

# Targeted therapy in biliary tract cancers—current limitations and potentials in the future

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**Abstract:** Biliary tract cancers (BTC)/Cholangiocarcinoma (CCA) is an aggressive biliary tract epithelial malignancy from varying locations within the biliary tree with cholangiocyte depreciation., including intrahepatic cholangiocarcinoma (iCCA) (iCCA), extrahepatic cholangiocarcinoma (eCCA) and gallbladder carcinoma (GBC). The disease is largely heterogeneous in etiology, epidemiology, and molecular profile. There are limited treatment options and low survival rates for those patients with advanced or metastatic disease. Systemic treatment is confined to cytotoxic chemotherapy with the combination of gemcitabine and cisplatin. Lack of a stereotype genetic signature makes difficult in identification of potential actionable target directly, which may also explain lack of obvious clinic benefit with target oriented agents from current studies. It is crucial to understand of BTC carcinogenesis, tumor-stroma interactions, and key molecular pathways, and herald to establish targeted, individualized therapies for the heterogeneous disease, and eventually to improve the survival and overall outcome of patients.

**Keywords:** Biliary cancer; hepatocellular carcinoma; abnormal liver function; chemotherapy

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## Introduction

Biliary tract cancers (BTC)/Cholangiocarcinoma (CCA) are resulting from malignant transformation of epithelial cells within the bile system as related but distinct malignancies along the intrahepatic and extrahepatic biliary tree. Besides the gallbladder carcinoma (GBC), CCA is also refers to cancers of the entire biliary tree. CCA is commonly classified as intrahepatic cholangiocarcinoma (iCCA) and extrahepatic cholangiocarcinoma (eCCA), with extrahepatic further divided into perihilar cholangiocarcinoma (pCCA) and distal cholangiocarcinoma (dCCA) tumors based on anatomical location with the cystic duct as the point of distinction (1). There is increasing recognition that these subtypes separated by anatomic origins of BTCs distinct

not only in desmoplasia markers of cholangiocytes, but also in biological and clinical characteristics (2). BTC has an aggressive natural course without early specific warning sign of the disease; therefore, potential curative surgery is only suitable for a limited number of patients. So far, systemic treatment is confined to the gemcitabine and cisplatin combination as the practice standard for patients with advanced and metastatic disease (3). The overall outcome is disappointing with limited response rate (RR) and low 5-year survival rate. It is crucial to understand BTC/CCA carcinogenesis, tumor-stroma interactions, and key molecular pathways, and herald to establish targeted, individualized therapies for the heterogeneous disease, and eventually to improve the survival and overall outcomes of patients. In this review, we are going to summarize the

results of systemic cytotoxic chemotherapy, and review the data from studies with ‘target-oriented’ agents. Since most of these studies are not targeting the distinguished subgroups of the disease specifically, they are more likely as ‘target intended’ rather than ‘target-oriented’. We will also discuss the current knowledge regarding the genetic basis of this disease, including molecular pathways involved in its carcinogenesis, and potential targeted therapies that hold promise in the future research and practice.

### Cytotoxic chemotherapy

Cytotoxic chemotherapy remains the mainstay of treatment for patients with advanced unresectable or metastatic BTC. Given the rarity of this disease, clinical studies have been small and have almost always been combined with ‘lumping’ various BTCs with very few randomized trials have been conducted. The majority of trials have been performed with either fluoropyrimidine-based or gemcitabine-based combination. 5-fluorouracil (5-FU) had been tested in small trials, both as monotherapy and in combinations. Overall RRs in these studies varied from 10% to 40%; median survival also varied notably, from 5 to 16 months (4-21).

A phase III study randomized 54 patients with previously untreated advanced biliary cancer between ECF (epirubicin, cisplatin, 5-FU) and FELV (5-FU/LV, etoposide) (14). The median OS was not significantly different between the two arm (9.02 months in ECF *vs.* 12.03 months in FELV,  $P=0.2059$ ). Objective RRs were also similar (19.2% in ECF *vs.* 15% in FELV,  $P=0.72$ ). The interesting point is greater than 60% of patients in each arm demonstrated resolution of pain, anorexia, weight loss, and nausea.

Based on the data of studies from advanced and metastatic pancreatic cancer, gemcitabine has also been evaluated biliary cancers at the similar setting (7,12,15,16,22-41) (*Table 1*). As a single agent, gemcitabine showed only moderate efficacy with RRs ranging from 0% to 30% at varied dosing schemes demonstrated. Efforts had been tried in combination of gemcitabine with multiple other cytotoxic agents, including 5-FU, capecitabine, cisplatin, oxaliplatin, and irinotecan with significant variations in RRs and survival.

In order to assess the overall efficacy of systematic chemotherapy, a pooled analysis was performed including 104 trials published between 1985 and 2006 (42). The overall RR was 22.6%; and the tumor control rate (TCR) was 57.3%. Significant correlations of RR and TCR with survival times were found. Subgroup analysis showed superior RRs for GBC compared with CCA (RR

35.5% *vs.* 17.7%,  $P=0.008$ ), but shorter OS for GBC (median 9.3 in CC *vs.* 7.2 months in GBC,  $P=0.048$ ). Based on treatment type subgroup analyses, the analysis showed that regimens containing both gemcitabine and a platinum agent had significantly higher response and TCRs compared to either fluoropyrimidine or gemcitabine monotherapy or fluoropyrimidine-plus-platinum regimens. Based on this information, United Kingdom-based Advanced Biliary Care (ABC)-02 trials was conducted to validate the combination (3). In this study, 410 patients with non-resectable, recurrent, or metastatic BTC were randomized to receive either gemcitabine alone or gemcitabine and cisplatin combination. The trial included patients with CCA, gallbladder cancer, or ampullary cancer. Patients in the combination arm received cisplatin 25 mg/m<sup>2</sup> and gemcitabine 1,000 mg/m<sup>2</sup> on days 1 and 8 of a 3-week cycle, for eight cycles; patients in the gemcitabine monotherapy arm received gemcitabine 1,000 mg/m<sup>2</sup> on days 1, 8, and 15 of a 4-week cycle, for six cycles. After a median follow-up of 8.2 months, the median OS was 11.7 months in the gemcitabine/cisplatin combination arm, compared to 8.1 months in the gemcitabine arm (HR =0.64,  $P<0.001$ ). The median progression-free survival (mPFS) was improved with the combination (8.0 *vs.* 5.0 months; HR =0.64;  $P<0.001$ ) as well. Severe hematologic toxicities were seen more frequently in the combination arm. However, there were more severe liver toxicities reported in the gemcitabine-alone arm for unclear reason.

### Target intended therapy

Although the combination of gemcitabine and cisplatin has achieved some advances in the treatment of advanced and metastatic biliary track cancers and has been accepted as the standard treatment option as the first line therapy since 2010, the overall outcome is still disappointing. Clinic studies of target-oriented agents (most of them in combination with gemcitabine based regimen) have been attempted for improving the outcomes of the disease. Those target-oriented agents primarily are monoclonal antibodies and tyrosine kinase inhibitors against EGFR and vascular endothelial growth factor (VEGF) (43-57) (*Table 2*). However, no or only marginal benefits showed from those trials, which is likely because of a mixed cohort of BTC patients (iCCA, eCCA, and GBC), and the underlying genetic variability of the disease.

One phase III trial randomized 268 patients with

**Table 1** Gemcitabine-based studies in advanced BTCs

Authors	N	Treatment regimen	RR (%)	mPFS (mo)	mOS (mo)
Raderer <i>et al.</i> (7)	19	Gem	16	2.5	6.5
Penz <i>et al.</i> (22)	32	Gem	22	5.6	11.5
Gebbia <i>et al.</i> (23)	18	Gem	22	3.4	8
	22	Gem + 5-FU + LV	36	4.1	11
Kuhn <i>et al.</i> (24)	43	Gem + docetaxel	9	NR	11.0
Bhargava <i>et al.</i> (25)	14	Gem + irinotecan	14	1.5	NR
Kornek <i>et al.</i> (12)	25	Gem + MMC	20	4.2	6.7
André <i>et al.</i> (26)	23	Gem + Ox	22	3.9	7.6
Knox <i>et al.</i> (27)	27	Gem + 5-FU	33	3.7	5.3
Alberts <i>et al.</i> (28)	42	Gem + 5-FU + LV	10	4.6	9.7
Thongprasert <i>et al.</i> (29)	40	Gem + Cis	28	4.8	8.4
Knox <i>et al.</i> (30)	45	Gem + Cape	31	7	14
Cho <i>et al.</i> (15)	44	Gem + Cape	32	6.0	14
Cho <i>et al.</i> (16)	24 <sup>b</sup>	Gem + Cape	33	6.0	16
Lee <i>et al.</i> (31)	24 <sup>a</sup>	Gem + Cis	21	5.0	9.3
Kim <i>et al.</i> (32)	29	Gem + Cis	34	3.0	11.0
Harder <i>et al.</i> (33)	31	Gem + Ox	26	6.5	11
Manzione <i>et al.</i> (34)	34	Gem + Ox	41	NR	10
Alberts <i>et al.</i> (35)	58	Gem + Pem	NR	3.8	6.6
Riechelmann <i>et al.</i> (36)	75	Gem + Cape	29	6.2	12.7
Lee <i>et al.</i> (37)	39	Gem + Cis	17	3.2	8.6
André <i>et al.</i> (38)	67	Gem + Ox	15	349	8.8
Meyerhardt <i>et al.</i> (39)	33	Gem + Cis	21	6.3	9.7
Kim <i>et al.</i> (40)	40	Gem + Ox	15	4.2	8.5
Jang <i>et al.</i> (41)	53	Gem + Ox	19	4.8	8.3

<sup>a</sup>, CCA patients only; <sup>b</sup>, gallbladder cancer patients only. RR, response rate; mPFS, median progression-free survival; mOS, median overall survival; Gem, gemcitabine; NR, not reported; 5-FU, 5-fluorouracil; LV, leucovorin; MMC, mitomycin-C; Cis, cisplatin; Cape, capecitabine; Ox, oxaliplatin; Pem, pemetrexed; BTC, biliary tract cancers; CCA, cholangiocarcinoma.

metastatic biliary tract adenocarcinoma to the combination of gemcitabine and oxaliplatin (GemOx) with erlotinib or chemotherapy (GemOx) alone (49). The combination showed increased RR (30% *vs.* 16%,  $P=0.005$ ), however, there were no significant difference in mPFS (5.8 *vs.* 4.2 months,  $P=0.087$ ) and median overall survival (mOS) (9.5 months for both arms). A randomized phase II study evaluated the GemOx with or without cetuximab in 150 patients

with advanced and metastatic BTC (51). There were no differences between the two arms in RR, mPFS or mOS. The TCOG trial evaluated the same regimen in advanced and metastatic BTC patients with stratification of iCCA/eCCA/GBC (71.3%/16.4%/12.3%) and KRAS status (36.1% KRAS mutation). On the intent-to-treat analysis, it showed some benefit of mPFS in the arm of GemOX + cetaximab (7.1 *vs.* 4.0 months,  $P=0.0069$ ), but no difference

**Table 2** Selected studies of targeted agents in advanced BTCs

Study	Number of patients	RR (%)	mPFS (mo)	mOS (mo)	References
Bevacizumab (VEGF-A)					
Bev + Gem + Ox	35	40	7.0	12.7	Zhu <i>et al.</i> (43)
Sorafenib (VEGFR-2/3, PDGFR, RAF)					
Sorafenib	46	2	2.3	4.4	Bengala <i>et al.</i> (44)
Sorafenib	31	0	3	9	El-Khoueiry <i>et al.</i> (45)
Sunitinib (VEGFR, PDGFR, cKit)					
Sunitinib	56	9	1.7	4.8	Yi <i>et al.</i> (46)
Erlotinib (EGFR)					
Erlotinib	43	8	2.6	7.5	Philip <i>et al.</i> (47)
Erlot + docetaxel	11	0	NR	5.7	Chiorean <i>et al.</i> (48)
Erlot + Gem + Ox	135	30	5.8	9.5	Lee <i>et al.</i> (49)
Gem + Ox	133	16	4.2	9.5	
Cetuximab (EGFR)					
Cet + Gem + Ox	9 <sup>a</sup>	11	4	7	Paule <i>et al.</i> (50)
Cet + Gem + Ox	76	24	6.1	11.0	Malka <i>et al.</i> (51)
Gem + Ox	74	23	5.5	12.4	
Panitumumab (EGFR)					
Pan + Gem + Ox	46	33	8.3	10.0	Jensen <i>et al.</i> (52)
Pan + Gem + irinotecan	35	31	9.7	12.9	Sohal <i>et al.</i> (53)
Lapatinib (HER2)					
Lapatinib	17	0	1.8	5.2	Ramanathan <i>et al.</i> (54)
Selumetinib (MEK1/2)					
Selumetinib	28	12	3.7	9.8	Bekaii-Saab <i>et al.</i> (55)
Target-oriented agent combination					
Erlot + Bev	49	12	4.4	9.9	Lubner <i>et al.</i> (56)
Erlot + Soraf	34	6	2	6	El-Khoueiry <i>et al.</i> (57)

<sup>a</sup>, CCA patients only. RR, response rate; mPFS, median progression-free survival; mOS, median overall survival; EGFR, epidermal growth factor receptor; Erlot, erlotinib; NR, not reported; Gem, gemcitabine; Ox, oxaliplatin; Cet, cetuximab; Pan, panitumumab; VEGF, vascular endothelial growth factor; Bev, bevacizumab; VEGFR, vascular endothelial growth factor receptor; PDGFR, platelet-derived growth factor receptor; RAF, rapidly accelerated fibrosarcoma; Soraf, sorafenib; HER2, human epidermal growth factor receptor 2; MEK, mitogen-activated protein kinase/extracellular-signal regulated kinase; BTC, biliary tract cancers; CCA, cholangiocarcinoma.

in RR (27.3% *vs.* 15%,  $P=0.1223$ ), and mOS (10.3 *vs.* 8.8 months,  $P=0.4057$ ). Planned subgroup analysis showed that potential more benefits of cetuximab with GemOX in KRAS mutated patients with mPFS of 7.1 *vs.* 1.9 months ( $P=0.0351$ ), however, no statistical significance in mOS (10.3 *vs.* 6.6 months,  $P=0.6924$ ) (58). Interestingly, a pooled

analysis with 161 trials comprising 6,337 patients (trials published in English between 1/2000 and 1/2014 as well as ASCO abstracts 2010 to 2013) showed some potential survival benefits of gemcitabine-based chemotherapy with targeted therapy (predominantly EGFR inhibitor as either monoclonal antibody or TKI) (59).

### New insights into the molecular pathogenesis and therapeutic opportunities

Recent studies have begun to uncover the genetic and molecular processes underlying carcinogenesis using advanced technologies such as next-generation sequencing (NGS) in the BTC. The emerging knowledge and data generated from studies of epidemiology, genome profiling, and laboratory based investigations provide new insights into risk factors, genomic composition, cellular origins and contribution of the tumor microenvironment to the pathogenesis of BTC. The remarkable genetic heterogeneity of BTC may be the result of a complex interplay among different factors—some of them are shared by most human cancers, while others may be unique for this malignancy. Emerging data supports that iCCA, eCCA, and GBC represent distinct tumors arising from different genetic backgrounds (60,61).

#### Cancer-associated fibroblasts (CAF) in the tumor stroma

CAF secondary to desmoplastic response is a prominent tumor microenvironment characteristics of biliary track cancers, especially iCCA. CAF may impede access of therapeutic agents to the tumor and pose therapeutic challenges further besides the genetic heterogeneity of this malignancy (1,62). Preclinical studies have demonstrated a reduction in fibrosis and carcinogenesis in BTC/CCA with 1D11, a transforming growth factor  $\beta$  (TGF- $\beta$ ) antagonist, as well as curcumin, a nutraceutical agent (63). A pre-clinic study with an orthotopic CCA model showed that the BH3 (BCL-2 family protein, pro-apoptosis) mimetic, Navitoclax, enhanced selective CAF apoptosis, decreased expression of  $\alpha$ -SMA, and reduced tumor burden and metastasis while improving survival (64). Further preclinical and clinical studies are needed to explore the role of antifibrotic therapies in CCA chemoprevention.

#### Inflammation and carcinogenesis

Chronic inflammation plays a significant role in the development of BTC. Chronic inflammatory pathways not only are key components in carcinogenesis, but also promote tumor invasion and migration. Primary sclerosing cholangitis (PSC), hepatobiliary flukes *Opisthorchis viverrini* (*O. viverrini*) and *Clonorchis sinensis* characterized by chronic biliary tract inflammation and liver injury, are common predisposing conditions for BTC. In addition, prolonged

hepatolithiasis may promote CCA development by calculi occurring proximal to the hepatic duct confluence (2,65). Inducible nitric oxide synthase (iNOS) activation by inflammatory cytokines contributes to nitrosative stress by generation of excess nitric oxide, which then results in inhibition of DNA repair proteins, and single-stranded, double-stranded, and oxidative DNA lesions. Oxidative stress via generation of oxysterols, cholesterol oxidation products present in human bile, creates a milieu favorable for tumor development and progression by activating Hedgehog signaling pathway. Oxysterols, bile acids, and iNOS all stimulate over-expression of cyclooxygenase-2, which has been implicated carcinogenesis of BTC (66-68). Therefore, inflammatory processing control may play a significant role in management of BTC, especially in the prevention.

#### Genomic profiling studies

Genomic profiling has demonstrated characteristic profiles for iCCA and eCCA. Next generation sequencing (NGS) of a BTC series showed key variations in certain mutations based on tumor location (69). Mutations in the isocitrate dehydrogenase 1 and 2 (IDH1 and IDH2) genes have consistently been shown to be more frequent in iCCA versus eCCA or GBC. IDH mutations caused inhibition of HNF-4 $\alpha$  (hepatocyte nuclear factor 4 alpha, a nuclear receptor; also known as NR2A1- nuclear receptor subfamily 2, group A, member 1), leading to impaired hepatocyte differentiation and increased cell proliferation, and associated with poorly differentiated histology. FGFR (Fibroblast growth factor receptor) mutations, specifically in FGFR2 are associated with the production of two fusion kinase genes, FGFR2-AHCYL1 and FGFR2-BICC1 that are mutually exclusive with KRAS/BRAF mutations. The median cancer specific survival is suggested significantly longer for patients whose tumors contained FGFR2 translocations (70).

It appears that mutational profiles may be influenced by the etiology of BTC. A study revealed the results of exome sequencing of 209 CCA samples from Asia and Europe with 108 cases of *O. Viverrini* infection related and the other 101 cases has non-*O. viverrini*-related etiologies (71). The study showed that TP53 was mutated in 40% of *O. viverrini* infection related CCA, and only in 9% of the non-*O. viverrini*-related cases ( $P < 0.001$ ). On the other hand, SMAD4 mutation was more frequently in non-*O. viverrini*-related CCA (6% vs. 19%,  $P = 0.006$ ). IDH1 and



IDH2 mutations were tested in 22.2% of non-*O. viverrini* iCCA and only 3.2% of *O. viverrini*-related iCCA.

Activating mutations in cell proliferation oncogenes lead to uncontrolled cell growth and survival. The Ras/MAPK signaling pathway plays a key role in cell growth, differentiation, survival, and migration. Gain-of-function mutations in KRAS are present in approximately 45–55% of iCCA and 10–15% of eCCA. One study showed that BRAF, an important downstream effector of KRAS, was found to be mutated in 22% of iCCA, no BRAF mutation was found in those cases with KRAS mutations; however, KRAS mutations were seen in 20% of tumors with BRAF mutations (72). The ErbB family consists of four receptor kinases, including ErbB1 (or EGFR) and ErbB2 [or human epidermal growth factor receptor 2 (HER2)]. Mutations in the EGFR gene were seen in 15% of CCA cases (73). MET is an oncogene that encodes for the hepatocyte growth factor (HGF) receptor. HGF/MET pathway is less established but may be important in development and progression of BTC/CCA. MET is a key regulator of invasive growth. Interaction of HGF and its receptor MET can activate many pathways including MAPK, PI3K and STAT. Overexpression of MET occurs in 12–58% of cases of iCCA and has been linked to overexpression of members of the EGFR family and shown the capacity of HGF to stimulate migration and invasion in CC cells (74).

Loss-of-function mutations in tumor suppressor genes also play a role in CCA. CDKN2A (p16INK4a) negatively regulates proliferation in normal cells and is capable of cell cycle arrest. This tumor suppressor gene was highly mutated in reports with 55% loss-of-function in iCCA and 83% in eCCA. TP53, a principal regulator of cell division, appears to be inactivated in approximately one-third of BTC, both iCCA and eCCA. SMAD4, in conjunction with the other SMAD proteins, is an end effector in the TGF $\beta$  pathway with promotes epithelial-mesenchymal transition, directly regulating the activity of genes controlling cell proliferation. Mutations in SMAD4 were described in up to 40% in iCCA with the relationship of disease staging (75). A recent large cohort study with 103 iCCA in identification of an iCCA-specific mutation signature that is associated with liver inflammation, fibrosis and cirrhosis (76). The study found that TP53-defection is more likely to be HBsAg-seropositive, whereas KRAS mutations are nearly exclusively in HBsAg-seronegative CCA patients. The study demonstrated that three pathways (Ras/PI3K, p53/cell cycle, and TGF $\beta$ /Smad), genes important for epigenetic regulations and oxidative phosphorylation are substantially affected in iCCA.

### **Potential role of stem and progenitor cells**

Besides the different carcinogenetic mechanisms driven by risk factors in BTC, the distinct genetic profiles are also reflecting two distinct stem cell niches, the canals of Hering harboring hepatic stem cells (HpSCs) and the peribiliary glands harboring biliary tree stem/progenitor cells (BTSCs) (77). Cell populations from HpSCs and BTSCs lineages may represent distinct candidate cells of origin during CC carcinogenesis, susceptible to distinct risk factors and responsible for the development of the different iCCA and eCCA or GBC subtypes: e.g., BTSC lineage may be activated under pathological conditions affecting the large intrahepatic and extrahepatic bile ducts (including liver flukes, cholangitis, PSC, hepatolithiasis, etc.), giving rise to large bile duct pure mucin secreting iCCA and eCCA; conversely, the hHpSC lineage has been suggested to be activated in response to parenchymal liver diseases (such as chronic viral/non-viral liver disease, schistosomiasis and liver cirrhosis) and to be involved in the development of combined hepatocellular carcinoma-iCCA, bile ductular iCCA and mixed iCC A (with a focal hepatocytic differentiation, ductular reaction and mucin-secreting adenocarcinoma). This stem cell compartment is probably activated also during nonalcoholic steatohepatitis (NASH) and asbestos exposure, as these two risk factors are exclusively associated with the development of ICC.

### **Summary and future directions**

Despite some advances in treatment of BTCs, the overall outcomes of the disease remain poor. With learning from ‘target intended’ studies and emerging understanding of the heterogeneity and the molecular landscape of BTC, future target oriented research/studies should be based on the underlining etiology, the specific genetic profile of each subgroup, and cancer-stroma microenvironment of the selected particular disease population. A recent retrospective study showed the promising therapy with blockage of Her-2/neu in BTC (with both GBC and CCA) patients who with the gene amplification (78). Advances in immunotherapy may also provide new opportunities for treating BTC (79). A complete response (CR) was reported in a chemotherapy refractory metastatic iCCA patient with mismatch-repair deficiency (dMMR) after being treated with PD-1 inhibitor (80). Most current on-going ‘target-oriented’ studies are less distinctive regarding the specificity of the characteristic mechanism of the disease (Tables 3,4). However, we are certain that the future studies will be more precisely to meet our goal.

Table 3 Current targeted studies (clinicaltrials.gov)

Treatment	Phase	Disease	Target	Schedule	Start date	Complete date	Sponsor	Clinicaltrials.gov ID
Ramucirumab	II	Advanced, pre-treated biliary	VEGFR1	8 mg/kg IV on day 1 of each 14 day cycle	12/2015	12/2019	M.D. Anderson Cancer Center	NCT02520141
Regorafenib	II	Refractory advanced biliary	—	—	04/2014	10/2018	H. Lee Moffitt Cancer Center and Research Institute	NCT02115542
Selumetinib (AZD6244) with cisplatin/gemcitabine (Cis/Gem) versus Cis/Gem alone	Phase II/ randomized	Advanced, biliary, first line	MEK 1/2 inhibitor	Selumetinib, orally, BID from days 1–21 of every 28 day cycle. Cisplatin/gemcitabine, IV, on days 1 and 8 of every 28 day cycle	11/2014	05/2017	University Health Network, Toronto	NCT02151084
Afatinib dimaleate (BIBW 2992) and capecitabine	II	Advanced refractory pancreatic cancer or biliary cancer	EGFR/HER2 tyrosine kinase inhibitor	Afatinib dimaleate PO QD on days 1–28 and capecitabine PO BID on days 1–14. Courses repeat every 21 days	11/2015	11/2018	University of Washington	NCT02451553
Ponatinib hydrochloride	II	Advanced biliary cancer with FGFR2 fusions	FGFR2 blockage	Ponatinib hydrochloride PO QD on days 1–28	12/2014	10/2019	NCI	NCT02265341
Lenvatinib (E7080)	II	BTC, failed gemcitabine-based therapy	—	24 mg lenvatinib, orally	10/2015	10/2017	Eisai Co., Ltd.	NCT02579616
Ramucirumab (LY3009806) or merestininb (LY2801653) or placebo plus gem/cis	—	Advanced or metastatic BTC	—	Ramucirumab plus cisplatin and gemcitabine IV on days 1 and 8, merestininb orally each day, plus cisplatin and gemcitabine IV on days 1 and 8, placebo plus cisplatin and gemcitabine IV on days 1 and 8, every 21 days	04/2016	04/2018	Eli Lilly and Company	NCT02711553

Table 3 (continued)

Table 3 (continued)

Treatment	Phase	Disease	Target	Schedule	Start date	Complete date	Sponsor	Clinicaltrials.gov ID
Regorafenib in combination with mGEMOX	II	-	-	Regorafenib: X mg/d, PO, from days 1 to 14, days 15 to 20; gemcitabine 900 mg/m <sup>2</sup> IV in 30 minutes; oxaliplatin 80 mg/m <sup>2</sup> IV in 120 minutes, days 1 and 8	09/2014	04/2019	Institut du Cancer de Montpellier - Val d'Aurelle	NCT02386397
ADH-1, gemcitabine hydrochloride and cisplatin	I	Metastatic pancreatic or BTC	-	ADH-1 IV over 20–80 minutes on days 1, 4, 8, 11, 15, and 18, cisplatin IV and gemcitabine hydrochloride IV over 30 minutes on days 1 and 8	04/2013/12/2018	-	NCI; Adherex Technologies, Inc.	NCT01825603
Regorafenib	II	Advanced and metastatic biliary	-	Regorafenib 120 mg orally once daily 21 days on and 7 days off in the 28-day cycle	01/2014	02/2018	University of Pittsburgh	NCT020533376
BIBW 2992 with Gem/Cis	I/II	Advanced and metastatic biliary	-	-	08/2012	-	Johannes Gutenberg University Mainz	NCT01679405
Copanlisib (BAY 80-6946) in combination with gemcitabine and cisplatin	II	Advanced and metastatic CCA	-	Cisplatin (25 mg/m <sup>2</sup> ) + gemcitabine (1,000 mg/m <sup>2</sup> ) + copanlisib (60 mg) on days 1 and 8 with days 15 off to be administered on an every 21-days schedule	05/2016	12/2018	H. Lee Moffitt Cancer Center and Research Institute	NCT02631590
Refametinib	II	Advanced biliary, 2 <sup>nd</sup> line	MEK inhibitor	Orally at the starting dose of 50 mg twice daily on a continuous daily dosing schedule	01/2015	07/2016	Ho Yeong Lim, Samsung Medical Center	NCT02346032
RRX-001	II	Advanced cholangiocarcinoma, 2 <sup>nd</sup> line	Epigenetic agent	-	05/2015	05/2018	EpigentRx, Inc.	NCT02452970

Table 3 (continued)



Table 3 (continued)

Treatment	Phase	Disease	Target	Schedule	Start date	Complete date	Sponsor	Clinicaltrials.gov ID
ARQ 087	I/II	Advanced solid tumors with FGFR genetic alterations	FGFR inhibitor	-	12/2012	12/2016	ArQule	NCT01752920
CX-4945 in combination with gemcitabine and cisplatin	I/II	Advanced cholangiocarcinoma, 1 <sup>st</sup> line	-	-	06/2014	12/2016	Senhwa Biosciences, Inc.	NCT02128282
Pazopanib/gemcitabine	II	Biliary tree cancer (CCA or GBC)	-	-	06/2013	11/2016	Hellenic Cooperative Oncology Group/GSK	NCT01855724

BTC, biliary tract cancers; CCA, cholangiocarcinoma; GBC, gallbladder carcinoma; IV, intravenously; NCI, National Cancer Institute.

Table 4 Immunotherapy of BTC (clinicaltrials.gov)

Treatment	Phase	Disease	Target	Schedule	Start date	Complete date	Sponsor	Clinicaltrials.gov ID
Pembrolizumab and GM-CSF	I/II	Advanced Biliary	-	-	04/2016	09/2019	University of California, San Francisco	NCT02703714
Precision T cells specific to personalized neo-antigen	II	Advanced Biliary	DC-PNAT combined with gemcitabine treatment: Gemcitabine: once a week with a total of 6 times before 60 days prior to the start of drawing blood. DC-PNAT: once per 3 weeks with a total of three periods	-	09/2015	09/2017	Second Military Medical University (China)	NCT02632019

BTC, biliary tract cancers; DC-PNAT, dendritic cell-precision T cell for neo-antigen.

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## Footnote

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

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