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Journal Journal of Clinical Oncology, 35(2)

ISSN 0732-183X

Authors

Nogova, Lucia Sequist, Lecia V Garcia, Jose Manuel Perez <u>et al.</u>

Publication Date

2017-01-10

DOI

10.1200/jco.2016.67.2048

Peer reviewed

JOURNAL OF CLINICAL ONCOLOGY

Evaluation of BGJ398, a Fibroblast Growth Factor Receptor 1-3 Kinase Inhibitor, in Patients With Advanced Solid Tumors Harboring Genetic Alterations in Fibroblast Growth Factor Receptors: Results of a Global Phase I, Dose-Escalation and Dose-Expansion Study

Lucia Nogova, Lecia V. Sequist, Jose Manuel Perez Garcia, Fabrice Andre, Jean-Pierre Delord, Manuel Hidalgo, Jan H.M. Schellens, Philippe A. Cassier, D. Ross Camidge, Martin Schuler, Ulka Vaishampayan, Howard A. Burris, G. Gary Tian, Mario Campone, Zev A. Wainberg, Wan-Teck Lim, Patricia LoRusso, Geoffrey I. Shapiro, Katie Parker, Xueying Chen, Somesh Choudhury, Francois Ringeisen, Diana Graus-Porta, Dale Porter, Randi Isaacs, Reinhard Buettner, and Jürgen Wolf

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Author affiliations appear at the end of this article

Published at ascopubs.org/journal/jco on November 21, 2016.

Support information appears at the end of this article.

Presented in part at the American Association for Cancer Research Annual Meeting, March 31-April 4, 2012, Chicago, IL; the American Association for Cancer Research Annual Meeting, April 5-9, 2014. San Diego, CA; and the American Society of Clinical Oncology Annual Meeting, May 30-June 3, 2014, Chicago, IL.

Clinical trial information: NCT01004224.

Corresponding author: Jürgen Wolf, MD, Lung Cancer Group Cologne, Department of Internal Medicine, Center for Integrated Oncology, University Hospital Cologne, D-50937 Cologne, Germany; e-mail: juergen.wolf@uk-koeln.de.

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0732-183X/17/3502w-157w/\$20.00

ASSOCIATED CONTENT

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Appendix

DOI: 10.1200/JCO.2016.67.2048

DOI: 10.1200/JCO.2016.67.2048

Purpose

This two-part, first-in-human study was initiated in patients with advanced solid tumors harboring genetic alterations in fibroblast growth factor receptors (FGFRs) to determine the maximum tolerated dose (MTD), the recommended phase II dose (RP2D), and the schedule, safety, pharmacokinetics, pharmacodynamics, and antitumor activity of oral BGJ398, a selective FGFR1-3 tyrosine kinase inhibitor.

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Patients and Methods

Adult patients were treated with escalating dosages of BGJ398 5 to 150 mg once daily or 50 mg twice daily continuously in 28-day cycles. During expansion at the MTD, patients with FGFR1amplified squamous cell non-small-cell lung cancer (sqNSCLC; arm 1) or other solid tumors with FGFR genetic alterations (mutations/amplifications/fusions) received BGJ398 daily on a continuous schedule (arm 2), or on a 3-weeks-on/1-week-off schedule (arm 3).

Results

Data in 132 patients from the escalation and expansion arms are reported (May 15, 2015, cutoff). The MTD, 125 mg daily, was determined on the basis of dose-limiting toxicities in four patients (100 mg, grade 3 aminotransferase elevations [n = 1]; 125 mg, hyperphosphatemia [n = 1]; 150 mg, grade 1 corneal toxicity [n = 1] and grade 3 aminotransferase elevations [n = 1]). Common adverse events in patients treated at the MTD (n = 57) included hyperphosphatemia (82.5%), constipation (50.9%), decreased appetite (45.6%), and stomatitis (45.6%). A similar safety profile was observed using the 3-weeks-on/1-week-off schedule (RP2D). However, adverse event-related dose adjustments/ interruptions were less frequent with the 3-weeks-on/1-week-off (50.0%) versus the continuous (73.7%) schedule. Antitumor activity (seven partial responses [six confirmed]) was demonstrated with BGJ398 doses \geq 100 mg in patients with FGFR1-amplified sqNSCLC and FGFR3-mutant bladder/urothelial cancer.

Conclusion

BGJ398 at the MTD/RP2D had a tolerable and manageable safety profile and showed antitumor activity in several tumor types, including FGFR1-amplified sqNSCLC and FGFR3-mutant bladder/ urothelial cancers.

J Clin Oncol 35:157-165. © 2016 by American Society of Clinical Oncology

INTRODUCTION

The discovery of targetable genomic aberrations underlying cancer has led to impressive improvements in the treatment of a subset of patients with

advanced cancers harboring such oncogenic drivers (eg, lung adenocarcinoma, melanoma).¹⁻³ However, novel personalized treatment strategies are missing in the vast majority of cancers.

Fibroblast growth factor receptor (FGFR) isoforms 1-4 and their 22 fibroblast growth factor ligands interact in a tissue-specific manner to initiate downstream signaling pathways that regulate cell proliferation, migration, differentiation, and survival—processes required for a variety of cell functions including angiogenesis and calcium/phosphate homeostasis.⁴⁻⁷ Genomic alterations in *FGFR1-3* (eg, gene amplifications, gain-of-function mutations, and chromosomal translocations) that trigger pathway activation^{4,5,8} have been identified in bladder cancer,^{9,10} squamous cell non–small-cell lung cancer (sqNSCLC),^{11,12} squamous cell cancer of the head and neck,¹³ endometrial cancer,^{8,14} cholangiocarcinoma,¹⁵ and breast cancer.^{4,16-18}

Although inhibitors targeting multiple receptor tyrosine kinases, including FGFR, are clinically active in several cancers,⁵ no FGFR-selective tyrosine kinase inhibitor (TKI) has been approved, and no TKI has been approved in a disease with a defined *FGFR* genetic alteration. BGJ398, an orally bioavailable, selective FGFR1 to 3 inhibitor (half maximal inhibitory concentration values range from 0.9 to 1.4 nM for FGFR1-3 to 60 nM for FGFR4),¹⁹ inhibits proliferation and tumor growth in preclinical cancer models bearing *FGFR1-3* genetic alterations.^{19,20}

On the basis of these preclinical data, we conducted a global, personalized phase I single-agent study to determine the maximum tolerated dose (MTD), recommended phase II dose (RP2D), schedule, safety, pharmacokinetics (PK), pharmacodynamics (PD), and antitumor activity of BGJ398 in patients with solid tumors bearing *FGFR* alterations (ClinicalTrials.gov identifier NCT01004224).

PATIENTS AND METHODS

Patients

Adults with solid tumors harboring *FGFR* alterations (eg, amplification, mutation, fusion) for whom no effective standard therapy exists were enrolled.

Patient selection criteria relative to *FGFR* genetic alterations are defined in the Appendix (online only). *FGFR* genetic alterations not specified in the protocol were compared with those in public databases (eg, COSMIC and dbSNP) and were adjudicated to determine suitability for enrollment, allowing for continual review and enrollment of patients with newly reported *FGFR* alterations suggestive of potential sensitivity to FGFR inhibition (Appendix Table A1, online only).

Patients with measurable/evaluable disease per Response Evaluation Criteria in Solid Tumors (RECIST) v1.0²¹ and a WHO performance status $\leq 2^{22}$ were eligible (Appendix). Patients with prior FGFR inhibitor (except TKI258) or MEK inhibitor treatment, a history or evidence of endocrine alteration of calcium/phosphate homeostasis or ectopic calcification/mineralization, or evidence of corneal disorder/keratopathy were excluded. Concomitant therapies increasing calcium/phosphate serum levels were not permitted.

Patient screening was conducted in seven countries at 18 study sites, with protocol and amendment approvals granted by each institution's review board/independent ethics committee. All applicable local regulations and principles of the Declaration of Helsinki were followed.²³ Patients provided written, informed consent before enrollment.

Study Design

The primary objective of this two-part phase I study was to determine the MTD, RP2D, and schedule of oral BGJ398. Secondary objectives included BGJ398 safety, tolerability, PK, PD, and antitumor activity. During dose escalation, sequential patient cohorts received BGJ398 5 (starting dose), 10, 20, 40, 60, 100, 125, and 150 mg once daily in continuous 28-day cycles (Fig 1). A twice daily 50-mg dose was also explored (after MTD/RP2D was established). At least three evaluable patients were treated in each cohort, and at least six patients were treated at the MTD/RP2D. During dose expansion, patients enrolled in one of three arms were treated at the MTD/RP2D: daily treatment of patients with *FGFR1*-amplified sqNSCLC (arm 1) or other *FGFR*-altered, advanced solid tumors (arm 2); and daily treatment on a 3-weeks-on/1-week-off schedule of patients with advanced solid tumors (excluding sqNSCLC; arm 3).

Dose-escalation decisions were guided by two adaptive Bayesian logistic regression models using the escalation-with-overdose-control principle, which estimated the rate of dose-limiting toxicities (DLTs), whether specific DLTs (Appendix) or adverse events (AEs)/laboratory abnormalities possibly related to BGJ398, resulting in failure to meet retreatment criteria. These models were reviewed, together with PK, PD, and safety assessments, and site investigator/personnel input to determine subsequent dose cohorts; intrapatient dose escalation was not allowed. The MTD was defined as the highest BGJ398 dose administered for ≥ 21 days resulting in DLTs in $\leq 33\%$ of patients during cycle 1. The probability of the dose with excessive toxicity (> 33%)was < 25%. DLT characterization and MTD determination were based on data from the dose-determining set, including patients who received the planned dose for ≥ 21 days in cycle 1 and were evaluated for safety for \geq 28 days after the first dose or who experienced a DLT during cycle 1.

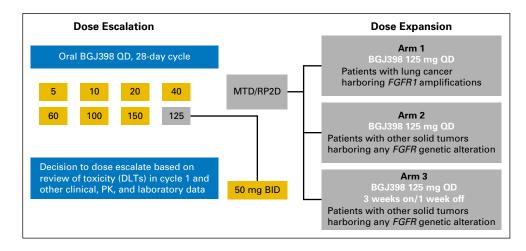


Fig 1. Study scheme. DLT, dose-limiting toxicity; MTD, maximum tolerated dose; PK, pharmacokinetics; RP2D, recommended phase II dose. BID, twice daily; QD, once daily.

Study Evaluations

Investigators assessed tumors at baseline and at every second treatment cycle until discontinuation (Appendix). Tumor evaluations were performed according to RECIST (version 1.0).²¹ A best overall response (BOR) (either a complete response or a partial response [PR]) required a response lasting \geq 4 weeks. A BOR of stable disease (SD) required a tumor assessment demonstrating SD for \geq 6 weeks after treatment initiation.

Safety was assessed using Common Terminology Criteria for Adverse Events (version 3),²⁴ with monitoring of AEs until 28 days after the final dose, and periodic physical examinations, laboratory evaluations, and electrocardiograms. FGFRs contribute to phosphate/vitamin D metabolism^{6,7}; thus, ophthalmologic examinations and assessments of calcium/phosphate homeostasis and renal function were performed (Appendix). PK and PD analyses are described in the Appendix.

Statistical Analyses

Descriptive statistics and/or contingency tables were used to summarize patient characteristics, efficacy and safety measurements, and PK. In a prespecified analysis, data from patients treated during dose escalation were used to determine the MTD/RP2D; subsequently, data were pooled with data from patients in the expansion phase who had the same dose and treatment schedule.

RESULTS

Patient Disposition and Baseline Characteristics

Data are reported from 132 heavily pretreated patients who started treatment between December 21, 2009, and April 7, 2015 (Table 1). At data cutoff (May 15, 2015), most patients (96.2%) had discontinued treatment, primarily because of disease progression (PD; 67.4%), AEs (12.9%), patient decision (10.6%), or death (3.8%; Appendix Table A2, online only).

Safety

The median duration of BGJ398 exposure was 7.1 weeks (range, 0.6 to 101.0 weeks; N = 132). Treatment-emergent AEs (TEAEs) were reported in 131 patients (99.2%), with most (95.5%) experiencing at least one AE suspected to be treatment related. Hyperphosphatemia (74.2%), constipation (40.2%), and decreased appetite (40.2%) were the most commonly reported TEAEs across all doses (Table 2). The most common AE suspected to be treatment related was asymptomatic hyperphosphatemia (72.7%); other frequent treatment-related AEs included stomatitis (36.4%), decreased appetite (28.8%), diarrhea (27.3%), fatigue (25.0%), alopecia (23.5%), and nausea (22.0%). Grade 3/4 TEAEs were reported in 69 patients (52.3%). Forty-one patients (31.1%) experienced a grade 3/4 AE suspected to be treatment related (Table 2). Seventy-eight patients (59.1%), primarily at \geq 100-mg doses, experienced an AE requiring dose adjustment or interruption. Eighteen patients (13.6%) had an AE leading to discontinuation (eye related in six patients [4.5%]). Fifteen patients died while receiving treatment or within 28 days of the last BGJ398 dose (Appendix Table A3, online only); 11 were attributed to PD or to related AEs. One death was suspected to be BGJ398 related (Appendix Table A3).

DLTs, experienced by four of the 34 patients (11.8%) in the dose-determining set, included grade 3 increases in ALT/AST (n = 1 each at 100 and 150 mg), hyperphosphatemia (n = 1, 125 mg),

Table 1. Patient Demographics and Disease History				
Baseline Characteristic	All Patients (N = 132)			
Age Median (range), years ≥ 65 years Sex, male	60 (25-86) 37 (28.0) 62 (47.0)			
WHO performance status 0 1 2	64 (48.5) 65 (49.2) 3 (2.3)			
Prior antineoplastic regimens 1 2 3 4 > 4	26 (19.7) 33 (25.0) 21 (15.9) 13 (9.8) 39 (29.5)			
Primary site of cancer Lung Breast Bladder/urothelial Colon Liver Head and neck Other*	48 (36.4) 43 (32.6) 12 (9.1) 5 (3.8) 3 (2.3) 3 (2.3) 18 (13.6)			
Dose level, mg once daily 5 10 20 40 60 100 125 125, 3 weeks on/1 week off 150 50†	$\begin{array}{c} 3 & (2.3) \\ 3 & (2.3) \\ 4 & (3.0) \\ 6 & (4.5) \\ 3 & (2.3) \\ 6 & (4.5) \\ 57 & (43.2) \\ 40 & (30.3) \\ 6 & (4.5) \\ 4 & (3) \end{array}$			

NOTE. Data are presented as No. (%) unless indicated otherwise. *Includes bone sarcoma, cervix, esophagus, gall bladder ducts, kidney, oral cavity, ovary, prostate, rectum, and other. †Twice daily.

and grade 1 corneal toxicity (n = 1, 150 mg). DLTs were reversible after BGJ398 interruption and/or concomitant medication. The BGJ398 MTD was determined as 125 mg once daily. Common AEs in patients treated at this dose and schedule (n = 57) included hyperphosphatemia (82.5%), constipation (50.9%), decreased appetite (45.6%), and stomatitis (45.6%). Dose adjustments/interruptions and AEs leading to discontinuations occurred in 73.7% and 10.5% of patients, respectively.

The DLT of hyperphosphatemia and the observation that most patients treated with doses ≥ 100 mg experienced AEs of hyperphosphatemia (Table 2) prompted the initiation of additional analyses to evaluate BGJ398 dose/schedule adjustment. Hyperphosphatemia was managed through dietary restrictions, phosphatelowering therapy, and drug interruptions. Earlier data from 43 patients treated at 125 mg once daily revealed a median time to first dose interruption of 22 days and a median duration of interruption of 7 days. Considering these data and the properties of BGJ398 PK, an intermittent 3-weeks-on/1-week-off schedule of 125 mg once daily was introduced as a dose-expansion arm in a protocol amendment (Appendix Table A1).

The safety profile of BGJ398 125 mg in patients treated on the intermittent schedule was similar to that observed in patients treated continuously. Fewer patients experienced AEs requiring dose adjustment/interruption (50%), but the rate of AEs leading to

	BGJ398 5-60 mg Once Dailv	BGJ398 100 mg Once Daily	BGJ398 125 mg Continuous Once Dailv	BGJ398 125 mg Once Daily 3 Weeks On/1 Week Off	BGJ398 150 mg Once	BGJ398 50 mg Twice Daily	All Patients (N = 132)	(N = 132)
AE	(n = 19)	(n = 6)	(n = 57)	(n = 40)	Daily $(n = 6)$	(n = 4)	All Grades	Grade 3/4
Any TEAE	19 (100)	6 (100)	57 (100)	39 (97.5)	6 (100)	4 (100)	131 (99.2)	69 (52.3)
Grade 3/4 treatment-related AEs	3 (15.8)	3 (50.0)	18 (31.6)	14 (35.0)	2 (33.3)	1 (25.0)	I	41 (31.1)
TEAEs leading to dose adjustment or interruption	4 (21.1)	5 (83.3)	42 (73.7)	20 (50.0)	4 (66.7)	3 (75.0)	78 (59.1)	I
TEAEs leading to treatment	4 (21.1)	2 (33.3)	6 (10.5)	5 (12.5)	1 (16.7)	0	18 (13.6)	I
Serious TEAEs	3 (15.8)	2 (33.3)	23 (40.4)	16 (40.0)	3 (50.0)	1 (25.0)	48 (36.4)	I
Eve-related TEAEs	6 (31.6)	4 (66.7)	27 (47.4)	19 (47.5)	2 (33.3)	2 (50.0)	60 (41.4)	4 (3.0)
TEAEs reported in $> 10\%$ of								
patients			i					:
Hyperphosphatemia	5 (26.3)	6 (100)	47 (82.5)	31 (77.5)	5 (83.3)	4 (100)	98 (74.2)	7 (5.3)
Constipation	1 (5.3)	3 (50.0)	29 (50.9)	18 (45.0)	0	2 (50.0)	53 (40.2)	1 (0.8)
Appetite decreased	2 (10.5)	3 (50.0)	26 (45.6)	17 (42.5)	3 (50.0)	2 (50.0)		5 (3.8)
Stomatitis	0	4 (66.7)	26 (45.6)	15 (37.5)	2 (33.3)	1 (25.0)		3 (2.3)
Diarrhea	7 (36.8)	5 (83.3)	16 (28.1)	14 (35.0)	1 (16.7)	3 (75.0)	46 (34.8)	0
Nausea	6 (31.6)	4 (66.7)	16 (28.1)	13 (32.5)	3 (50.0)	3 (75.0)	45 (34.1)	2 (1.5)
Fatigue	6 (31.6)	0	16 (28.1)	18 (45.0)	1 (16.7)	2 (50.0)	43 (32.6)	4 (3.0)
Alopecia	0	2 (33.3)	16 (28.1)	13 (32.5)	1 (16.7)	0	32 (24.2)	1 (0.8)
Creatinine increased	1 (5.3)	1 (16.7)	15 (26.3)	9 (22.5)	1 (16.7)	0	27 (20.5)	0
Dry mouth	4 (21.1)	1 (16.7)	11 (19.3)	10 (25.0)	0	1 (25.0)	27 (20.5)	0
Dyspnea	2 (10.5)	2 (33.3)	11 (19.3)	10 (25.0)	1 (16.7)	1 (25.0)	27 (20.5)	7 (5.3)
Vomiting	2 (10.5)	1 (16.7)	13 (22.8)	10 (25.0)	0	1 (25.0)	27 (20.5)	3 (2.3)
Asthenia	1 (5.3)	2 (33.3)	11 (19.3)	8 (20.0)	1 (16.7)	2 (50.0)	25 (18.9)	1 (0.8)
Dry eye	1 (5.3)	2 (33.3)	9 (15.8)	10 (25.0)	1 (16.7)	0	23 (17.4)	1 (0.8)
Dyspepsia	3 (15.8)	2 (33.3)	9 (15.8)	5 (12.5)	2 (33.3)	2 (50.0)	23 (17.4)	0
Alanine aminotransferase	2 (10.5)	1 (16.7)	12 (21.1)	5 (12.5)	1 (16.7)	0	21 (15.9)	9 (6.8)
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Asuartate aminotransferase	2 (10.5) 2 (10.5)	1 (16.7)	0 (14.0) 12 (21 1)	0 (13.0) 4 (10.0)	U 1 (16 7)	0.0.010	20 (15.2)	6 (4 5)
increased	10.011 2		11.12/21	0.01 +		þ	12:01/02	().t)
Dysgeusia	1 (5.3)	3 (50.0)	9 (15.8)	6 (15.0)		0	19 (14.4)	0
Back pain	4 (21.1)	0	5 (8.8)	5 (12.5)	2 (33.3)	2 (50.0)	18 (13.6)	0
Pyrexia	3 (15.8)	1 (16.7)	7 (12.3)	7 (17.5)	0	0	18 (13.6)	1 (0.8)
Anemia	1 (5.3)	2 (33.3)	6 (10.5)	8 (20.0)	0	0	17 (12.9)	0
Cough	3 (15.8)	1 (16.7)	6 (10.5)	5 (12.5)	1 (16.7)	0	16 (12.1)	0
Epistaxis	0	1 (16.7)	9 (15.8)	4 (10.0)	1 (16.7)	0	15 (11.4)	0
Lipase increased	3 (15.8)	1 (16.7)	6 (10.5)	4 (10.0)	0	1 (25.0)	15 (11.4)	7 (5.3)
Arthralgia	0	1 (16.7)	6 (10.5)	5 (12.5)	1 (16.7)	1 (25.0)	14 (10.6)	0
Hypercalcemia	2 (10.5)	0	6 (10.5)		0	1 (25.0)	14 (10.6)	3 (2.3)
Weight decreased	0	0	10 (17.5)	3 (7.5)	1 (16.7)	0	14 (10.6)	0

discontinuation (12.5%) and the most common AEs were similar in the two schedules (Table 2). On the basis of these data, the RP2D for BGJ398 was chosen as 125 mg once daily administered in the 3-weeks-on/1-week-off schedule. With this regimen, the majority of patients achieved exposures above the threshold associated with preclinical evidence of FGFR pathway inhibition and in vivo efficacy (Fig 2).²⁰

РК

BGJ398 mean plasma concentration time profiles and area under the plasma concentration-time curve by cohort, starting at dosages \geq 20 mg once daily, are presented in Fig 2. Estimated PK parameters are reported in Appendix Table A4 (online only). Plasma concentrations for BGJ398 at dosages of 5 mg and 10 mg once daily were frequently below the lower limit of quantification (data not shown).

The median time to reach maximum plasma concentration after a single dose was approximately 2 to 3 hours. The mean area under the plasma concentration-time curve from 0 to 24 hours on day 1 increased by approximately nine-fold, from 20 to 150 mg once daily. Despite the relatively short median terminal elimination half-life on day 1 for these doses (range, 2.69 to 5.90 hours), accumulation was observed with dosing of \geq 60 mg once daily;

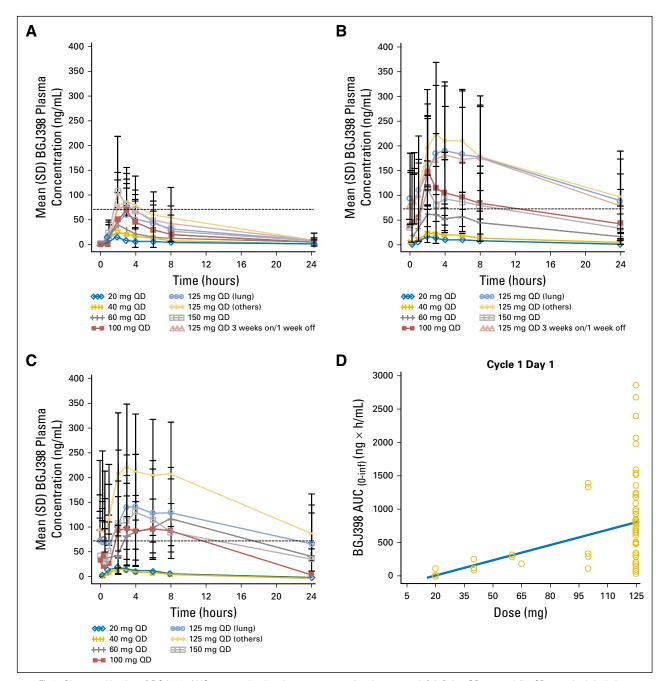


Fig 2. Pharmacokinetics of BGJ398. AUC, area under the plasma concentration-time curve; inf, infinity; QD, once daily; SD, standard deviation.

mean accumulation ratios ranged from 3.78 to 6.60 (day 15) and from 2.99 to 7.86 (day 28). After 125 mg once daily dosing, the unbound average steady-state BGJ398 concentration on day 28 of cycle 1 was 6.93 nM. Because dose interruptions occurred frequently after continuous once daily dosing, PK parameters on day 28 should be interpreted with caution.

Clinical Activity

Among all the patients treated with BGJ398 (N = 132), 42 had a BOR of SD, six achieved PR, and one achieved an unconfirmed PR (confirmatory scan performed 1 day early). Among the 85 patients treated at \geq 100 mg with evaluable data, 28 (32.9%) had reduced tumor burden assessed as the best percentage change from baseline in the sum of the longest diameter in the target lesion or lesions (Fig 3).

The disease control rate (DCR: PR + SD) in 36 patients with *FGFR1*-amplified sqNSCLC treated at doses of \geq 100 mg continuously was 50%; four patients (11.1%) achieved PRs (100 mg [n = 1], 125 mg [n = 3, 2 confirmed]) and 14 patients had SD. In the subset of 31 patients treated at 125 mg continuously, 12 continued to receive treatment for > 8 weeks, and one half of these patients continued to receive therapy for \geq 16 weeks (Fig 4).

Twenty-seven of the 36 patients treated at \geq 100 mg had preand post-treatment target lesion assessments; of these, 11 (40.7%) had reduced tumor burden (Fig 3; Appendix Table A5, online only). The responders remained on study for 39.9 to 76.6 weeks (confirmed PRs) and for 26.3 weeks (unconfirmed PR).

The DCR in eight patients with *FGFR3*-mutated bladder/ urothelial cancer treated at doses \geq 100 mg was 75%; three patients (37.5%) achieved PRs (125 mg continuously [n = 1], 125 mg 3-weeks-on/1-week-off [n = 2]) and three patients had SD. Five patients (62.5%) had reduced tumor burden and two of three with SD had reductions in tumor measurements nearing PR (27% to 28%; Fig 3). Of the six patients with disease control, the time receiving treatment ranged from 15.1 to \geq 101 weeks, with one ongoing at data cutoff (Fig 4). One patient withdrew after 2 weeks; this patient had PD in an assessment performed 25 days after the last dose. Of note, one patient with *FGFR1*-amplified bladder cancer treated at 125 mg progressed rapidly in cycle 1 after 11 days of treatment.

Although no other PRs were observed at doses \geq 100 mg, 10 of 32 patients (31%) with breast cancer treated at \geq 100 mg had a best response of SD. Of 26 patients with breast cancer (*FGFR1/2* amplified [n = 25]; *FGFR3* mutant [n = 1]) with pre- and post-treatment target lesion measurements, four (15.4%) had reduced tumor burden. In addition, all three patients with *FGFR2*-altered (fusion [n = 2] or mutation [n = 1]) cholangiocarcinoma with pre- and post-treatment target lesion assessments had reduced tumor burden (Fig 3).

PD

BGJ398 treatment led to increased serum phosphate levels (Appendix Fig A1A, online only), as well as to dose- and exposurerelated hyperphosphatemia, in most patients treated at doses \geq 100 mg (Table 2). The median percentage change in FGF23 plasma levels ranged from approximately -25% to 80%, with a trend toward greater increases in patients treated at higher doses (Appendix Fig A1B).

DISCUSSION

This global first-in-human study of the FGFR1-3 inhibitor BGJ398 demonstrated a tolerable safety profile in patients with advanced

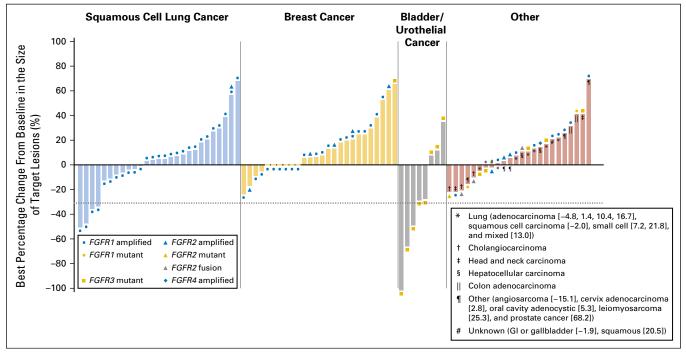


Fig 3. Waterfall plots of best change from baseline in the size of target lesions for patients treated at \geq 100 mg BGJ398.

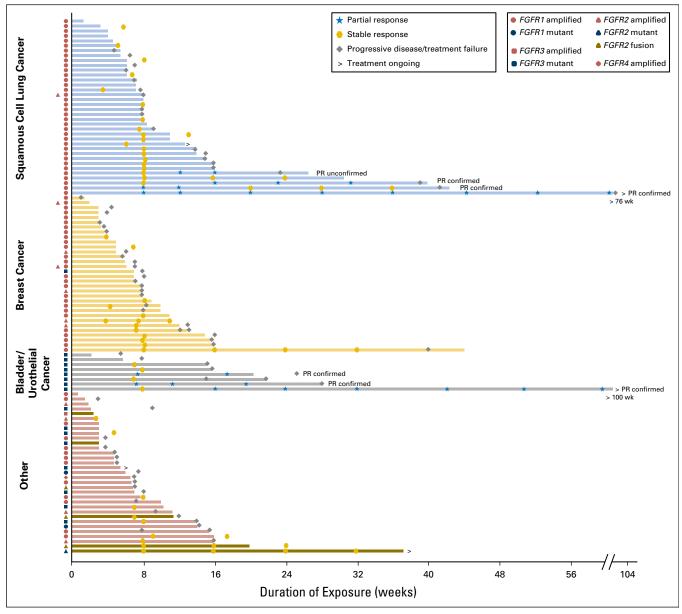


Fig 4. Response and duration of exposure in patients treated at BGJ398 doses \geq 100 mg. Among the patients with other cancer types were five patients with cholangiocarcinoma (green bars). PR, partial response.

solid tumors bearing *FGFR* amplifications, mutations, or fusions. The BGJ398 MTD/RP2D was determined to be 125 mg daily; the recommended schedule at the RP2D was 3 weeks on/1 week off on the basis of DLTs observed in four patients treated at doses \geq 100 mg, BGJ398 PK, and safety data in patients treated on the intermittent schedule. This BGJ398 dose resulted in an average unbound steady-state concentration of 6.93 nM, which is similar to the half maximal inhibitory concentration of BGJ398 in the RT112 human tumor-derived *FGFR* gene fusion bladder cancer cell line (4 nM).²⁵ Commonly reported AEs included hyperphosphatemia, constipation, decreased appetite, and stomatitis. A similar safety profile was observed with the pan-FGFR inhibitor JNJ-42756493.²⁶

The FGFR pathway plays an important role in FGF23-mediated phosphate homeostasis.^{6,7} Accordingly, hyperphosphatemia, the

most common AE, was experienced by most patients receiving $\geq 100 \text{ mg BGJ398.}^{27}$ An intermittent dosing arm (125 mg once daily 3 weeks on/1 week off) was opened during dose expansion to manage hyperphosphatemia-related interruptions, resulting in fewer patients requiring an AE-related dose adjustment/interruption (50.0% v 73.7% [continuous]). Importantly, objective responses and reductions in tumor measurements were observed with both schedules, with similar overall safety profiles. Further exploration of BGJ398 dosing may be performed in the context of ongoing or future trials.

Because the FGF23/FGFR pathway mediates renal tubular phosphate secretion,⁷ FGF23 levels were used initially as a biomarker for FGFR pathway inhibition. Median FGF23 levels increased by \leq 80% in patients treated with BGJ398, with generally higher levels at higher doses. Elevated serum phosphate levels,

consistently observed with BGJ398 doses \geq 100 mg, proved to be a sensitive, dose-dependent PD indicator of on-target activity and a key biomarker for FGFR pathway inhibition.

BGJ398 treatment provided disease control in 49 of 132 patients across all doses, with PRs observed in patients with FGFR1-amplified sqNSCLC (n = 4) and FGFR3-mutant bladder/ urothelial cancer (n = 3). The clinical activity observed in *FGFR1*amplified sqNSCLC (11% overall response rate, 50% DCR) is notable given the unfavorable prognosis for patients who relapse after chemotherapy.²⁸ Moreover, the duration on study for responding patients (6.1 to 17.7 months) is comparable to that achieved with nivolumab, an anti-PD-1 antibody approved in metastatic sqNSCLC after failure of platinum-based chemotherapy.²⁹⁻³² To date, no oncogenic driver-targeted drug has shown activity as a monotherapy, raising the possibility of a molecularly definable sqNSCLC subtype sensitive to FGFR inhibition. Despite preselecting for FGFR1 amplification, the response rate was lower than expected on the basis of preclinical data, suggesting that FGFR1 amplification may not function as a sole biomarker predicting clinical benefit.¹¹ Although the basis for this discrepancy (eg, tumor heterogeneity, different amplification parameters, and/ or presence of required cofactors)^{33,34} remains unknown, BGJ398 use as a personalized treatment approach in sqNSCLC warrants further investigation. Future whole genome analyses may help define the exploitable molecular differences between responders and nonresponders,³⁴ providing further insight into additional oncogenic drivers in saNSCLC.

Responses observed in *FGFR3*-mutant bladder/urothelial cancer after failure of platinum-based chemotherapy (38% overall response rate, 75% DCR) strongly support a role for *FGFR3* mutations as driver alterations in this molecular subgroup and the potent inhibitory function of BGJ398. In light of the < 1-year overall survival for patients with metastatic urothelial/bladder cancer after relapse after first-line chemotherapy,³⁵ BGJ398-targeted treatment warrants further investigation. To this end, a fourth expansion arm was opened to further evaluate BGJ398 activity in patients with urothelial cancer harboring an *FGFR3* mutation or fusion.

SD with reduced tumor burden was also observed in patients with cholangiocarcinoma (*FGFR2* fusions [n = 2], *FGFR2* mutation [n = 1]) and *FGFR1*-amplified squamous head and neck cancer. The disease control observed in cholangiocarcinoma is notable given the limited treatment options available for patients who progress after chemotherapy.^{36,37} Of interest, another patient with cholangiocarcinoma who was enrolled with a presumed *FGFR3* mutation progressed rapidly and was later identified to be wild type for *FGFR* but as having a mutation in *KRAS*, a negative predictor (preclinically) for BGJ398 sensitivity.²⁰ A phase II study exploring BGJ398 as a second-line or later therapy in patients with *FGFR2*-altered advanced/metastatic cholangiocarcinoma is ongoing (ClinicalTrials.gov identifier NCT02150967).

The lack of objective responses and the limited disease control observed with BGJ398 in patients with breast cancer challenge the idea of *FGFR* amplification as a sole oncogenic driver in this disease; however, BGJ398 may prove more effective against advanced breast cancer when combined with other endocrine or targeted agents. Certain *FGFR* alterations (eg, *FGFR3* mutations/gene fusions in bladder/urothelial carcinoma and *FGFR2* gene

fusions in cholangiocarcinoma) are dominant oncogenic drivers and confer sensitivity to BGJ398-mediated FGFR inhibition, whereas *FGFR1* amplification, observed in a number of tumor types including sqNSCLC and breast cancer, may not be sufficient to identify a BGJ398-sensitive population. It is unclear whether alternative or additional biomarkers (eg, FGFR1 protein levels or FGFR pathway activity) would better predict responders, or whether FGFR signaling is less essential for tumor growth in breast cancer.

When this study was initiated in 2009, predictors of FGFR inhibitor sensitivity were limited. As the knowledge of FGFR biology and driver genetic alterations increased and assays for patient selection became available over the 6-year enrollment period, patient inclusion criteria were amended accordingly (Appendix Table A1). This study allowed the clinical evaluation of multiple preclinical hypotheses related to BGJ398-mediated FGFR pathway sensitivity and established which patient populations were likely to benefit from treatment with an FGFR-selective inhibitor.

Taken together, treatment with BGJ398 in patients with advanced solid tumors bearing *FGFR* alterations was tolerable, with manageable toxicity. The MTD/RP2D was determined to be 125 mg once daily on a 3-weeks-on/1-week-off schedule. BGJ398 demonstrated antitumor activity in *FGFR1*-amplified sqNSCLC, *FGFR3*-mutant bladder/urothelial cancer, and *FGFR2*-gene fusion/ mutant cholangiocarcinoma, strongly supporting further biologic and clinical investigation. BGJ398 clinical development is ongoing, including adding a fourth expansion arm to this study for patients with urothelial carcinoma and *FGFR3* mutation/gene fusion and a phase II study in cholangiocarcinoma with *FGFR2* gene fusion/ other *FGFR* genetic alterations.^{38,39}

AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

Disclosures provided by the authors are available with this article at ascopubs.org/journal/jco.

AUTHOR CONTRIBUTIONS

Conception and design: Jose Manuel Perez Garcia, Manuel Hidalgo, Jan H.M. Schellens, D. Ross Camidge, Howard A. Burris, Somesh Choudhury, Francois Ringeisen, Randi Isaacs, Reinhard Buettner, Jürgen Wolf **Administrative support:** Martin Schuler

Provision of study materials or patients: Lucia Nogova; Lecia V. Sequist, Manuel Hidalgo, D. Ross Camidge, Martin Schuler, Ulka Vaishampayan, Howard A. Burris, G. Gary Tian, Geoffrey I. Shapiro, Jürgen Wolf

Collection and assembly of data: Lucia Nogova, Lecia V. Sequist, Jean-Pierre Delord, Jan H.M. Schellens, D. Ross Camidge, Martin Schuler, Ulka Vaishampayan, Howard A. Burris, G. Gary Tian, Mario Campone,

Wan-Teck Lim, Patricia LoRusso, Geoffrey I. Shapiro, Katie Parker, Somesh Choudhury, Francois Ringeisen, Randi Isaacs, Reinhard Buettner, Jürgen Wolf

Data analysis and interpretation: Lucia Nogova, Lecia V. Sequist, Fabrice Andre, Manuel Hidalgo, Jan H.M. Schellens, Philippe A. Cassier, D. Ross Camidge, Ulka Vaishampayan, Howard A. Burris, Zev A. Wainberg, Wan-Teck Lim, Patricia LoRusso, Geoffrey I. Shapiro, Katie Parker, Xueying Chen, Somesh Choudhury, Francois Ringeisen, Diana Graus-Porta, Dale Porter, Randi Isaacs, Reinhard Buettner, Jürgen Wolf Manuscript writing: All authors

Final approval of manuscript: All authors

REFERENCES

1. Kris MG, Johnson BE, Berry LD, et al: Using multiplexed assays of oncogenic drivers in lung cancers to select targeted drugs. JAMA 311:1998-2006, 2014

 Clinical Lung Cancer Genome Project (CLCGP); Network Genomic Medicine (NGM): A genomics-based classification of human lung tumors. Sci Transl Med 5: 209ra153, 2013

 Muñoz-Couselo E, Garcia JS, Pérez-Garcia JM, et al: Recent advances in the treatment of melanoma with BRAF and MEK inhibitors. Ann Transl Med 3:207, 2015

4. Turner N, Grose R: Fibroblast growth factor signalling: From development to cancer. Nat Rev Cancer 10:116-129, 2010

 Helsten T, Schwaederle M, Kurzrock R: Fibroblast growth factor receptor signaling in hereditary and neoplastic disease: Biologic and clinical implications. Cancer Metastasis Rev 34:479-496, 2015

6. Perwad F, Zhang MY, Tenenhouse HS, et al: Fibroblast growth factor 23 impairs phosphorus and vitamin D metabolism in vivo and suppresses 25-hydroxyvitamin D-1alpha-hydroxylase expression in vitro. Am J Physiol Renal Physiol 293:F1577-F1583, 2007

7. Gattineni J, Bates C, Twombley K, et al: FGF23 decreases renal NaPi-2a and NaPi-2c expression and induces hypophosphatemia in vivo predominantly via FGF receptor 1. Am J Physiol Renal Physiol 297: F282-F291, 2009

8. Dienstmann R, Rodon J, Prat A, et al: Genomic aberrations in the FGFR pathway: Opportunities for targeted therapies in solid tumors. Ann Oncol 25: 552-563, 2014

9. Cappellen D, De Oliveira C, Ricol D, et al: Frequent activating mutations of FGFR3 in human bladder and cervix carcinomas. Nat Genet 23:18-20, 1999

10. Williams SV, Hurst CD, Knowles MA: Oncogenic FGFR3 gene fusions in bladder cancer. Hum Mol Genet 22:795-803, 2013

11. Weiss J, Sos ML, Seidel D, et al: Frequent and focal FGFR1 amplification associates with therapeutically tractable FGFR1 dependency in squamous cell lung cancer. Sci Transl Med 2:62ra93, 2010

12. Dutt A, Ramos AH, Hammerman PS, et al: Inhibitor-sensitive FGFR1 amplification in human non-small cell lung cancer. PLoS One 6:e20351, 2011

 Freier K, Schwaenen C, Sticht C, et al: Recurrent FGFR1 amplification and high FGFR1 protein expression in oral squamous cell carcinoma (OSCC). Oral Oncol 43:60-66, 2007

14. Dutt A, Salvesen HB, Chen TH, et al: Drugsensitive FGFR2 mutations in endometrial carcinoma. Proc Natl Acad Sci U S A 105: 8713-8717, 2008

15. Arai Y, Totoki Y, Hosoda F, et al: Fibroblast growth factor receptor 2 tyrosine kinase fusions define a unique molecular subtype of cholangiocarcinoma. Hepatology 59:1427-1434, 2014

 Courjal F, Cuny M, Simony-Lafontaine J, et al: Mapping of DNA amplifications at 15 chromosomal localizations in 1875 breast tumors: Definition of phenotypic groups. Cancer Res 57:4360-4367, 1997

17. Jacquemier J, Adelaide J, Parc P, et al: Expression of the FGFR1 gene in human breastcarcinoma cells. Int J Cancer 59:373-378, 1994

 Turner N, Lambros MB, Horlings HM, et al: Integrative molecular profiling of triple negative breast cancers identifies amplicon drivers and potential therapeutic targets. Oncogene 29:2013-2023, 2010

19. Guagnano V, Furet P, Spanka C, et al: Discovery of 3-(2,6-dichloro-3,5-dimethoxy-phenyl)-1-{6-[4-(4-ethyl-piperazin-1-yl)-phenylamino]-pyrimidin-4yl)-1-methyl-urea (NVP-BGJ398), a potent and selective inhibitor of the fibroblast growth factor receptor family of receptor tyrosine kinase. J Med Chem 54:7066-7083, 2011

20. Guagnano V, Kauffmann A, Wöhrle S, et al: FGFR genetic alterations predict for sensitivity to NVP-BGJ398, a selective pan-FGFR inhibitor. Cancer Discov 2:1118-1133, 2012

21. Therasse P, Arbuck SG, Eisenhauer EA, et al: New guidelines to evaluate the response to treatment in solid tumors. European Organization for Research and Treatment of Cancer, National Cancer Institute of the United States, National Cancer Institute of Canada. J Natl Cancer Inst 92:205-216, 2000

22. World Health Organization: WHO Handbook for Reporting Results for Cancer Treatment. Geneva, Switzerland, World Health Organization, 1979

23. World Medical Association. Declaration of Helsinki. Ethical Principles for Medical Research Involving Human Subjects. Helsinki, Finland, World Medical Association, 1964

24. Trotti A, Colevas AD, Setser A, et al: CTCAE v3.0: Development of a comprehensive grading system for the adverse effects of cancer treatment. Semin Radiat Oncol 13:176-181, 2003

25. Graus-Porta D: BGJ398, a pan-FGFR inhibitor for the treatment of FGFR genetically altered cancers and beyond. Presented at Mildred Scheel Cancer Conference, Königswinter, Germany, June 5-7, 2013

26. Tabernero J, Bahleda R, Dienstmann R, et al: Phase I dose-escalation study of JNJ-42756493, an oral pan-fibroblast growth factor receptor inhibitor, in patients with advanced solid tumors. J Clin Oncol 33: 3401-3408, 2015 27. Sequist LV, Cassier P, Varga A, et al: Phase I study of BGJ398, a selective pan-FGFR inhibitor in genetically preselected advanced solid tumors. Cancer Res 74, 2014 (suppl; abstr CT326)

28. Shepherd FA, Dancey J, Ramlau R, et al: Prospective randomized trial of docetaxel versus best supportive care in patients with non-small-cell lung cancer previously treated with platinum-based chemotherapy. J Clin Oncol 18:2095-2103, 2000

29. Brahmer J, Reckamp KL, Baas P, et al: Nivolumab versus docetaxel in advanced squamous-cell nonsmall-cell lung cancer. N Engl J Med 373:123-135, 2015

30. Rizvi NA, Mazières J, Planchard D, et al: Activity and safety of nivolumab, an anti-PD-1 immune checkpoint inhibitor, for patients with advanced, refractory squamous non-small-cell lung cancer (CheckMate 063): A phase 2, single-arm trial. Lancet Oncol 16:257-265, 2015

31. Bustamante Alvarez JG, González-Cao M, Karachaliou N, et al: Advances in immunotherapy for treatment of lung cancer. Cancer Biol Med 12:209-222, 2015

32. Opdivo (nivolumab) [package insert]. Princeton, NJ, Bristol-Myers Squibb, 2015. http://www. accessdata.fda.gov/drugsatfda_docs/label/2015/ 125527s000lbl.pdf

33. Malchers F, Dietlein F, Schöttle J, et al: Cellautonomous and non-cell-autonomous mechanisms of transformation by amplified FGFR1 in lung cancer. Cancer Discov 4:246-257, 2014

34. Wynes MW, Hinz TK, Gao D, et al: FGFR1 mRNA and protein expression, not gene copy number, predict FGFR TKI sensitivity across all lung cancer histologies. Clin Cancer Res 20:3299-3309, 2014

35. Yafi FA, North S, Kassouf W: First- and secondline therapy for metastatic urothelial carcinoma of the bladder. Curr Oncol 18:e25-e34, 2011

36. Ang C: Role of the fibroblast growth factor receptor axis in cholangiocarcinoma. J Gastroenterol Hepatol 30:1116-1122, 2015

37. National Comprehensive Cancer Network: NCCN clinical practice guidelines in oncology. Hepatobiliary cancers, v2.2016. https://www.nccn.org/ professionals/physician_gls/pdf/hepatobiliary.pdf

38. Javle MM, Shroff RT, Zhu A, et al: A phase 2 study of BGJ398 in patients (pts) with advanced or metastatic FGFR-altered cholangiocarcinoma (CCA) who failed or are intolerant to platinum-based chemotherapy. J Clin Oncol 34, 2016 (suppl; abstr 335)

39. Pal SK, Rosenberg JE, Keam B, et al: Efficacy of BGJ398, a fibroblast growth factor receptor (FGFR) 1-3 inhibitor, in patients (pts) with previously treated advanced/ metastatic urothelial carcinoma (mUC) with FGFR3 alterations. J Clin Oncol 34, 2016 (suppl; abstr 4517)

Affiliations

Lucia Nogova, Reinhard Buettner, and Jürgen Wolf, University Hospital Cologne, Cologne; Martin Schuler, University Hospital Essen, and German Cancer Consortium (DKTK), Essen, Germany; Lecia V. Sequist, Massachusetts General Hospital; Geoffrey I. Shapiro, Dana-Farber Cancer Institute, Boston, MA; Jose Manuel Perez Garcia, Vall d'Hebron Institut d'Oncologia, Barcelona; Manuel Hidalgo, START Madrid; Manuel Hidalgo, Spanish National Cancer Research Centre, Madrid, Spain; Fabrice Andre, Gustave Roussy, Villejuif; Jean-Pierre Delord, Institute Universitaire du Cancer de Toulouse, Toulouse; Philippe A. Cassier, Centre Regional Leon-Berard, Lyon; Mario Campone, Institute de Cancerologie de l'Ouest-René Gauducheau, Nantes, France; Jan H.M. Schellens, Netherlands Cancer Institute, Amsterdam, and Utrecht University, Utrecht, the Netherlands; D. Ross Camidge, University of Colorado, Aurora, CO; Ulka Vaishampayan, Barbara Ann Karmanos Cancer Institute, Detroit, MI; Howard A. Burris, Sarah Cannon Research Institute, Nashville; G. Gary Tian, The West Clinic, Memphis, TN; Zev A. Wainberg, UCLA School of Medicine, Los Angeles, CA; Wan-Teck Lim, National Cancer Centre, Singapore; Patricia LoRusso, Yale Cancer Center, New Haven, CT; Katie Parker, Xueying Chen, Somesh Choudhury, and Randi Isaacs, Novartis Pharmaceuticals Corporation, East Hanover, NJ; Dale Porter, Novartis Institutes for Biomedical Research, Cambridge, MA; Francois Ringeisen, Novartis Pharma AG; and Diana Graus-Porta, Novartis Institutes for Biomedical Research, Basel, Switzerland.

Support

Supported by Novartis Pharmaceuticals.

AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

Evaluation of BGJ398, a Fibroblast Growth Factor Receptor 1-3 Kinase Inhibitor, in Patients With Advanced Solid Tumors Harboring Genetic Alterations in Fibroblast Growth Factor Receptors: Results of a Global Phase I, Dose-Escalation and Dose-Expansion Study

The following represents disclosure information provided by authors of this manuscript. All relationships are considered compensated. Relationships are self-held unless noted. I = Immediate Family Member, Inst = My Institution. Relationships may not relate to the subject matter of this manuscript. For more information about ASCO's conflict of interest policy, please refer to www.asco.org/rwc or ascopubs.org/jco/site/ifc.

Lucia Nogova

Honoraria: Pfizer Consulting or Advisory Role: Boehringer Ingelheim, Roche Research Funding: Pfizer Travel, Accommodations, Expenses: Novartis, Pfizer

Lecia V. Sequist

Honoraria: AstraZeneca

Consulting or Advisory Role: Clovis Oncology, Novartis, Merrimack, AstraZeneca, ARIAD Pharmaceuticals, Genentech

Research Funding: Boehringer Ingelheim (Inst), Clovis Oncology (Inst), Genentech (Inst), Merrimack (Inst), ArQule (Inst), Novartis (Inst), AstraZeneca (Inst), Johnson & Johnson (Inst), Eli Lilly (Inst), Merck (Inst), Taiho Pharmaceutical (Inst)

Jose Manuel Perez Garcia

No relationship to disclose

Fabrice Andre

Research Funding: AstraZeneca (Inst), Novartis (Inst), Pfizer (Inst), Eli Lilly (Inst)

Jean-Pierre Delord No relationship to disclose

Manuel Hidalgo

Stock or Other Ownership: Champions Oncology
Honoraria: Pfizer, Novartis, Merck Sharp & Dohme, Celgene
Consulting or Advisory Role: Novartis, Pfizer, Celgene, Merck, Eli Lilly/ ImClone Systems
Research Funding: Pfizer, Celgene, Merck Serono

Travel, Accommodations, Expenses: Pfizer, Celgene, Merck

Jan H.M. Schellens No relationship to disclose

Philippe A. Cassier Honoraria: Novartis Research Funding: Novartis (Inst) Travel, Accommodations, Expenses: PharmaMar

D. Ross Camidge

Honoraria: Novartis

Martin Schuler

Honoraria: Alexion Pharmaceuticals, Boehringer Ingelheim, Celgene, Eli Lilly, Novartis, Pfizer, Bristol-Myers Squibb

Consulting or Advisory Role: AstraZeneca, Boehringer Ingelheim, Bristol-Myers Squibb, Celgene, Eli Lilly, Novartis

Research Funding: Boehringer Ingelheim (Inst), Bristol-Myers Squibb (Inst), Novartis (Inst)

Patents, Royalties, Other Intellectual Property: University of Duisburg-Essen

Travel, Accommodations, Expenses: Merck Sharp & Dohme

Ulka Vaishampayan

Honoraria: Astellas Pharma, Pfizer, Janssen Pharmaceuticals, Novartis, Bayer AG, Sanofi, Bristol-Myers Squibb

Consulting or Advisory Role: Pfizer, Novartis, Astellas Pharma, Sanofi, Bristol-Myers Squibb

Speakers' Bureau: Pfizer, Novartis, Astellas Pharma, Bayer AG, Bristol-Myers Squibb

Research Funding: Astellas Pharma, Novartis, Exelixis, Pfizer

Howard A. Burris

Employment: HCA Healthcare/Sarah Cannon **Leadership:** HCA Healthcare/Sarah Cannon

Stock and Other Ownership Interests: HCA Healthcare/Sarah Cannon Consulting or Advisory Role: Mersana Therapeutics (Inst), AstraZeneca (Inst), FORMA Therapeutics (Inst), Janssen (Inst), Novartis (Inst), Roche/ Genentech (Inst), TG Therapeutics (Inst), MedImmune (Inst), Bristol-Myers Squibb (Inst)

Research Funding: Roche/Genentech (Inst), Bristol-Myers Squibb (Inst), Incyte (Inst), Tarveda Therapeutics (Inst), Mersana Therapeutics (Inst), AstraZeneca (Inst), MedImmune (Inst), Macrogenics (Inst), Novartis (Inst), Boehringer Ingelheim (Inst), Eli Lilly (Inst), Seattle Genetics (Inst), AbbVie (Inst), Bayer (Inst), Celldex Therapeutics (Inst), Merck (Inst), Celgene (Inst), Agios Pharmaceuticals (Inst), Jounce Therapeutics (Inst), Moderna Therapeutics (Inst), CytomX Therapeutics (Inst), GlaxoSmithKline (Inst), Verastem (Inst), Tesaro (Inst), Immunocore (Inst), Takeda (Inst), Millennium (Inst), BioMed Valley Discoveries (Inst), Pfizer (Inst), PTC Therapeutics (Inst), TG Therapeutics (Inst), Loxo (Inst), Vertex (Inst), eFFECTOR Therapeutics (Inst), Janssen (Inst), Gilead Sciences (Inst), Valent Technologies (Inst), BioAtla (Inst), CicloMed (Inst), Harpoon Therapeutics (Inst), Jiangsu Hengrui Medicine (Inst), Revolution Medicines (Inst), Daiichi Sankyo (Inst), H3 Biomedicine (Inst), Neon Therapeutics (Inst), OncoMed (Inst), Regeneron (Inst), Sanofi (Inst) Expert Testimony: Novartis

G. Gary Tian

Consulting or Advisory Role: Cardinal Health, Merrimack **Travel, Accommodations, Expenses:** Cardinal Health

Mario Campone

Honoraria: Novartis, Servier, Menarini Consulting or Advisory Role: Novartis, Servier, Menarini, Cellgen Speakers' Bureau: Novartis, Amgen Research Funding: Novartis (Inst) Travel, Accommodations, Expenses: Novartis

Zev A. Wainberg

Consulting or Advisory Role: Taiho Pharmaceutical, Sirtex Medical, Amgen Speakers' Bureau: Genentech Research Funding: Novartis (Inst), Plexxikon (Inst), Pfizer (Inst), Biomarin (Inst), Merck (Inst)

Travel, Accommodations, Expenses: Genentech, Eli Lilly, Amgen

Wan-Teck Lim

Consulting or Advisory Role: Boehringer Ingelheim, Bristol-Myers Squibb, Novartis

Travel, Accommodations, Expenses: Boehringer Ingelheim, Roche

Patricia LoRusso

Consulting or Advisory Role: Astex Pharmaceuticals, Novartis, Astellas Pharma, Genentech, ARIAD Pharmaceuticals, Alexion Pharmaceuticals, Celgene

Research Funding: Genentech (Inst), Novartis (Inst), Merrimack (Inst), Immunogen (Inst), Tensha Therapeutics (Inst), Taiho Pharmaceutical (Inst), Roche (Inst), Incyte (Inst)

Geoffrey I. Shapiro

Consulting or Advisory Role: Vertex, Chugai Pharmaceutical, G1 Therapeutics, Eli Lilly, EMD Serono, Takeda Pharmaceuticals, Tesaro **Research Funding:** Pfizer (Inst), Genentech (Inst), Bayer AG (Inst), Immune Design (Inst), Vertex (Inst), Millennium Pharmaceuticals (Inst), Puma Biotechnology (Inst), Tensha Therapeutics (Inst), Covidien (Inst), Novartis (Inst), Cellceutix (Inst), Sanofi (Inst), Cyclacel (Inst), Mirati Therapeutics (Inst), AstraZeneca (Inst), GlaxoSmithKline (Inst), Eli Lilly, Aileron (Inst), PharmaMar (Inst), PTC Therapeutics (Inst)

Katie Parker

Employment: Novartis Stock or Other Ownership: Novartis (stock options)

Xueying Chen Employment: Novartis

Somesh Choudhury Employment: Novartis Stock or Other Ownership: Novartis Travel, Accommodations, Expenses: Novartis

Francois Ringeisen Employment: Novartis Stock or Other Ownership: Novartis

Diana Graus-Porta Employment: Novartis Stock or Other Ownership: Novartis **Dale Porter**

Employment: Novartis Stock or Other Ownership: Novartis Patents, Royalties, Other Intellectual Property: Novartis

Randi Isaacs

Employment: Novartis **Stock or Other Ownership:** Novartis **Patents, Royalties, Other Intellectual Property:** Use patent for a particular drug in a specific indication

Reinhard Buettner

Stock or Other Ownership: Targos Molecular Pathology Speakers' Bureau: Roche, Boehringer Ingelheim, Novartis, Pfizer, Merck Sharp & Dohme, Merck, Bristol-Myers Squibb, QIAGEN, Eli Lilly

Jürgen Wolf

Honoraria: AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Merck Sharp & Dohme, Clovis Oncology, Novartis, Pfizer, Roche Consulting or Advisory Role: AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Merck Sharp & Dohme, Clovis Oncology, Novartis, Pfizer, Roche

Research Funding: Novartis (Inst), Pfizer (Inst), Roche (Inst), Bristol-Myers Squibb (Inst)

Acknowledgment

We thank all patients and their families and the study investigators and personnel from each participating center. We also thank the members of the Lung Cancer Group Cologne and the Network Genomic Medicine Germany (supported by Nationales Genomforschungsnetz, grant number 01GS08101), particularly Masyar Gardizi, Matthias Scheffler, Marc Bos, Sebastian Michels, Rieke Fischer, Elisabeth Bitter, Roman Thomas, Monika Serke, Kato Kambartel, and Stefan Krüger. In addition, we thank Kon Skordos and Swarupa Kulkarni from the Novartis Clinical Pharmacology group. Medical writing assistance was provided by Christopher Reina and Jillian Brechbiel of ArticulateScience LLC.

Appendix

Methods

Molecular Prescreening. Molecular prescreening to assess *FGFR* genetic alteration status was implemented after a protocol amendment and before enrollment of cohort 4 during the dose-escalation phase (BGJ398 40 mg once daily). Patient eligibility was determined on the basis of *FGFR* alteration status as assessed centrally or locally using fresh and/or archival tumor samples. Data on the number of patients prescreened versus the number of patients enrolled are not available, because this information was not reported by study sites that performed local assessment of eligibility.

Patients were eligible for enrollment if one of the following genetic alteration criteria was met: (1) *FGFR1* or *FGFR2* amplification was identified using fluorescence in situ hybridization (defined as a ratio of the respective FGFR to chromosome enumeration probe 8 [*FGFR1*] or chromosome enumeration probe 10 [*FGFR2*] of \geq 2.2 or an average *FGFR* copy number of six or more signals/nucleus in \geq 20 contiguous cells from two tumor areas); chromogenic or silver-enhanced in situ hybridization (defined as an average respective *FGFR* copy number of six or more signals/nucleus or a large gene cluster in \geq 30% of tumor cells from \geq 100 contiguous cells from two tumor areas); or quantitative polymerase chain reaction (defined as a respective *FGFR* copy number of at least six); (2) *FGFR3* mutations were detected in exon 7 (R248C, S249C), exon 10 (G372C, A393E, Y375C), or exon 15 (K652M/T, K652E/Q); or (3) other *FGFR* genetic alterations, including gene fusions, were identified.

Alterations other than those described above were reviewed against public databases (eg, COSMIC and dbSNP), and enrollment was approved by study personnel for appropriate alterations (eg, to exclude known single-nucleotide polymorphisms and synonymous substitutions, or to include more recently identified somatic missense mutations). The intent of including unspecified *FGFR* alterations, such as fusions, in the prescreening evaluation was to adapt to the rapidly evolving understanding of the role of *FGFR* in various cancer types and to allow the enrollment of patients with previously unknown activating *FGFR* alterations (Appendix Table A1).

Additional Patient Enrollment Criteria. Patients were required to have adequate bone marrow (absolute neutrophil count $\geq 1,500/\mu L$ [$\geq 1.5 \times 10^9/L$]; platelets $\geq 75,000/\mu L$ [$\geq 75 \times 10^9/L$]; and hemoglobin ≥ 10 mg/dL [≥ 100 g/L]) and renal and hepatic function (serum creatinine $\leq 1.5 \times$ the upper limit of normal [ULN]; calculated or measured creatinine clearance > 75% lower limit of normal; and proteinuria grade ≤ 1 , total bilirubin $\leq 1.5 \times$ ULN, AST and ALT $\leq 2.5 \times$ ULN, and serum albumin is greater than or equal to the lower limit of normal). Further requirements for inclusion included balanced calcium-phosphate homeostasis and adequate cardiovascular function, including a heart rate–corrected QT interval ≤ 470 ms and blood pressure within the normal range.

Patients with primary CNS tumors or CNS tumor involvement were not eligible for enrollment, unless clinically stable. Concomitant therapies prolonging the QT interval or associated with a risk of torsades de pointes were not permitted.

Study Design Rationale. During dose escalation, sequential patient cohorts received BGJ398 5 (starting dose), 10, 20, 40, 60, 100, 125, and 150 mg once daily in continuous 28-day cycles. Fifty milligrams twice daily was also investigated after the maximum tolerated dose (MTD)/recommended phase II dose was established. Patients treated at 100 mg once daily had drug exposures approaching the predicted efficacious level (unbound BGJ398 concentration of 70 ng/mL) on the basis of preclinical data²⁰; however, a high degree of variability in exposure among patients was observed, and one patient experienced a dose-limiting toxicity of reversible grade 3 AST/ALT that resulted in dose interruption and modification during cycle 1. This variability in exposure was not observed at 60 mg once daily. These data supported further exploration of alternative dosing schedules for BGJ398, and the twice daily dose cohort was added in Amendment 5 (Jan 2012; Appendix Table A1). Enrollment in the 50 mg twice daily cohort began after the MTD was established, and was completed before patients were treated in expansion arms 1 and 2. The twice daily dosing schedule was not investigated further, because variability in exposure was also observed with this schedule, and suboptimal concentrations were achieved relative to predicted levels associated with preclinical efficacy.

Tumor Evaluations. Tumors were evaluated, using chest and abdomen computed tomography and, in patients with a history of brain metastases, cranial computed tomography or brain magnetic resonance imaging, at baseline and within 14 days of every second treatment cycle until discontinuation.

Pharmacokinetic Assessments. Serial blood samples were collected predose and up to 24 hours postdose on days 1, 15, and 28 of cycle 1. Samples were processed, and plasma was frozen at $\leq -60^{\circ}$ C until analysis.

BGJ398 plasma concentrations were measured using a validated liquid chromatography-tandem mass spectrometry method with a 1.0 ng/mL lower limit of quantification. Pharmacokinetic (PK) parameters, including maximum plasma concentration, area under the plasma concentration-time curve, time to reach maximum plasma concentration, and half-life, were calculated using noncompartmental methods with Phoenix (Pharsight, Mountain View, CA). Descriptive statistics (mean, median, and standard deviation) were estimated for PK parameters in each cohort. Median values and ranges were provided for half-life and maximum plasma concentration.

Pharmacodynamic Assessments. Blood samples for determination of fibroblast growth factor (FGF) 23 levels were collected predose on days 1, 2, and 15 of cycle 1 and day 1 on cycles 2 and 3, as well as 4 and 8 hours postdose on day 1. Fibroblast growth factor 23 levels were assessed using standard methods. Median percentage changes in fibroblast growth factor 23 levels from baseline were plotted by dose.

Specific Dose-Limiting Toxicities. Specific dose-limiting toxicities modeled in the second adaptive Bayesian logistic regression model included corneal opacity, ectopic mineralization/calcification, or elevated serum creatinine (grade 1 for > 7 consecutive days) and serum ionized phosphorus (Pi) > 5.5 mg/dL and/or total serum calcium × Pi > 55 mg²/dL² despite Pi-lowering therapy for \geq 14 days.

Hyperphosphatemia Grading, Management, and Prevention. Because grading for hyperphosphatemia is not defined in the Common Terminology Criteria for Adverse Events (version 3) guidelines,²⁴ hyperphosphatemia was defined in this study as any value above the ULN, and severity was based on investigator discretion; severe hyperphosphatemia was considered grade 3. During dose escalation, treatment with phosphate binders was initiated when serum phosphate levels exceeded 5.5 mg/dL, with the goal of maintaining levels at \leq 7 mg/dL. Prophylactic use of phosphate binders was instituted after the MTD determination and was used routinely in the dose expansion cohorts. Dose interruption was indicated when phosphate levels remained > 7 mg/dL despite the use of phosphate binders for \geq 14 days.

Results

The BGJ398 125-mg dose level was determined to be the MTD on the basis of clinical safety and PK data and was supported quantitatively by the Bayesian logistic regression models used in this study. The 3-weeks-on/1-week-off intermittent schedule was investigated on the basis of the observation that the majority of patients being treated with the 125 mg once daily dosing schedule required dose interruptions by day 22 to control hyperphosphatemia, with interruptions lasting 7 days (median). The 3-weeks-on/1-week-off schedule was evaluated in an attempt to establish a regimen that allowed control of hyperphosphatemia over the course of the cycle. In addition, although exposure was variable, the majority of patients treated at 125 mg once daily achieved BGJ398 exposures above the threshold associated with preclinical evidence of fibroblast growth factor receptor pathway inhibition and in vivo efficacy.²⁰ Although the data are limited, this was not the case for patients treated at the 100 mg once daily dosage. It was recognized that many patients required dose reduction over the course of their therapy for the management of chronic toxicities; however, given the fact that patients with *FGFR* alterations are rare and are not easily identified, we chose to initiate treatment with a dose most likely to provide clinical benefit. Moreover, the rate of hyperphosphatemia and the overall safety profile between the 100-mg and 125-mg dose levels was not substantially different; therefore, 125 mg once daily 3 weeks on/1 week off was selected as the optimal BGJ398 dose and schedule for further exploration.

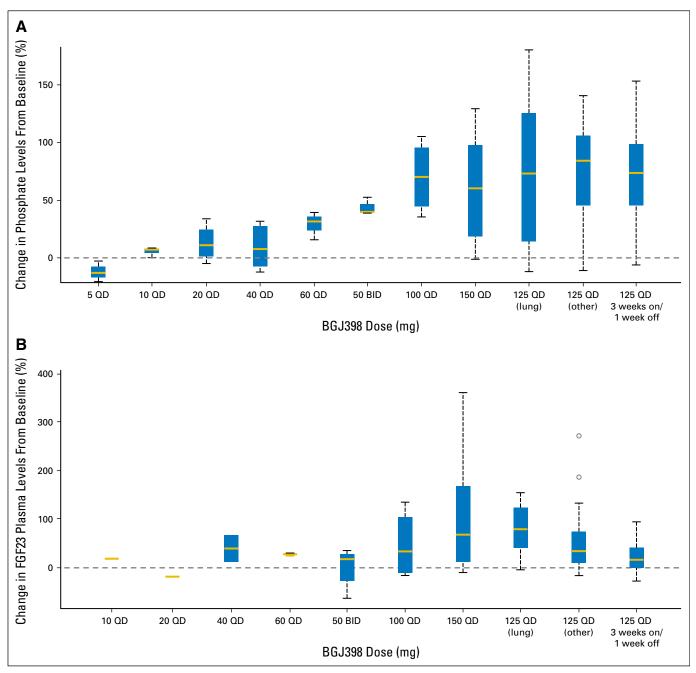


Fig A1. Pharmacodynamics of BGJ398. Percentage change from baseline in (A) phosphate and (B) FGF23 plasma levels on cycle 1 day 15 by dose. Median values are shown as box plots. BID, twice daily; FGF, fibroblast growth factor; QD, once daily.

		Dationala for Destant 1.4
Date Released	Patient Prescreening/Molecular Eligibility	Rationale for Protocol Amendment/ Reference(s)
Aug 2009	No molecular preselection	
Feb 2010	No molecular preselection	
May 2010	With the availability of new clinical biomarker tests, enrollment was limited to patients with advanced solid tumors harboring <i>FGFR1</i> or <i>FGFR2</i> gene amplification or <i>FGFR3</i> gene mutation. The molecular selection criteria were implemented before enrollment of patients at the 40-mg dose level during dose escalation.	 This amendment allowed for an earlier readout of pharmacodynamic marker in the patient population harboring genetic alterations in the FGFR pathway, made feasible by the clini availability of biomarker tests and emerging epidemiology of the different subpopulations. The state of understanding at that tim regarding the genetic basis of sensitivity to BGJ398 was describe by Guagnano et al,²⁰ with evidence BGJ398 sensitivity in preclinical models of <i>FGFR</i> amplification in breat and lung cancer and hotspot mutatic in bladder cancer.
Apr 2011	An option was added for central laboratory evaluation to detect <i>FGFR1</i> amplification in squamous cell lung cancer. Before the amendment, patients with squamous cell lung cancer with <i>FGFR1</i> amplification were eligible for enrollment on the basis of a molecular analysis performed by a local laboratory at an individual study site.	 This amendment was implemented to improve enrollment, because local laboratory testing was not available some study sites. Both local and central laboratories use stringent guidelines to define amplification, as described in the Appendix.
Sep 2011	 This amendment allowed the opening of a dedicated expansion arm at the MTD/RP2D for patients with squamous cell carcinoma harboring <i>FGFR1</i> amplification. Quantitative PCR was added as a methodology for detection of <i>FGFR1</i> amplification. 	On the basis of newly published evidence, ^{11,12} this amendment allowed probing for early signs of activity in a specific population of patients with squamous cell lung cancer and <i>FGFR1</i> amplification. Quantitative PCR was added to the approved study methodologies to expand the ability of study sites to perform local testing.
Jan 2012	This amendment introduced an alternative twice daily BGJ398 dosing schedule (50 mg).	With exposure levels in the predicted efficacious range at 100 mg once da but also variable PK parameters, the I profile achieved with an alternative twice daily schedule was compared with the PK profile after 100 mg on daily dosing.
Jun 2012	This amendment allowed the enrollment of patients with any <i>FGFR</i> genetic alteration. The change also allowed patients to be enrolled on the basis of <i>FGFR</i> amplification identified using methods other than fluorescence in situ hybridization, chromogenic in situ hybridization, silver-enhanced in situ hybridization, or reverse transcriptase PCR.	To address the evolving understanding <i>FGFR</i> contributions to human cance this change allowed exploration of potential BGJ398 activity in patients with advanced tumors carrying <i>FGF</i> amplification or other <i>FGFR</i> genetic alterations that are potentially oncogenic drivers to be enrolled. Patients with <i>FGFR</i> gene fusions were also permitted to be enrolled on the basis of a report of <i>FGFR3-TACC</i> fusions in patients with glioblastom, published later in 2012* and person communications in advance of a 20 publication t regarding the use of MiOncoseq to identify <i>FGFR2</i> fusion in patients with cholangiocarcinoma. The underlying mechanism of previous observed sensitivity of bladder cance cell lines to BGJ398 ²⁰ was later explained by a publication in early 20 reporting <i>FGFR3</i> gene fusions in human bladder cancer. ¹⁰ In addition, patient enrollment was enhanced by allowing additional options for genetic screening.
		optiono foi gonotio obroching.
	Aug 2009 Feb 2010 May 2010 Apr 2011 Sep 2011 Jan 2012	Aug 2009 No molecular preselection May 2010 With the availability of new clinical biomarker tests, enrollment was limited to patients with advanced solid tumors harboring <i>FGFR1</i> or <i>FGFR2</i> gene amplification or <i>FGFR3</i> gene mutation. The molecular selection criteria were implemented before enrollment of patients at the 40-mg dose level during dose escalation. Apr 2011 An option was added for central laboratory evaluation to detect <i>FGFR1</i> amplification in squamous cell lung cancer. Before the amendment, patients with squamous cell lung cancer with <i>FGFR1</i> amplification were eligible for enrollment on the basis of a molecular analysis performed by a local laboratory at an individual study site. Sep 2011 This amendment allowed the opening of a dedicated expansion arm at the MTD/RP2D for patients with squamous cell carcinoma harboring <i>FGFR1</i> amplification. Jan 2012 This amendment introduced an alternative twice daily BGJ398 dosing schedule (50 mg). Jun 2012 This amendment allowed the enrollment of patients with any <i>FGFR1</i> amplification, identified using methods other than fluorescence in situ hybridization, chromogenic in situ hybridization, silver-enhanced in situ hybridization, or bromegenic in situ hybridization, o

Protocol/Amendment	Date Released	Patient Prescreening/Molecular Eligibility	Rationale for Protocol Amendment, Reference(s)
Amendment 7	Jun 2013	This amendment allowed the opening of an expansion arm (arm 3), which used an alternative BGJ398 dosing schedule of 125 mg once daily 3 weeks on/1 week off for patients with advanced solid tumors harboring any <i>FGFR</i> genetic alteration (including gene amplification, mutation, or fusion).	Initiated a new expansion arm using an alternative dosing schedule implemented to aid management of hyperphosphatemia.
Amendment 8	Apr 2014	This amendment allowed the opening of an expansion arm (arm 4) dedicated to evaluating BGJ398 in patients with advanced or metastatic urothelial carcinoma with an <i>FGFR3</i> gene mutation or fusion.	 Expansion arm 4 specifically evaluated BGJ398 in a patient population deemed sensitive on the basis of phase I study results as detailed in th article. With this amendment, new study site: were opened in current countries ar in an additional eight countries, bringing the total number of study site to 57 and enabling patient enrollment an <i>FGFR3</i>-altered urothelial carcinom study arm.
Amendment 9	Apr 2015	This amendment made no change to molecular screening/eligibility.	

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 *Singh D, et al: Science 337:1231-1235, 2012.

 †Wu, et al: Cancer Discov 3:636-647, 2013.

	BGJ398 Daily Dose						
Treatment Status	5-60 mg (n = 19)	100 mg (n = 6)	125 mg Continuous (n = 57)	125 mg 3 Weeks On/ 1 Week Off (n = 40)	150 mg (n = 6)	BGJ398 50 mg Twice Daily (n = 4)	All Patients (N = 132)
Treatment ongoing	19 (100)	6 (100)	55 (96.5)	37 (92.5)	6 (100)	4 (100)	127 (96.2)
Treatment discontinued	0	0	2 (3.5)	3 (7.5)	0	0	5 (3.8)
Primary reason for discontinuation							
Disease progression	13 (68.4)	4 (66.7)	36 (63.2)	29 (72.5)	4 (66.7)	3 (75.0)	89 (67.4)
Adverse event	4 (21.1)	2 (33.3)	6 (10.5)	4 (10.0)	1 (16.7)	0	17 (12.9)
Patient decision	2 (10.5)	0	8 (14.0)	3 (7.5)	1 (16.7)	0	14 (10.6)
Death	0	0	3 (5.3)	1 (2.5)	0	1 (25.0)	5 (3.8)
Protocol deviation	0	0	1 (1.8)	0	0	0	1 (0.8)
Administrative problem	0	0	1 (1.8)	0	0	0	1 (0.8)

Table A3. Deaths						
Patient	Tumor Type	Dose (mg once daily)	Study Day of Last Dose	Study Day of Death	Primary Cause of Death	
1	Lung	125	27	29	Sepsis	
2	Lung	125	50	51	Cardiac arrest*	
3	Lung	125	7	10	Sepsis	
4	Lung	125	96	114	Respiratory failure secondary to progressive disease	
5	Lung	125	64	77	Squamous cell carcinoma	
6	Lung	125	57	75	Lung cancer	
7	Squamous cell carcinoma	125	110	114	Squamous cell carcinoma	
8	Anal	125†	21	31	Respiratory failure	
9	Breast	125†	14	27	Breast cancer	
10	Esophageal	125†	14	42	Tumor progression	
11	Bladder	125†	155	171	Bladder cancer	
12	Renal	125†	11	39	Respiratory failure caused by progressive disease	
13	Prostate/urothelial	125†	21	32	Urothelial cancer	
14	Lung	150	104	115	Tumor progression	
15	Lung	50‡	29	34	Not reported (unknown disease-related event)	

*This 75-year-old male with advanced non-small-cell lung cancer died as a result of cardiac arrest 1 day after the last dose of the study drug. The patient had no clinical symptoms for myocardial infarction. Disease progression was suspected to be the cause; however, in the absence of a follow-up computed tomography scan to confirm, and without any other clear cause of death, this patient's death was reported as suspected to be related to BGJ398 treatment. †Three weeks on/1 week off.

‡Twice daily.

BGJ398 Once Daily Dose/Parameter	$AUC_{(0-24 h)}$ (h × ng/mL)	$AUC_{(0-inf)}$ (h $ imes$ ng/mL)	C _{max} (ng/mL)	T _{max} (hours)	T _{1/2} (hours)	R _{acc}
Cycle 1 day 1						
20 mg(n = 4)						
No.	3	3	4	4	3	_
Mean (SD)	71,79 (38,29)	78.30 (48.35)	15.80 (3.71)	_	_	_
Median	58.84	60.05	16.00	2.04	2.69	_
40 mg (n = 6)						
No.	5	4	6	6	4	_
Mean (SD)	167.39 (61.57)	185.97 (71.66)	27.22 (11.83)	_	_	_
Median	159.96	179.23	25.10	2.08	4.15	_
60 mg (n = 3)	100100	170120	20.10	2.00		
No.	3	2	3	3	2	_
Mean (SD)	255.18 (74.95)	310.79 (32.37)	42.37 (25.92)	_	_	_
Median	277.66	310.79	53.10	2.00	5.61	_
100 mg (n = 6)	277.00	010.70	00.10	2.00	0.01	
No.	6	5	6	6	6	_
Mean (SD)	473.05 (445.10)	469.35 (527.10)	76.42 (82.66)	_	_	_
Median	320.30	318.50	54.10	3.00	5.71	_
125 mg lung (n = 30)	020.00	010.00	01.10	0.00	0.71	
No.	23	18	29	29	22	_
Mean (SD)	581.35 (481.59)	660.88 (566.64)	71.59 (60.76)			
Median	495.59	568.39	51.40	3.00	5.05	_
125 mg other (n = 26)	100100	000.00	01110	0.00	0.00	
No.	22	16	26	26	19	_
Mean (SD)	938.47 (603.45)	1057.38 (683.66)	115.98 (75.00)			_
Median	810.28	929.43	121.50	3.00	5.90	_
125 mg 3 weeks on/1 week off (n = 39)	010.20	020110	121100	0.00	0.00	
No.	33	27	36	36	30	_
Mean (SD)	846.50 (658.44)	788.55 (606.72)	92.81 (58.43)			_
Median	708.74	669.10	71.90	3.00	4.88	_
150 mg (n = 6)	700.71	000.10	71.00	0.00	1.00	
No.	6	6	6	6	6	_
Mean (SD)	677.90 (343.78)	706.82 (353.33)	118.65 (105.74)	_	_	_
Median	710.07	726.87	91.05	2.63	5.20	_
ycle 1 Day 15						
20 mg (n = 4)						
No.	4	3	4	4	4	3
Mean (SD)	212.58 (210.41)	115.14 (47.99)	20.85 (4.28)	· 	_	2.82 (1.5
Median	133.18	131.51	19.35	1.99	5.91	2.02 (1.0
		inued on following page)			0.01	20

Та	ble A4. BGJ398 Pharmacok	inetic Parameters (once o	daily doses) (continu	ued)		
BGJ398 Once Daily Dose/Parameter	$\rm AUC_{(0-24~h)}$ (h \times ng/mL)	$\rm AUC_{(0-inf)}$ (h $ imes$ ng/mL)	C _{max} (ng/mL)	T _{max} (hours)	T _{1/2} (hours)	R_{acc}
40 mg (n = 6)						
No.	4	2	5	5	3	3
Mean (SD)	183.48 (113.07)	229.95 (144.09)	30.24 (18.64)	_	_	0.82 (0.50)
Median	191.98	229.95	28.00	2.07	6.17	0.91
60 mg (n = 3)						
No.	3	2	3	3	2	3
Mean (SD)	1,007.57 (477.30)	1,081.47 (749.36)	80.37 (44.54)	_		4.37 (2.86)
Median	1260.03	1081.47	85.70	2.08	9.84	
	1260.03	1081.47	85.70	2.08	9.84	4.13
100 mg (n = 6)		_	_	_		
No.	4	2	5	5	4	4
Mean (SD)	1,375.03 (1,244.19)	685.42 (600.48)	136.70 (104.59)	—	—	5.65 (3.51)
Median	1,045.91	685.42	93.70	3.00	7.89	5.75
125 mg lung (n = 30)						
No.	15	4	23	23	12	11
Mean (SD)	3,483.83 (2,991.54)	732.88 (549.51)	209.14 (137.37)	_	_	8.18 (5.43)
Median	3,240.82	708.11	219.00	4.00	15.43	6.60
125 mg other (n = 26)	-,					
No.	19	2	22	22	12	16
Mean (SD)	3,902.68 (2,261.84)	2,950.53 (450.98)	22 261.44 (137.80)		12	5.88 (4.67)
Median	3,442.56	2950.53	235.00	3.54	11.12	4.49
125 mg 3 weeks on/1 week off (n = 39)						
No.	23	2	29	29	12	20
Mean (SD)	3,296.27 (1,998.18)	900.93 (440.79)	212.68 (115.07)	_	_	7.63 (6.85)
Median	3,183.30	900.93	244.00	4.12	11.55	4.78
150 mg (n = 6)						
No.	4	2	4	4	3	3
Mean (SD)	1,565.50 (1,671.84)	757.30 (727.61)	133.98 (139.23)	_	_	3.38 (2.92)
Median	1,025.44	757.30	85.30	3.00	9.34	3.78
Cycle 1 Day 28	1,020.11	707.00	00.00	0.00	0.01	0.70
20 mg (n = 4)						
	3	2	2	2	0	0
No.		2	3	3	2	2
Mean (SD)	146.81 (46.90)	133.59 (66.45)	22.80 (2.41)	—	_	2.52
Median	163.67	133.59	22.60	2.00	5.30	2.52
40 mg (n = 6)						
No.	3	3	4	4	3	2
Mean (SD)	179.03 (160.81)	201.66 (209.85)	18.35 (11.59)	—	—	1.35 (0.02)
Median	158.37	139.61	19.25	3.02	2.76	1.35
60 mg (n = 3)						
No.	1	0	2	2	1	1
Mean (SD)	2,192.88	- -	113.95 (49.57)	_	_	6.93
Median	2,192.88		113.95	4.09	12.46	6.93
	2,132.00		113.35	4.03	12.40	0.35
100 mg (n = 6)	,		-	-	0	0
No.	4	1	5	5	3	3
Mean (SD)	1,977.52 (1,475.18)	438.58	117.90 (82.26)	_		6.37 (5.95)
Median	1,765.09	438.58	112.00	3.08	20.14	2.99
125 mg lung (n = 30)						
No.	18	2	19	19	13	13
Mean (SD)	2,427.38 (1,934.85)	202.81 (80.69)	153.53 (107.66)	_	_	10.00 (13.76
Median	2,286.07	202.81	153.00	3.20	17.62	6.23
125 mg other (n = 26)	,					
No.	13	1	17	17	10	6
Mean (SD)			273.63 (137.04)			
	3,405.44 (1,819.50)	4,093.22				4.98 (1.66)
Median	4,091.88	4,093.22	274.00	4.00	15.88	5.43
150 mg (n = 6)						
No.	4	1	4	4	2	2
Mean (SD)	1,835.90 (581.88)	1,909.96	146.75 (19.67)	—	—	7.86 (8.79)
Median	1,784.66	1,909.96	151.50	2.99	12.44	7.86

Abbreviations: AUC, area under the plasma concentration-time curve; C_{max}, maximum plasma concentration; R_{acc}, accumulation ratio; T_{1/2}, terminal elimination half-life; T_{max}, time to reach maximum plasma concentration

	Table A5. FGFR Mutations and Fusions	
Best Percentage Change in Tumor	Tumor Type	FGFR Genetic Alteration
-15.1	Angiosarcoma	FGFR1 S125L mutation
-100	Bladder/urothelial carcinoma	FGFR3 S249C mutation
-65.0	Bladder/urothelial carcinoma	FGFR3 S249C mutation
-48.2	Bladder/urothelial carcinoma	FGFR3 S249C mutation
-28.2	Bladder/urothelial carcinoma	FGFR3 S249C mutation
11.4	Bladder/urothelial carcinoma	FGFR3 S249C mutation
-27.3	Bladder/urothelial carcinoma	FGFR3 K652E mutation
7.3	Bladder/urothelial carcinoma	FGFR3 K652E mutation
34.2	Bladder/urothelial carcinoma	FGFR3 Y373C mutation
64.3	Breast	FGFR3 F384L mutation
-21.7	Cholangiocarcinoma	FGFR2 N549S mutation
-20.2	Cholangiocarcinoma	FGFR2 10Q26.13 fusion (partner unknown)
-9.8	Cholangiocarcinoma	FGFR2 BICC1 fusion
40.4	Colon adenocarcinoma	FGFR1 S125L mutation
40.4	Head and neck carcinoma	FGFR3 R248C mutation
10.4	Hepatocellular carcinoma	FGFR2 BICC1 fusion
-4.8	Lung adenocarcinoma	FGFR3 F384L mutation
16.7	Lung adenocarcinoma	FGFR3 F384L mutation
10.4	Lung adenocarcinoma	FGFR3 F386L mutation
-2.0	Lung squamous cell carcinoma	FGFR3 S249C mutation

NOTE. Table includes patients treated at \geq 100 mg and who had an *FGFR* mutation or fusion noted by any methodology and were included in the evaluation of best change from baseline in the sum of longest diameters in target lesion(s). Abbreviation: FGFR, fibroblast growth factor receptor.