to the kinase ATP-binding cleft. This compound equally targets wild-type, mutated p110 and mTOR [28] as well as in cells with loss of PTEN expression [48]. We tested the growth inhibition effect of NVP-BEZ235 on a panel of CCA cell lines, which showed PTEN inactivated by phosphorylation. We found that NVP-BEZ235 could suppress CCA cell growth in the one nanomolar range of IC₅₀ values in a dose-dependent manner. Moreover, the agent exerted its anti-cancer activity by reducing AKT and mTOR phosphorylation. Furthermore, flow cytometric analysis demonstrated that NVP-BEZ235 was capable of inducing G1 cell cycle arrest while lacking significant apoptosis induction. On the other hand, NVP-BEZ235 activates the CCA cells undergoing autophagy, a type II cell death which was indicated by increased LC3-II, an autophagic marker level, in a dose-and time-dependent manner. In many cancers, NVP-BEZ235 had no apparent apoptosis induction [48-52]. Liu and coworker demonstrated previously the cell growth inhibition as well as autophagy induction of NVP-BEZ235 in human glioma cells [50]. This is not surprising because NVP-BEZ235 target is mTORC1, an autophagic blocker resulting in autophagy induction. Although many studies demonstrate the proapoptotic properties of NVP-BEZ235 in breast cancer [28], waldenstrom macroglobulinemia [53], and hepatocellular carcinoma [54]. Additionally, it is known that autophagy is a cellular response to stress conditions, which is implicated in both pro-survival and pro-death processes in

Fig. 2 NVP-BEZ235 inhibits CCA cell proliferation. a Seven CCA cell lines were treated with NVP-BEZ235 with the indicated concentration for 48 h. and cell proliferation was assessed using SRB assay. Data are shown as mean±SEM of one representative experiment. Each culture was done in three independent experiments. b Effects of NVP-BEZ235 on reducing of AKT and mTOR phosphorylation in CCA cells. Cells were treated with NVP-BEZ235 as indicated concentration for 48 h. Cell extracts were subjected to immunoblotting analysis for AKT, p-AKT (S473), mTOR, and p-mTOR (S2448), respectiviely. c NVP-BEZ235 causes cell cycle arrest at G1 phase. Cells were treated with NVP-BEZ235 for 48 h and subjected to propidium iodide staining analysis. The results were shown as the mean±SD from three independent experiments.*P<0.05 compared with control cells



normal cells, as well as in cancer. It is also defined as a mechanism that may enable tumor cells to survive antineoplastic therapy. In addition, there are several lines of evidence demonstrating that many anti-cancer agents can also induce autophagy [55–58]. There are studies suggesting that NVP-BEZ235 alone was not sufficient to induce apoptosis; hence, this compound should be combined with other therapeutic approaches, such as MEK inhibitor [59] or radiotherapy [49, 52]. Furthermore, some studies mentioned that combined NVP-BEZ235 with autophagy inhibition can enhance its anti-cancer efficacy by driving the cancer cells to undergo apoptosis [60, 61]. Since our IHC analysis demonstrated the expression of several key components in PI3K/AKT pathway is correlated with metastasis in CCA patients, we tested if NVP-BEZ235 could inhibit CCA cell migration. As shown in Fig. 4, treated CCA cells with 10 nM of NVP-BEZ235 resulted in reduced CCA cell migration. This result suggests that this pathway plays a role in metastasis of CCA. This is supported by the study of Menakongka and colleagues who reported that hepatocyte growth factor promotes CCA cell invasiveness through PI3K pathway [27]. Furthermore, there are several reports that are

Fig. 3 NVP-BEZ235 induces autophagy without remarkable effect on apoptosis in CCA cells. a Cells were treated with NVP-BEZ235 as indicated concentration for 48 h. then apoptosis cells were analyzed by using Annexin-V-FLUOS staining kit and flow cytometry. **b** NVP-BEZ235 induces autophagy in CCA cell lines. Cells were treated with NVP-BEZ235 for 24 and 48 h, respectively. Protein extracts were analyzed by Western blot with specific antibodies against LC3B (LC3-II), an autophagic marker. The apparent intensity of bands on the membranes was estimated using densitometry





Fig. 4 NVP-BEZ235 reduced CCA cell migration. The effect of NVP-BEZ235 on the migration of CCA cell lines was determined by in vitro wound healing assay. The extent of wound closure was observed and photographed using a phase-contrast microscope with a digital camera at 0, 12 and 24 h incubation

consistent with our finding that NVP-BEZ235 can inhibit cancer cell migration [51, 52]. Moreover, Manara and coworkers also reported that NVP-BEZ235 suppressed cell metastasis of Ewing sarcoma xenografts in vivo mouse models [51]. Therefore, the PI3K pathway may be used as a potential target for treatment or prevention of metastasis in CCA.

Taken together, we demonstrated the constitutive activation of PI3K/AKT pathway in CCA patients as well as CCA cell lines. The possible mechanisms are PI3K overexpression as well as PTEN inactivation by both loss of expression and phosphorylation resulting in poor prognosis in human CCA studied. Therefore, this particular pathway represents a possible useful target for CCA treatment. This work also indicates that inhibition of PI3K and mTOR activity by a novel inhibitor, NVP-BEZ235, has anti-proliferation as well as migration inhibition effects on CCA cells. However, this small inhibitor failed to induce apoptosis of CCA cells while autophagy induction could be observed. Based on these findings, NVP-BEZ235 might be applied in a therapeutic strategy for CCA; however, using this particular drug in combination with other therapies should be considered.

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Conflicts of interest None

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