Cancer Therapy: Clinical

Randomized Phase Ib/II Trial of Rilotumumab or Ganitumab with Panitumumab versus Panitumumab Alone in Patients with Wild-type KRAS Metastatic Colorectal Cancer

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Abstract

**Purpose:** Panitumumab, a fully human anti-epidermal growth factor receptor monoclonal antibody (mAb), has demonstrated efficacy in patients with wild-type KRAS metastatic colorectal cancer (mCRC). Rilotumumab and ganitumab are investigational, fully human mAbs against hepatocyte growth factor (HGF)/scatter factor and IGF1R, respectively. Here we evaluate combining rilotumumab or ganitumab with panitumumab in previously treated patients with wild-type KRAS mCRC.

**Experimental Design:** Part 1 was a phase Ib dose-finding study of panitumumab plus rilotumumab. The primary endpoint was the incidence of dose-limiting toxicities (DLT). Part 2 was a randomized phase II trial of panitumumab in combination with rilotumumab, ganitumab, or placebo. The primary endpoint was objective response rate (ORR); safety, progression-free survival (PFS), and overall survival (OS) were secondary endpoints. Archival tissue specimens were collected for exploratory correlative work.

**Results:** In part 1, no DLTs were reported. A recommended phase II dose of 10 mg/kg rilotumumab was selected. In part 2, for the panitumumab plus rilotumumab (n = 48), panitumumab plus ganitumab (n = 46), and panitumumab plus placebo arms (n = 48), the ORRs were 31%, 22%, and 21%, respectively. The median PFS was 5.2, 5.3, and 3.7 months and median OS 13.8, 10.6, and 11.6 months, respectively. Adverse events were tolerable. Exploratory biomarker analyses, including MET and IGF-related protein expression, failed to indicate conclusive predictive evidence on efficacy endpoints.

**Conclusions:** Panitumumab plus rilotumumab met the prespecified criterion for improvement in ORR whereas ganitumab did not. This is the first study to suggest a benefit for combining an HGF inhibitor (rilotumumab) with panitumumab in previously treated patients with wild-type KRAS mCRC. Clin Cancer Res; 20(16); 4240–50. ©2014 AACR.

Introduction

Worldwide, colorectal cancer remains the third most common malignancy in men and the second most common malignancy in women, affecting a total of 1.2 million individuals and resulting in 608,000 deaths (1). With few treatment options for patients who have progressed on chemotherapy, patients with metastatic colorectal cancer (mCRC) continue to search for innovative therapies. Panitumumab, a fully human monoclonal antibody (mAb) against the epidermal growth factor receptor (EGFR), has demonstrated efficacy as a single agent and in combination with chemotherapy in wild-type (WT) KRAS mCRC tumors (2–5).

MET and the insulin-like growth factor 1 receptor (IGF1R) are tyrosine kinases that share multiple downstream pathways with EGFR, including the Ras–Raf–MAPK and PI3K/Akt pathways (6–9). Hepatocyte growth factor/scatter factor (HGF/SF) is the only known ligand for MET. This pathway is required for normal physiologic...
Translational Relevance

Upregulation of the hepatocyte growth factor (HGF)/MET and the IGF1R pathways have been suggested as potential mechanisms of signal escape in colorectal tumors upon treatment with epidermal growth factor receptor (EGFR) inhibitors. In this clinical study, it is suggested for the first time that the dual inhibition of the EGFR by panitumumab and HGF by rilotumumab translates in incremental antitumor activity over panitumumab as a single agent in patients with refractory wild-type KRAS metastatic colorectal cancer.

Rilotumumab, Ganitumab, and Panitumumab in Treatment of mCRC

Materials and Methods

Patients

Eligible patients were ≥18 years of age, had an Eastern Cooperative Oncology Group (ECOG) performance score of 0 or 1, and had histologically or cytologically confirmed metastatic adenocarcinoma of the colon or rectum. Radiographic evidence of disease progression during or following prior treatment with irinotecan- and/or oxaliplatin-based chemotherapy for mCRC was required. At least 1 unidimensionally measurable lesion per modified RECIST 1.0 was required. Archival tumor tissue was confirmed by a central laboratory to be WT KRAS using a validated test method.

Patients were excluded if they received prior treatment with an anti-EGFR inhibitor (e.g., panitumumab, cetuximab, erlotinib, and/or gefitinib) unless treatment was received in the adjuvant setting ≥6 months before enrollment. Prior treatment with MET or IGF1R inhibitors was not allowed. Additional exclusion criteria included the use of systemic chemotherapy or radiotherapy ≤21 days before enrollment and the use of any targeted therapies ≤30 days before enrollment (including bevacizumab).

The study protocol was approved by the institutional review boards or the ethics committees at all participating institutions. Written, informed consent was provided by all patients before any study-related treatment was conducted. Amgen Inc. maintained the database for this study. All authors had access to the complete data.

Study design and treatment

This is an exploratory, global, multicenter 3-part phase Ib/II trial designed to evaluate the efficacy and safety of rilotumumab or ganitumab in combination with panitumumab versus panitumumab plus placebo in previously treated patients with WT KRAS mCRC (NCT00788957; Supplementary Fig. S1). Exploratory correlative work was conducted only in the phase II portion of the study.

Part 1 (phase Ib) consisted of a dose de-escalation study to evaluate and select a tolerable dose of rilotumumab in combination with panitumumab to be used in part 2.

Part 2 (phase II) consisted of a 3-arm randomized trial to evaluate the efficacy and safety of panitumumab in combination with rilotumumab, ganitumab, or placebo in previously treated patients with WT KRAS mCRC.

In part 3 (phase II), patients in the panitumumab plus placebo arm of part 2 could be randomized to receive either rilotumumab or ganitumab monotherapy, if the patient ended their treatment because of disease progression or intolerability but remained eligible by protocol standards for additional therapy. Part 3 of the study is ongoing and results will be reported at a later date.

Phase Ib: Evaluation of DLT(s) in part 1

Patients received 6 mg/kg panitumumab in combination with 10 mg/kg rilotumumab intravenously once every 2 weeks (Q2W). The starting dose of rilotumumab was based on unpublished pharmacokinetic data. A human plasma concentrations of 10 mg/kg rilotumumab administered Q2W showed a median trough concentration of 158.6 μg/mL, which exceeds the IC90 values (approximately 75 μg/mL) predicted from in vitro cell proliferation assays and therefore was likely to be an effective dose. Based on the incidence of dose-limiting toxicities (DLT), dose de-escalation to 5 mg/kg rilotumumab was allowed. No dose reduction in panitumumab was planned. A DLT evaluable patient was defined as a patient who had received at least 2 doses of panitumumab and rilotumumab as scheduled (i.e., week 1 and 3) and had a minimum 4 weeks (28 days) of follow-up for safety or had received at least 1 dose of panitumumab and rilotumumab and had a DLT within the first 4 weeks (28 days) on study.

If ≤1 of the first 6 evaluable patients experienced a DLT(s), the 6 mg/kg panitumumab plus 10 mg/kg rilotumumab dose was determined tolerable and used in part 2.
If 2 of the first 6 evaluable patients experienced a DLT(s), then 3 additional patients were enrolled to receive the 6 mg/kg panitumumab plus 10 mg/kg rilotumumab dose. If ≥3 of the first 6 evaluable patients experienced a DLT (s), then the dose of panitumumab plus rilotumumab was determined not tolerable and a lower dose of 5 mg/kg rilotumumab was to be evaluated in combination with panitumumab.

DLTs were defined as any grade 3 or 4 therapy-related adverse event (AE) or laboratory abnormality that was deemed clinically significant by the investigator. Fatigue, nausea, vomiting, diarrhea, hypomagnesemia, rash, neutropenia, thrombocytopenia, and increased aspartate aminotransferase (AST)/alanine aminotransferase (ALT) were not considered to be DLTs unless the following criteria were met: grade 3 fatigue that persisted for >7 days, or grade 4 fatigue; grade 3 or 4 nausea, diarrhea, or vomiting despite maximum supportive care; grade 4 hypomagnesemia despite IV maximal magnesium replacement; grade 4 rash/desquamation; grade 3 or 4 neutropenia with persistent fever ≥38.5°C; grade 4 neutropenia or thrombocytopenia >7 days; elevated ALT or AST >10× upper limit of normal.

DLT-eligible patients received ≥2 doses of panitumumab plus rilotumumab as scheduled and had a minimum 28 days of follow-up for safety or received ≥1 dose of panitumumab plus rilotumumab and had a DLT within the first 28 days on study.

Phase II: Randomization in part 2
In part 2, patients were randomized in a ratio of 1:1:1 to treatment arms: 6 mg/kg panitumumab plus 10 mg/kg rilotumumab; 6 mg/kg panitumumab plus 12 mg/kg ganitumab; and 6 mg/kg panitumumab plus placebo. Randomization was stratified by prior chemotherapy regimen (treated with irinotecan or oxaliplatin vs. both). Panitumumab was open-label; rilotumumab, ganitumab, and placebo were double-blinded. Treatments were administered Q2W until disease progression, intolerability, withdrawal, or death.

Tumor response was assessed by the investigator using RECIST 1.0 criteria at weeks 8, 12, 16, 24, 32, 40, 48, and every 12 weeks until disease progression. Responses were confirmed no earlier than 28 days after the criteria for response were first met.

Statistical analysis
In part 1, the primary objective was to identify a tolerable dose of panitumumab plus rilotumumab based on the incidence of DLTs. A secondary objective was to evaluate the safety of panitumumab plus rilotumumab.

In part 2, the primary objective was to evaluate the efficacy as measured by the objective response rate (ORR) of panitumumab plus rilotumumab and panitumumab plus ganitumab versus panitumumab plus placebo. Key secondary objectives included safety (AEs and laboratory values), duration of response, disease control [complete response plus partial response plus stable disease (SD); SD required patients to be on study for ≥49 days before SD], progression-free survival (PFS), and overall survival (OS).

Bayesian analysis was used for the statistical interpretation of ORR. This method calculates the posterior distribution of the odds ratio (OR) of ORRs for the experimental arms and the control arm using prior distributions of ORRs. It was prespecified that if there was ≥90% posterior probability that the combination therapy was better than panitumumab alone (i.e., OR > 1) as evaluated by objective tumor response, the combination was considered promising. If there was between 50% and 90% probability, the combination was considered indeterminate. If there was <50% probability, the combination was considered not promising. An ORR prior distribution for panitumumab monotherapy was derived from 4 prior studies in which patients had received both prior oxaliplatin and irinotecan (5, 19). The ORR prior distributions for the combination arms were assumed to have the same mean as the panitumumab alone arm and the ORR posterior distribution for each arm combines the prior distributions with observed ORRs from the study.

Results
Part 1: Phase Ib
Eleven patients from 3 sites (1 each in the USA, Belgium, and Spain) were enrolled to the starting dose of 10 mg/kg rilotumumab. Accrual was not halted until the dose decision for part 2 was made and therefore ≥6 patients were enrolled. No DLTs were reported for the first 6 DLT-evaluable patients, thus the Q2W dose of 6 mg/kg panitumumab plus 10 mg/kg rilotumumab was used in part 2. Capillary leak syndrome was reported in one of the first 6 DLT-evaluable patients (64 days after the first dose), but this occurred outside of the DLT reporting window. DLTs, defined as any grade 3 or 4 treatment-related AE or laboratory abnormality that was deemed clinically significant by the investigator, were not reported beyond the first 6 DLT-evaluable patients. However, AEs were assessed for all 11 patients in part 1 (described below).

Patient demographics and disease characteristics are summarized in Table 1. Treatment-related grade 3 or 4 AEs consisted of acneiform dermatitis/rash (55%), paronychia (18%), infection (9%), capillary leak syndrome (9%), erythema (9%), nail disorder (9%), and pruritus (9%). Treatment-emergent grade 3 or 4 AEs that occurred in ≥15% of patients were acneform dermatitis/rash (55%), fatigue (18%), and paronychia (18%). Five patients had serious AEs. None were fatal and consisted of acneform dermatitis (grade 3) and capillary leak syndrome (grade 3), both considered to be related to study treatment, intestinal obstruction (grade 4), anemia/general health deterioration (grade 3), and cerebrovascular accident (grade 4). There was 1 death on study (within 30 days of the last dose of protocol-specified therapy) because of progressive disease.
Part 2: Phase II patient demographics

From June 9, 2009 to February 5, 2010, 142 patients were randomly assigned from 37 sites in 11 countries (Table 1). The mean number of patients enrolled per site was 3.8 and the median was 3.0. All patients received at least one dose of study therapy. Median follow-up time (from the date of first dose to the last on-study or long-term follow-up visit or death date) was 7.2 months (range, 0.9–12.2), 6.8 months (range, 0.7–13.4), and 6.9 months (range, 0.5–12.0 months), in the panitumumab plus rilotumumab, panitumumab plus ganitumab, and panitumumab plus placebo arms, respectively.

Demographics and baseline characteristics were generally balanced across treatment arms (Table 1), with the exception of fewer patients with ECOG 1 performance status in the panitumumab plus rilotumumab arm and approximately twice as many patients ≥65 years old in the combination arms compared with the panitumumab plus placebo arm. One patient had an ECOG performance score of 2 and was in the panitumumab plus rilotumumab arm. Two patients in the control arm had not received first-line therapy for mCRC; both patients had received oxaliplatin-based chemotherapy for non-mCRC and progressed on therapy before entering the study.

Demographics and baseline characteristics at baseline

<table>
<thead>
<tr>
<th></th>
<th>Part 1: Phase Ib</th>
<th>Part 2: Phase II</th>
<th>Panitumumab + ganitumab</th>
<th>Panitumumab + placebo</th>
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<tbody>
<tr>
<td></td>
<td>Panitumumab</td>
<td>Panitumumab</td>
<td>Panitumumab</td>
<td>Panitumumab</td>
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<tr>
<td></td>
<td>(AMG 102)</td>
<td>(AMG 102)</td>
<td>(AMG 479)</td>
<td>(n = 48)</td>
</tr>
<tr>
<td>Men, n (%)</td>
<td>5 (45)</td>
<td>29 (60)</td>
<td>25 (54)</td>
<td>28 (58)</td>
</tr>
<tr>
<td>Age, mean years (range)</td>
<td>56.5 (37–75)</td>
<td>62.1 (45–78)</td>
<td>62.0 (33–81)</td>
<td>55.0 (19–75)</td>
</tr>
<tr>
<td>Age ≥ 65 years, n (%)</td>
<td>4 (36)</td>
<td>20 (42)</td>
<td>18 (39)</td>
<td>9 (19)</td>
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<tr>
<td>ECOG status, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>0</td>
<td>6 (55)</td>
<td>24 (50)</td>
<td>18 (39)</td>
<td>15 (31)</td>
</tr>
<tr>
<td>1</td>
<td>5 (45)</td>
<td>23 (48)a</td>
<td>28 (61)</td>
<td>33 (69)</td>
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<tr>
<td>Metastatic sites, n (%)</td>
<td></td>
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<tr>
<td>Liver only</td>
<td>3 (27)</td>
<td>5 (10)</td>
<td>4 (9)</td>
<td>5 (10)</td>
</tr>
<tr>
<td>Liver + other sites</td>
<td>5 (45)</td>
<td>32 (67)</td>
<td>29 (63)</td>
<td>27 (56)</td>
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<tr>
<td>Prior line of therapy for mCRC, n (%)</td>
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</tr>
<tr>
<td>First-line therapy</td>
<td>11 (100)</td>
<td>48 (100)</td>
<td>46 (100)</td>
<td>46 (96)b</td>
</tr>
<tr>
<td>Second-line therapy</td>
<td>8 (73)</td>
<td>33 (69)</td>
<td>26 (57)</td>
<td>31 (65)</td>
</tr>
<tr>
<td>Third-line therapy and later</td>
<td>3 (27)</td>
<td>16 (33)</td>
<td>12 (26)</td>
<td>14 (29)</td>
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<td>Prior therapies for mCRC, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>OXaliplatin</td>
<td>8 (73)</td>
<td>42 (88)</td>
<td>40 (87)</td>
<td>39 (81)</td>
</tr>
<tr>
<td>Irinotecan</td>
<td>11 (100)</td>
<td>32 (67)</td>
<td>26 (57)</td>
<td>30 (63)</td>
</tr>
<tr>
<td>OXaliplatin and irinotecan</td>
<td>8 (73)</td>
<td>26 (54)</td>
<td>20 (44)</td>
<td>23 (48)</td>
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<tr>
<td>Bevacizumab</td>
<td>5 (45)</td>
<td>17 (35)</td>
<td>12 (26)</td>
<td>19 (40)</td>
</tr>
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</table>

aOne patient with ECOG performance score of 2 was enrolled and was not counted as either ECOG performance score of 0 or 1; data from this patient were included in all efficacy and safety analyses.
bTwo patients had not received first-line therapy for mCRC; both patients had received oxaliplatin-based chemotherapy for non-mCRC in the adjuvant setting and progressed on therapy before entering the study.

Part 2: Efficacy

Objective response rate. ORR was 31% for patients receiving panitumumab plus rilotumumab versus 21% for patients receiving panitumumab plus placebo (Table 2). No complete responses were noted. For the panitumumab plus rilotumumab arm, the posterior probability that the OR is greater than 1 was 0.93 and therefore met the prespecified criteria that the combination regimen was considered promising.

The ORR was 22% for patients receiving panitumumab plus ganitumab versus 21% for patients receiving panitumumab plus placebo. For the panitumumab plus ganitumab arm, the posterior probability that the OR is greater than 1 was 0.63 and therefore the combination was indeterminate.

Nineteen patients in the panitumumab plus rilotumumab arm (15 of which had an objective response), 12 patients in the panitumumab plus ganitumab arm (10 of which had an objective response), and 15 patients in the panitumumab alone arm (10 of which had an objective response) had a sum of the longest diameter (SLD) reduction >30% on RECIST defined target lesions (Supplementary Fig. S2).

Median durations of response were 5.1 months [95% confidence interval (CI), 3.7–5.6 months], 3.7 months (95% CI, 3.6–5.8 months), and 3.7 months (95% CI, 3.6
to not estimable months) in the panitumumab plus rilotumumab, panitumumab plus ganitumab, and panitumumab plus placebo arms, respectively. The disease control rates were 71%, 61%, and 56% in the panitumumab plus rilotumumab, panitumumab plus ganitumab, and panitumumab plus placebo arms, respectively (Table 2).

**Progression-free survival.** Median PFS for the 3 treatment groups was 5.2 months (95% CI, 3.6–5.4 months), 5.3 months (95% CI, 2.7–5.7 months), and 3.7 months (95% CI, 2.5–5.3 months) in the panitumumab plus rilotumumab, panitumumab plus ganitumab, and panitumumab plus placebo arms, respectively (Table 2). The PFS HR comparing the panitumumab plus rilotumumab arm with the panitumumab plus placebo arm was 0.96 (95% CI, 0.61–1.51) and that for the panitumumab plus ganitumab arm with the panitumumab plus placebo arm was 0.89 (95% CI, 0.52–1.53).

**Overall survival.** OS outcomes were immature at the time of the primary analysis with 40 (28%) deaths reported. An additional follow-up in June 2011 was conducted when the OS data were more mature, with 90 (63%) deaths reported. Median OS for the 3 treatment groups was 13.8 months (95% CI, 11.0–18.0 months), 10.6 months (95% CI, 7.3–14.3 months), and 11.6 months (95% CI, 9.8–18.0 months) in the panitumumab plus rilotumumab, panitumumab plus ganitumab, and panitumumab plus placebo arms, respectively (Fig. 2). The OS HR comparing the panitumumab plus rilotumumab arm with the panitumumab plus placebo arm was 0.89 (95% CI, 0.52–1.53), and that for the panitumumab plus ganitumab arm with the panitumumab plus placebo arm was 1.15 (95% CI, 0.68–1.93). Twenty-seven patients in the panitumumab plus placebo arm (56%) received subsequent therapy with 24 of these patients enrolling in part 3 of this phase II study where they received either subsequent rilotumumab or ganitumab monotherapy. Seven patients (15%) in the panitumumab plus rilotumumab arm and 12 (26%) patients in the panitumumab plus ganitumab arm received subsequent therapy.

**Part 2: Safety**

Twenty percent of patients receiving treatment had experienced at least one serious AE with 19% in the panitumumab plus rilotumumab arm, 20% in the panitumumab plus ganitumab arm, and 23% in the panitumumab plus placebo arm. There were 9 deaths on study: 4 occurred in each of the combination arms and 1 occurred in the panitumumab plus placebo arm. Eight were because of disease progression and 1 was because of staphylococcal sepsis (panitumumab plus ganitumab arm). None were considered related to investigational treatment.

Grade 3 and 4 AEs, including events that occurred with ≥15% occurrence, are shown in Table 3. The safety profiles were similar for the 3 arms with the following exceptions: grade 3/4 hypomagnesemia (highest in the panitumumab alone arm) and grade 3/4 rash (highest in the panitumumab plus rilotumumab arm).

Twelve percent of the total number of rilotumumab doses were withheld and 4% of the ganitumab doses. Sixteen percent, 8%, and 8% of the total number of panitumumab doses were withheld in the rilotumumab plus panitumumab, ganitumab plus panitumumab, and the panitumumab alone arms, respectively. Of patients who had panitumumab withheld, skin- or nail-related toxicities was the reason for 62%, 38%, and 50% of patients in the rilotumumab, ganitumab, and the panitumumab alone arms, respectively.

**Part 2: Phase II biomarker analysis**

Cytoplasmic MET expression levels by IHC were obtained for 47 of 48 patients (98%) in the panitumumab plus rilotumumab (AMG 102) (n = 48) group, 46 of 46 patients (100%) in the panitumumab plus ganitumab (AMG 479) (n = 46) group, and 48 of 48 patients (100%) in the panitumumab plus placebo (n = 48) group.

**Table 2. Primary endpoint: ORR**

<table>
<thead>
<tr>
<th></th>
<th>Panitumumab (AMG 102) (n = 48)</th>
<th>Panitumumab (AMG 479) (n = 46)</th>
<th>Panitumumab (n = 48)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Patients with baseline measurable disease, n (%)</td>
<td>48 (100)</td>
<td>46 (100)</td>
<td>48 (100)</td>
</tr>
<tr>
<td>Objective response, n (%)</td>
<td>15 (31)</td>
<td>10 (22)</td>
<td>10 (21)</td>
</tr>
<tr>
<td>Complete response</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>0 (0)</td>
</tr>
<tr>
<td>Partial response</td>
<td>15 (31)</td>
<td>10 (22)</td>
<td>10 (21)</td>
</tr>
<tr>
<td>Stable disease</td>
<td>19 (40)</td>
<td>18 (39)</td>
<td>17 (35)</td>
</tr>
<tr>
<td>Progressive disease</td>
<td>11 (23)</td>
<td>15 (33)</td>
<td>16 (33)</td>
</tr>
<tr>
<td>Unevaluable/not done</td>
<td>3 (6)</td>
<td>3 (6)</td>
<td>5 (10)</td>
</tr>
<tr>
<td>Disease control ratea, % (95% CI)</td>
<td>71 (56–83)</td>
<td>61 (45–75)</td>
<td>56 (41–71)</td>
</tr>
<tr>
<td>Duration of response, median months (95% CI)</td>
<td>5.1 (3.7–5.6)</td>
<td>3.7 (3.6–5.8)</td>
<td>3.7 (3.6–NE)</td>
</tr>
<tr>
<td>Posterior probability of odds ratio &gt; 1b</td>
<td>0.93</td>
<td>0.63</td>
<td>0.63</td>
</tr>
</tbody>
</table>

Abbreviation: NE, not estimable.

aDisease control rate = complete response + partial response + stable disease.

bOdds ratio is calculated based on ORR; an odds ratio of >1 favors the combination arm over panitumumab alone.
Rilotumumab, Ganitumab, and Panitumumab in Treatment of mCRC

Figure 1. Kaplan–Meier curves for PFS (central review) in patients treated with panitumumab plus rilotumumab (AMG 102; A) and panitumumab plus ganitumab (AMG 479; B) versus panitumumab plus placebo.

Rilotumumab, Ganitumab, and Panitumumab in Treatment of mCRC

rilotumumab arm and 44 of 48 patients (92%) in the panitumumab plus placebo arm (15% of samples were from metastatic lesions). No statistically significant differences were observed for the association of tumor MET cytoplasmic expression with PFS or OS in the panitumumab plus rilotumumab or panitumumab plus placebo arms (Fig. 3). Trends favoring high MET cytoplasmic levels were observed for ORR but the interaction P values indicated no statistically significant predictive treatment effects between the panitumumab plus rilotumumab and panitumumab plus placebo arms (Supplementary Fig. S3).

To identify possible predictive markers for ganitumab treatment, analysis of baseline circulating IGF and IGF-related proteins was performed. An association between lower values for IGFBP1 and IGFBP2 and improved OS was observed for all 3 treatment arms suggesting a prognostic effect. None of the analytes demonstrated a significant interaction with treatment (Supplementary Fig. S4). There were no strong associations between the tested analytes and ORR in ganitumab-treated patients (Supplementary Fig. S5).

There was an association between tumor EGFR staining and improved OS for the panitumumab plus ganitumab arm, and a similar trend was seen in the panitumumab plus placebo arm but the effect was not statistically significant. Several studies have indicated an association of PTEN with anti-EGFR therapy (20, 21). An analysis of PTEN IHC levels suggested a correlation between lower PTEN expression and improved OS but only in the panitumumab plus placebo arm. Taken together, the predictive ability of these biomarkers for the efficacy of the combination of panitumumab
plus rilotumumab or ganitumab was not evident in this study.

Discussion

This is the first phase Ib/II study evaluating the safety and efficacy of the fully human mAbs against HGF/SF, rilotumumab, and IGF1R, ganitumab, in combination with panitumumab in previously treated patients with WT KRAS mCRC. The rationale for this unique trial design was that inhibition of the EGFR pathway, combined with MET or IGF1R pathway inhibition, could provide additive anticancer effects. To our knowledge, the data reported here are the first to suggest promising evidence of efficacy by an HGF (MET pathway) inhibitor (rilotumumab) when combined with an EGFR pathway inhibitor (panitumumab) in patients with WT KRAS mCRC who have progressed on standard chemotherapy.

The primary endpoint of ORR increased by 10 percentage points with the addition of rilotumumab to panitumumab compared with panitumumab alone (31% vs. 21%) and these results were considered statistically significant (per prospectively specified Bayesian criterion). The ORR in the panitumumab alone arm was 21% and similar to that previously reported in the panitumumab monotherapy setting (17% in the WT KRAS subset in trial 20020408; 2). The disease control rate and the duration of response also favored the rilotumumab combination arm. However, the improvement in ORR did not translate into significant PFS and OS benefits.

Figure 2. Kaplan–Meier curves for OS in patients treated with panitumumab plus rilotumumab (AMG 102; A) and panitumumab plus ganitumab (AMG 479; B) versus panitumumab plus placebo.
There was an imbalance of rash in the rilotumumab arm for all grade rash (58%, 52%, and 42% for the rilotumumab, ganitumab, and panitumumab alone arms, respectively) and for grade 3 and 4 rash (29%, 8%, 13% for the rilotumumab, ganitumab, and panitumumab alone arms, respectively). It is possible this may have led to unblinding of the treatment arms. However, all arms received panitumumab for which rash is a known toxicity, no prior or subsequent association of increased rash has been reported for rilotumumab, and it is not clear that the presence of rash alone would provide any strong indication of the treatment arm.

In part 1 (phase Ib) of this study, no DLTs were reported for the first 6 DLT-evaluable patients receiving panitumumab plus rilotumumab. In part 2 (phase II), the safety profile of the drug combination was similar to that of panitumumab plus placebo with the exception of a higher incidence of grade 3/4 rash in the panitumumab plus rilotumumab arm. The median PFS in the panitumumab plus ganitumab arm was similar to what was observed in the rilotumumab-containing arm. However, there was no improvement in OS. Treatment-related grade 3/4 AEs were similar between the arms with the exception of an increase in hypomagnesemia in the panitumumab plus ganitumab arm.

Overall, these results are consistent to those reported elsewhere (22), that the combination of an anti-EGFR and an anti-IGF1R agent do not provide strong additive anti-cancer effects in patients with WT KRAS mCRC. One explanation for these results is that there is compensatory signaling via IGF2 through the insulin receptor (23, 24) which is not downregulated by ganitumab. Additional patient selection markers beyond KRAS status may be required to determine which tumors are most dependent on combined signaling through EGFR and IGF1R as IGF1R expression levels alone do not seem to be sufficient to predict response (22).

Exploratory biomarker analysis was conducted to investigate the correlation of efficacy endpoints with tumor MET, EGFR, and PTEN and baseline circulating IGF1, IGF2, and IGFBP1-3 and -6 protein levels. For the analyzed biomarkers, there was no strong evidence of predictive potential on efficacy endpoints for rilotumumab or ganitumab in combination with panitumumab. Reasons for the lack of conclusive evidence may be attributed to the small number of patients per arm as well as potential imbalances in the arms for RAS mutations beyond KRAS exon 2, which are known negative predictive factors for anti-EGFR mAB therapies, and BRAF mutations, which are known prognostic factors in mCRC. Identification of patients in this study with additional RAS mutations is on-going.

### Table 3. Adverse events (any grade in ≥15% of patients)

<table>
<thead>
<tr>
<th>AE (preferred term), %</th>
<th>Panitumumab + rilotumumab (AMG 102) (n = 48)</th>
<th>Panitumumab + ganitumab (AMG 479) (n = 46)</th>
<th>Panitumumab + placebo (n = 48)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Any AE</td>
<td>Any grade 98</td>
<td>Grade 3/4 71</td>
<td>Any grade 94</td>
</tr>
<tr>
<td>Skin disorders</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Rash</td>
<td>58</td>
<td>29</td>
<td>52</td>
</tr>
<tr>
<td>Acneiform dermatitis</td>
<td>35</td>
<td>15</td>
<td>33</td>
</tr>
<tr>
<td>Pruritus</td>
<td>21</td>
<td>0</td>
<td>25</td>
</tr>
<tr>
<td>Skin fissures</td>
<td>15</td>
<td>2</td>
<td>17</td>
</tr>
<tr>
<td>Paronychia</td>
<td>31</td>
<td>4</td>
<td>15</td>
</tr>
<tr>
<td>Dry skin</td>
<td>23</td>
<td>2</td>
<td>15</td>
</tr>
<tr>
<td>Gastrointestinal disorders</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Constipation</td>
<td>10</td>
<td>0</td>
<td>25</td>
</tr>
<tr>
<td>Decreased appetite</td>
<td>21</td>
<td>2</td>
<td>17</td>
</tr>
<tr>
<td>Nausea</td>
<td>8</td>
<td>0</td>
<td>17</td>
</tr>
<tr>
<td>Vomiting</td>
<td>6</td>
<td>0</td>
<td>15</td>
</tr>
<tr>
<td>Abdominal pain</td>
<td>10</td>
<td>4</td>
<td>15</td>
</tr>
<tr>
<td>Diarrhea</td>
<td>15</td>
<td>4</td>
<td>10</td>
</tr>
<tr>
<td>Fatigue/asthenia</td>
<td>18</td>
<td>4</td>
<td>36</td>
</tr>
<tr>
<td>Hypomagnesemia</td>
<td>29</td>
<td>4</td>
<td>21</td>
</tr>
<tr>
<td>Anemia</td>
<td>4</td>
<td>0</td>
<td>17</td>
</tr>
<tr>
<td>Peripheral edema</td>
<td>19</td>
<td>2</td>
<td>13</td>
</tr>
<tr>
<td>Epistaxis</td>
<td>15</td>
<td>0</td>
<td>2</td>
</tr>
</tbody>
</table>

NOTE: There were nine grade 5 AEs. Four grade 5 AEs occurred in the panitumumab plus rilotumumab arm (all disease progression); 4 in the panitumumab plus ganitumab arm (3 disease progression, one staphylococcal sepsis); 1 in the panitumumab plus placebo arm (disease progression).
To date, review of the literature has not identified a uniform approach to evaluation of the HGF/MET pathway in mCRC. MET is believed to be a late event in carcinogenesis and expression levels may be higher at sites of metastatic disease (11, 12, 25). MET amplification is uncommon and was not measured (26). In this analysis, 85% of the archival tissue samples originated from primary tumors.

This study was not directly planned to examine MET biomarkers, however, results suggest that the MET and EGFR pathways may be related in their impact on outcome. Recent literature in patient-derived colorectal cancer xenograft models, as well as a limited set of patient plasma, demonstrated that MET amplification is associated with EGFR resistance. *In vitro* this resistance may be overcome by MET kinase inhibitors (27). In an updated analysis of a
phase II study in esophagogastric junction cancer (28). OS was 11.5 months in the rilotumumab plus chemotherapy arm and 5.7 months in the chemotherapy alone arm in patients with tumors expressing high MET (HR, 0.34; 95% CI, 0.15–0.78). Hence, for future clinical trial development in the HGF/MET pathway, we encourage the use of MET-positive patient populations. These studies would require baseline tissue from all patients to provide the power needed to address biomarker hypotheses. A specific definition of MET positivity would be one of the objectives as the definition of MET positivity would determine the specific biomarker hypothesis that would be tested.

Although combined inhibition of the HGF/MET pathway in combination with EGFR inhibition appeared promising, there are no current plans to pursue further development of panitumumab plus rilotumumab in mCRC. However, rilotumumab continues to undergo further development in multiple malignancies, including in combination with bevacizumab in recurrent malignant glioma (NCT01113398), in combination with erlotinib in recurrent or progressive non–small cell lung carcinoma (NCT01233687), and in combination with chemotherapy in naïve gastroesophageal patients (NCT01791374, NCT01443065, NCT01697072).

These data support further study of combined biologic blockade of cancer-related pathways without the requirement for cytotoxic agents and emphasize the need to identify patients that may benefit from combined EGFR and MET pathway inhibition.

Disclosure of Potential Conflicts of Interest

E. Van Cutsem reports receiving a commercial research grant from Amgen Inc. C. Eng and N. Tebbutt report receiving commercial research grants from and are consultants/advisory board members for Amgen Inc. I. McCaffery and D. Smethurst were employees of and have ownership interest (including patents) in Amgen Inc. K. Oliner, L. Chen, J.L. Gansert are employees of Amgen Inc. E. Lob is an employee of and has ownership interest (including patents) in Amgen Inc. J. Tabernero is a consultant/advisory board member for Amgen Inc. No potential conflicts of interest were disclosed by the other authors.

Authors’ Contributions


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Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): C. Eng, E. Nowara, H. Prenen, I. Gansert


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References


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