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Background

- Fibroblast growth factor receptor 2 (FGFR2) fusions or other rearrangements occur in approximately 14% of patients with intrahepatic cholangiocarcinoma (iCCA)^{1–4}
- Futibatinib is a novel and highly selective covalent inhibitor of FGFR1-4 that irreversibly inhibits FGF/FGFR signaling⁵
- In a global phase 2 study (FOENIX-CCA2), 42% of patients treated with futibatinib had a confirmed response (median duration of response: 9.7 months), with a median progression-free survival (PFS) and overall survival (OS) of 9.0 months and 21.7 months, respectively⁵
- These findings led to the accelerated approval of futibatinib by the U.S. Food and Drug Administration in September 2022 for adult patients with previously treated, unresectable, locally advanced or metastatic iCCA harboring FGFR2 gene fusions or other rearrangements⁶
- In April 2023, the European Medicines Agency's Committee for Medicinal Products for Human Use issued a positive opinion recommending the conditional marketing authorization of futibatinib for the second-line treatment of locally advanced or metastatic CCA harboring an *FGFR2* fusion or rearrangement⁷

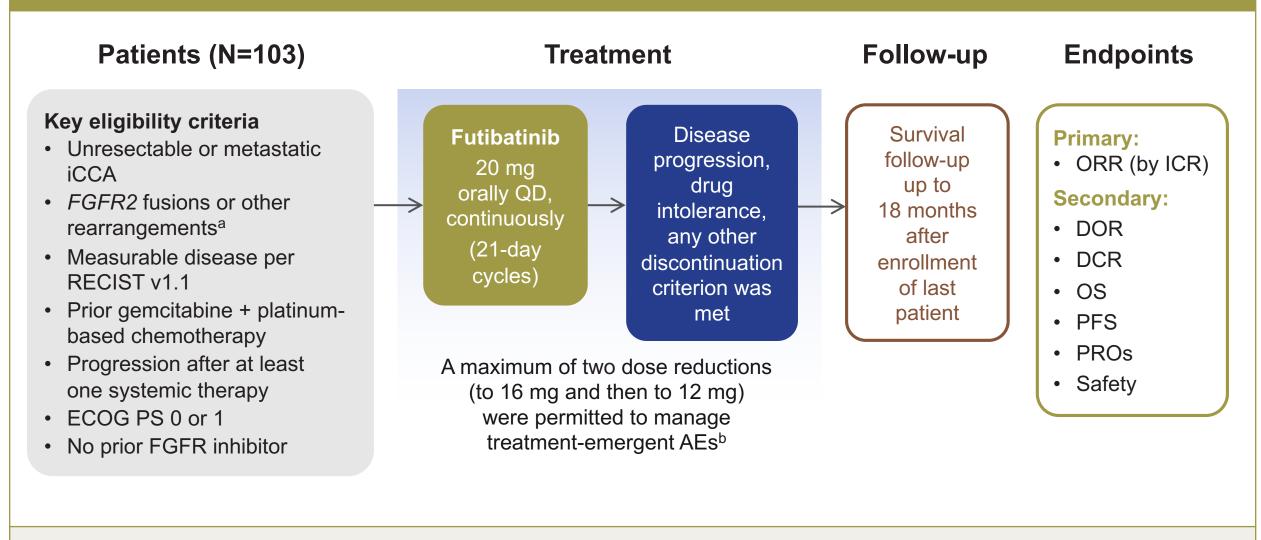
Objective

• To evaluate baseline demographics and measure clinical outcomes in patients with and without a confirmed response to futibatinib in the FOENIX-CCA2 study

Methods

- FOENIX-CCA2 was a multinational, single-arm, phase 2 study (NCT02052778) of futibatinib in patients with unresectable, locally advanced or metastatic, FGFR2 fusion or rearrangement-positive iCCA and disease progression after at least one previous line of systemic therapy, including gemcitabine plus platinum-based chemotherapy⁵
- Patients received oral futibatinib at a dose of 20 mg once daily in a continuous regimen over a 21-day cycle (**Figure 1**)
- The primary endpoint was objective response rate according to Response Evaluation Criteria in Solid Tumors version 1.1, as assessed by independent central review
- Samples were assessed for genetic coalterations by the TruSight Oncology 500 (Illumina) ctDNA sequencing assay
- In this post hoc analysis of the FOENIX-CCA2 study, baseline demographics, PFS, OS, and dose reductions/interruptions were evaluated descriptively for futibatinib responders (partial or complete response based on independent central review) and nonresponders (stable or progressive disease)
- A post hoc exploratory analysis of the differences in baseline genomic alterations between futibatinib responders and nonresponders was also conducted

Figure 1. FOENIX-CCA2 study design



^aIdentified locally or centrally in tumor tissue by Foundation Medicine or by local laboratory testing of tumor tissue or circulating tumor DNA. ^bTreatment was discontinued if treatment-emergent AEs did not resolve after two dose reductions or if the next cycle of treatment was delayed by

AE, adverse event; DCR, disease control rate; DOR, duration of response; ECOG PS, Eastern Cooperative Oncology Group performance status; FGFR, fibroblast growth factor receptor; iCCA, intrahepatic cholangiocarcinoma; ICR, independent central review; ORR, objective response rate; OS, overall survival; PFS, progression-free survival; PRO, patient-reported outcome; QD, once daily; RECIST, Response Evaluation Criteria in Solid Tumors.

Futibatinib in Patients With FGFR2-Rearranged Intrahepatic Cholangiocarcinoma: Responder Analyses of Efficacy and Safety From the Phase 2 FOENIX-CCA2 Study

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Results

Patients

- Overall, 103 patients were enrolled in FOENIX-CCA2; of these, 43 (42%) had a confirmed response to futibatinib, and 60 (58%) were nonresponders Baseline characteristics of responders and nonresponders were generally
- comparable (**Table 1**)

Table 1. Baseline characteristics

Characteristic	Responders ^a (n=43)	Nonresponders (n=60)
Median (range) age, years	60.0 (22–79)	56.0 (28–74)
Sex, n (%) Male Female	17 (39.5) 26 (60.5)	28 (46.7) 32 (53.3)
Race, n (%) White Asian Black Native Hawaiian or other Pacific Islander Unknown	21 (48.8) 13 (30.2) 4 (9.3) 0 5 (11.6)	30 (50.0) 17 (28.3) 4 (6.7) 1 (1.7) 8 (13.3)
Region, n (%) North America Europe Asia Pacific, excluding Japan Japan	19 (44.2) 13 (30.2) 7 (16.3) 4 (9.3)	28 (46.7) 15 (25.0) 7 (11.7) 10 (16.7)
ECOG PS, n (%) 0 1	22 (51.2) 21 (48.8)	26 (43.3) 34 (56.7)
Mean (SD) weight, kg	72.5 (18.2)	74.9 (22.7)

^aPatients with confirmed partial or complete response based on independent central review ECOG PS, Eastern Cooperative Oncology Group performance status; SD, standard deviation

Treatment

- At the data cutoff for this analysis (October 1, 2020), the median duration of treatment among responders and nonresponders was 10.9 months (range: 4.2–24.5) and 6.9 months (range: 0.5–18.9), respectively (**Table 2**)
- Treatment discontinuation was more frequent in nonresponders, most commonly because of disease progression
- Median time to first dose reduction/interruption and median duration of interruption due to adverse events (AEs) were both longer in responders

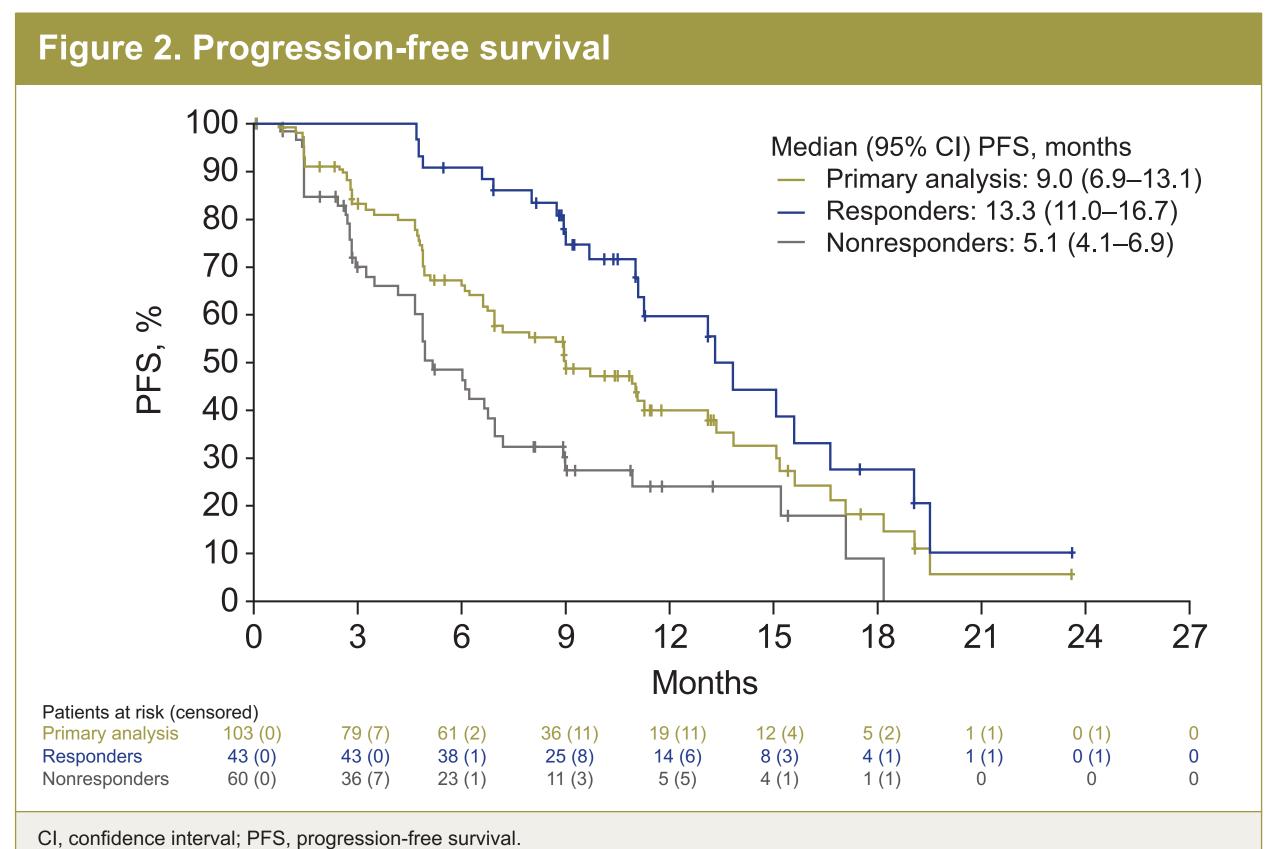
Table 2. Treatment among responders and nonresponders

	Responders ^a (n=43)	Nonresponders (n=60)
Median (range) duration of treatment, months	10.9 (4.2–24.5)	6.9 (0.5–18.9)
Treatment discontinuation, n (%)	22 (51.2)	50 (83.3)
Dose reduction/interruption due to AE, n (%) Dose reduction Dose interruption	28 (65.1) 30 (69.8)	31 (51.7) 38 (63.3)
Median (range) time to first dose reduction/interruption due to AEs, days Dose reduction Dose interruption	84.5 (5–316) 52.0 (4–253)	36.0 (5–332) 22.0 (4–325)
Median (range) duration of interruption, days	23.0 (1–140)	14.0 (1–214)

^aPatients with confirmed partial or complete response based on independent central review. AE, adverse event.

Progression-free survival

- Median PFS was 13.3 (95% confidence interval [CI]: 11.0–16.7) months in responders and 5.1 (95% CI: 4.1–6.9) months in nonresponders (Figure 2)
- The PFS rates at 6 and 12 months were 90.7% (95% CI: 77.1–96.4) and 59.8% (95% CI: 40.7–74.5), respectively, among responders and 46.4% (95% CI: 32.7– 59.1) and 24.1% (95% CI: 12.8–37.3), respectively, among nonresponders



Overall survival

- Median OS was 26.4 (95% CI: not evaluable-not evaluable) months in responders and 14.6 (95% CI: 10.3–21.7) months in nonresponders (**Figure 3**)
- The OS rates at 6 and 12 months were 100% (95% CI: 100–100) and 89.6%
- (95% CI: 74.5–96.0), respectively, among responders and 79.3% (95% CI: 66.4–87.7) and 59.2% (95% CI: 45.1–70.8), respectively, among nonresponders
- The impact of differences in poststudy therapies, including chemotherapy, resection in oligometastatic disease, and other local therapies, was not assessed

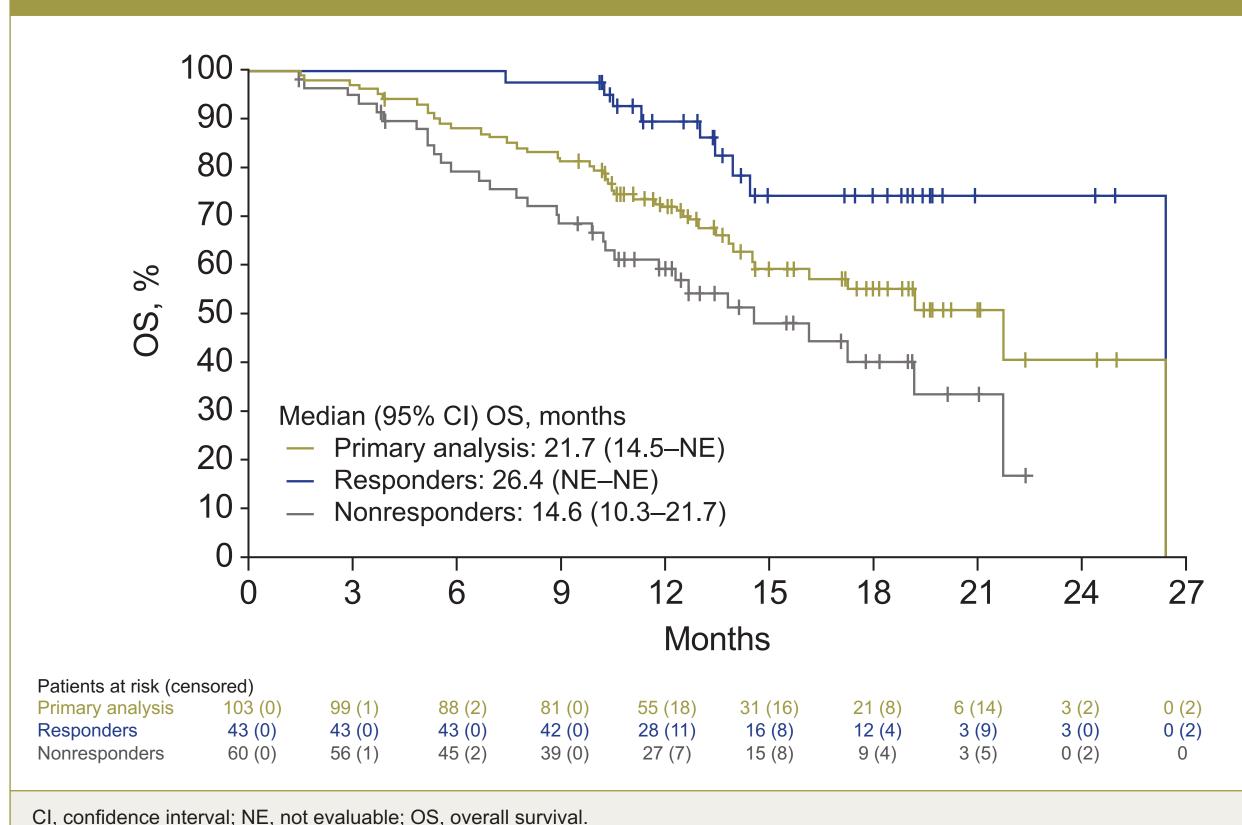


Figure 3. Overall survival

Genomic alterations

• While genomic subsets were small, an exploratory analysis showed that differences in co-occurring baseline genomic alterations between responders and nonresponders were varied and did not suggest a clear association with response (**Table 3**)

Table 3. Co-occurring baseline genomic alterations with >5% difference between responders and nonresponders

Patients with genomic alteration, n (%)	All treated patients (N=103)	Responders ^a (n=43)	Nonresponders (n=60)
BAP1	40 (38.8)	14 (32.6)	26 (43.3)
MLL2	11 (10.7)	7 (16.3)	4 (6.7)
PIK3C2B	11 (10.7)	7 (16.3)	4 (6.7)
IKBKE	10 (9.7)	6 (14.0)	4 (6.7)
MDM4	10 (9.7)	6 (14.0)	4 (6.7)
AKT3	9 (8.7)	8 (18.6)	1 (1.7)
CDC73	9 (8.7)	8 (18.6)	1 (1.7)
DDR2	9 (8.7)	7 (16.3)	2 (3.3)
PBRM1	9 (8.7)	2 (4.7)	7 (11.7)
BTG2	8 (7.8)	6 (14.0)	2 (3.3)
FH	8 (7.8)	7 (16.3)	1 (1.7)
RAD21	8 (7.8)	5 (11.6)	3 (5.0)
SDHC	8 (7.8)	5 (11.6)	3 (5.0)
H3F3A	7 (6.8)	5 (11.6)	2 (3.3)
CREBBP	6 (5.8)	4 (9.3)	2 (3.3)
NOTCH2	6 (5.8)	4 (9.3)	2 (3.3)
TET2	6 (5.8)	1 (2.3)	5 (8.3)
CALR	5 (4.9)	0 (0)	5 (8.3)
PARP1	5 (4.9)	4 (9.3)	1 (1.7)
JAK1	4 (3.9)	0 (0)	4 (6.7)
BCORL1	3 (2.9)	3 (7.0)	0 (0)
ERBB4	3 (2.9)	3 (7.0)	0 (0)
FANCC	3 (2.9)	3 (7.0)	0 (0)

^aPatients with confirmed partial or complete response based on independent central review.

CONCLUSIONS

- This post hoc analysis of the FOENIX-CCA2 study found that patients with a confirmed response to futibatinib had numerically longer PFS and OS versus nonresponders
- Although dose reductions and dose interruptions due to AEs were numerically more frequent in responders than nonresponders, they occurred later during treatment, suggesting a potential benefit of maintaining the starting dose for longer
- Further investigation of co-occurring genomic alterations as potential predictors of response to FGFR inhibitors is warranted

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