

Initial Report of Second-Line FOLFIRI in Combination with Ramucirumab in Advanced Gastroesophageal Adenocarcinomas: A Multi-Institutional Retrospective Analysis

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Disclosures of potential conflicts of interest may be found at the end of this article.

Key Words. Gastric cancer • GEJ adenocarcinoma • Ramucirumab • FOLFIRI • Second line

ABSTRACT

Background. The randomized phase III RAINBOW trial established paclitaxel (pac) plus ramucirumab (ram) as a global standard for second-line (2L) therapy in advanced gastric and gastroesophageal junction adenocarcinoma, together gastroesophageal adenocarcinoma (GEA). Patients (pts) receiving first-line (1L) FOLFOX often develop neuropathy that renders continued neurotoxic agents in the 2L setting unappealing and other regimens more desirable. As such, FOLFIRI-ram has become an option for patients with 2L GEA. FOLFIRI-ramucirumab (ram) has demonstrated safety and activity in 2L colorectal cancer, but efficacy/safety data in GEA are lacking.

Subjects, Materials, and Methods. Patients with GEA treated with 2L FOLFIRI-ram between August 2014 and April 2018 were identified. Clinicopathologic data including oxaliplatin neurotoxicity rates/grades (G), 2L treatment response, progression-free survival (PFS), overall survival (OS), safety, and molecular features were abstracted from three U.S.

academic institutions. Kaplan-Meier survival analysis was used to generate PFS/OS; the likelihood ratio test was used to determine statistical significance.

Results. We identified 29 pts who received 2L FOLFIRI-ram. All pts received 1L platinum + fluoropyrimidine, and 23 of 29 (79%) had post-1L neuropathy; 12 (41%) had G1, and 11 (38%) had G2. Patients were evenly split between esophagus/gastroesophageal junction (12; 41%) and gastric cancer (17; 59%). Among evaluable pts (26/29), the overall response rate was 23% (all partial response) with a disease control rate of 79%. Median PFS was 6.0 months and median OS was 13.4 months among all evaluable pts. Six- and 12-month OS were 90% ($n = 18/20$) and 41% ($n = 7/17$). There were no new safety signals.

Conclusion. We provide the first data suggesting FOLFIRI-ram is a safe, non-neurotoxic regimen comparing favorably with the combination of pac + ram used in the seminal RAINBOW trial. *The Oncologist* 2019;24:475–482

Implications for Practice: Results of this study provide initial support for the safety and efficacy of second-line (2L) FOLFIRI-ramucirumab (ram) after progression on first-line platinum/fluoropyrimidine in patients with gastroesophageal adenocarcinoma (GEA). The overall response, progression-free survival, overall survival, and toxicity profile compare favorably with paclitaxel (pac) + ram and highlight the importance of the ongoing phase II RAMIRIS trial examining FOLFIRI-ram versus pac + ram in 2L GEA (NCT03081143). FOLFIRI-ram may warrant consideration for inclusion as an alternate regimen in consensus guidelines for GEA.

INTRODUCTION

Gastroesophageal adenocarcinoma (GEA) accounts for over 1.1 million annual cancer-related deaths, representing nearly 14% of global cancer deaths [1]. Despite improved molecular characterization, the prevailing international

standard for advanced GEA remains doublet therapy with platinum-based and fluoropyrimidine (5FU)-based combination regimens [2, 3]. The addition of trastuzumab to first-line (1L) platinum/5FU in HER2-positive patients improves

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Table 1. Clinicopathologic features in a cohort of GEA patients treated with second-line FOLFIRI-ram

Clinicopathologic feature	n = 29, n (%)
Age at 2L initiation, median (range), yr	61.5 (36–80)
Sex	
Male	21 (72.4)
Female	8 (27.6)
ECOG at 2L initiation	
0	12 (45.2)
1	14 (48.0)
2	2 (6.9)
3	1 (3.4)
Ethnicity	
Hispanic/Latino	7 (24.1)
White	16 (55.0)
Black	3 (10.3)
Asian	2 (6.9)
Other	1 (3.4)
Tumor location	
Esophagus	4 (13.8)
GEJ	8 (27.6)
Gastric	17 (58.6)
Stage at diagnosis	
II	1 (3.4)
III	6 (20.7)
IV	22 (75.9)
Lauren histology	
Diffuse	14 (48.3)
Intestinal	13 (44.8)
NOS	2 (6.9)
Histologic grade	
1 (well differentiated)	1 (3.4)
2 (moderately differentiated)	6 (20.7)
3 (poorly differentiated)	22 (75.9)
Signet ring cell features	
Yes	14 (48.3)
No	15 (51.7)
Ascites	
Yes	13 (44.8)
No	16 (55.2)
Metastatic disease sites	
Visceral	15 (51.7)
Lymph node	20 (70.0)
Radiographic peritoneal	13 (44.8)
HER2 status at diagnosis	
IHC 0	17 (54.8)
IHC 1+	4 (12.9)
IHC 2+	4 (12.9)
IHC 2+, FISH/NGS amp	1
IHC 2+, FISH/NGS non-amp	3
IHC 3+	1 (3.2)
Unknown	2 (6.5)

Abbreviations: 2L, second-line; amp, amplification; ECOG, Eastern Cooperative Oncology Group; FISH, fluorescence in situ hybridization; GEA, gastroesophageal adenocarcinoma; GEJ, gastroesophageal junction; IHC, immunohistochemistry; NGS, next-generation sequencing; NOS, not otherwise specified; ram, ramucirumab.

survival in this molecularly selected subset [4]. Among U.S. patients there is significant 1L regimen heterogeneity, with FOLFOX being the most common platinum/5FU combination. Following 1L therapy, U.S. claims analyses suggest only 42%–54% of patients with GEA receive second-line (2L) therapy [5, 6]. The phase III RAINBOW trial established the combination of paclitaxel (pac) and the anti-vascular endothelial growth factor (VEGF) receptor 2 antibody ramucirumab (ram) as a global 2L standard after platinum/5FU failure [7]. However, the rates of neurotoxicity with 1L FOLFOX in GEA range from 30%–70%, making 2L taxane-containing therapy less appealing to a significant portion of patients with GEA [8, 9]. Moreover, the use of prior taxane regimens in the 1L setting, or rapid recurrence after perioperative FLOT therapy, makes determining an optimal “2L” taxane-free regimen of critical importance [10–13]. Irinotecan as either monotherapy or in combination with 5FU (FOLFIRI) has established activity in GEA as both a 1L and 2L option [14, 15]. A small ($n = 6$) Japanese phase Ib trial of irinotecan plus ramucirumab (no 5FU) in more heavily pretreated gastric cancer demonstrated safety and a response rate of 17% (1/6) [16]. Additional support for the safety and efficacy of FOLFIRI in combination with ramucirumab (FOLFIRI-ram) was established in a phase III trial in 2L advanced colorectal cancer after progression on FOLFOX with bevacizumab [17].

Despite the lack of data, FOLFIRI-ram has emerged as a reasonable 2L option for a significant portion of patients with advanced disease across U.S. institutions. We therefore sought to examine the safety and efficacy of 2L FOLFIRI-ram across a clinically annotated and molecularly characterized U.S. cohort with GEA.

SUBJECTS, MATERIALS, AND METHODS

Patients with advanced GEA who were treated with 2L FOLFIRI-ram were retrospectively identified from three tertiary U.S. medical centers. All patients fulfilled prespecified inclusion criteria, including histologically confirmed advanced GEA, 1L platinum + 5FU-based chemotherapy, receipt of at least two cycles of FOLFIRI-ram, and available clinicopathologic details. All patients were treated in accordance with the Declaration of Helsinki, and retrospective review was approved by the respective local institutional review board. Clinicopathologic details were abstracted, Common Terminology Criteria for Adverse Events version 4.0 was used to grade adverse events, and RECIST v1.1 per investigator assessment was utilized for response assessment. HER2 testing was performed locally at diagnosis per American Society of Clinical Oncology (ASCO)/College of American Pathologists (CAP) guidelines. Where possible, programmed death ligand 1 (PD-L1) status was determined using the Food and Drug Administration-approved pharmDx (Agilent; Santa Clara, CA) immunohistochemical (IHC) assay (PD-L1 IHC 22C3) combined positive score as previously described [18]. Comprehensive genomic profiling was performed using hybrid-capture-based genomic profiling as previously described (FoundationOne; Foundation Medicine, Cambridge, MA) [19].

Overall survival (OS) was calculated from the date of initiation of 2L therapy to death. Progression-free survival (PFS)

Table 2. First-line treatment outcomes in a cohort of GEA patients treated with second-line FOLFIRI + ram

Treatment feature	n = 29, n (%)
First-line therapy	
FOLFOX/XELOX backbone	23 (79.3)
FOLFOX + trastuzumab	2 (6.8)
FOLFOX + experimental agent (trial)	12 (41.3) ^a
Other 5FU + platinum	6 (20.7)
5FU + platinum-containing triplets	6 (20.7)
FLOT	3 (10.3)
DOF	1 (3.4)
Modified DCF	1 (3.4)
ECX + placebo/experimental agent (trial)	1 (3.4) ^a
Duration of 1L therapy, median (range), mo	7 (2–12)
Objective response to 1L	
Complete response	1 (3.4)
Partial response	13 (44.8)
Stable disease	10 (34.4)
Progressive disease	3 (10.3)
Unknown or nonmeasurable	2 (6.9)
Maximal neuropathy grade	
0	6 (20.7)
1	12 (41.4)
2	11 (37.9)

^aDenotes participation in a first line clinical trial. Total 45% (13/29) patients were treated on a first line clinical trial
Abbreviations: 1L, first-line; 5FU, fluoropyrimidine; DCF, docetaxel, cisplatin, 5-fluorouracil; DOF, docetaxel, oxaliplatin, 5-fluorouracil; ECX, epirubicin, cisplatin, and capecitabine.

was calculated from both the start of 1L therapy to disease progression and the start of 2L FOLFIRI-ram to progression or death. Times to event (PFS/OS) were estimated using the Kaplan-Meier method and compared using log-rank and Cox proportional hazards, with the likelihood ratio used to determine statistical significance. For genomic analyses, PFS and OS Cox univariate and multivariate regression models were developed to analyze the role of individual alterations and predetermined pathway activation on outcomes. All inferential analyses utilized two-sided methods ($\alpha = 0.05$), and statistical significance was determined when p value was $<.05$. All statistical analyses were conducted using R version 3.4.2.

RESULTS

A total of 29 patients met the prespecified inclusion criteria and received 2L FOLFIRI-ram between August 2014 and April 2018, and 26 of 29 patients were evaluable for PFS and OS analyses. Baseline demographic and pathologic features are shown in Table 1. The majority (75%) presented with stage IV disease, and poor prognostic features, including peritoneal disease (45%), ascites (45%), and signet ring cell features (45%), were common. Forty-five percent of patients participated in first-line clinical trials, and all patients received first-line platinum plus 5-fluorouracil (5FU)-containing regimens, with the majority (79%) receiving FOLFOX/XELOX (Table 2). The overall response rate to

Table 3. Response characteristics and toxicity profile in a cohort of advanced GEA patients receiving second line FOLFIRI in combination with ramucirumab

Treatment feature	(n = 29), n (%)
Second-line therapy	
FOLFIRI + ram	29 (100.0)
FOLFIRI-ram doses, median (range)	8 (2–40)
Objective response to 2L FOLFIRI + ram	
Complete response	0 (0.0)
Partial response	6 (20.7)
Stable disease	17 (58.6)
Progressive disease	5 (17.2)
Unknown or nonmeasurable	1 (3.4)
6-month OS rate (n = 20)	18 (90)
12-month OS rate (n = 17)	7 (41)
Reason for 2L FOLFIRI + ram	
1L neuropathy	6 (20.7)
1L taxane with platinum/5FU	5 (17.2)
Perceived efficacy	18 (62.1)
Other	0 (0.0)
Dose reductions needed at any time	
Yes	17 (58.6)
No	12 (41.4)
Adverse events of interest	
Diarrhea grade 1–2	9 (31.0)
Diarrhea grade >2	0 (0.0)
Anemia grade 1–2	7 (24.1)
Anemia grade >2	0 (0.0)
Neutropenia, any grade	4 (13.8)
Fatigue grade 1–2	20 (69.0)
Fatigue grade >2	2 (6.9)
Hypertension grade 1–2	2 (6.9)
Hypertension grade >2	0 (0.0)
Grade 3–4 adverse event of any type	2 (6.9)

Abbreviations: 1L, first-line; 2L, second-line; 5FU, fluoropyrimidine; GEA, gastroesophageal adenocarcinoma; OS, overall survival; ram, ramucirumab.

1L therapy was 48% with a median 1L therapy duration of 7 months, consistent with phase II–III platinum/5FU containing 1L trials (Table 2) [8, 9, 20, 21]. Post-1L neuropathy was seen in 79% of patients, and nearly 40% had grade 2 neuropathy.

The most common reason for selecting second-line FOLFIRI-ram was perceived efficacy (62%), followed by persistent post-1L neuropathy (21%) and receipt of 1L taxane (17%; Table 3). Amongst the 26 evaluable patients, median PFS and OS for 2L FOLFIRI-ram were 6 and 13.4 months, respectively (Fig. 1). Progression-free survival ranged from 2 to 24 months, and 2 patients remained on therapy at the time of censoring (June 1, 2018). The RECIST v1.1 disease control rate was 79% (23/26), and objective response rate in evaluable patients was 23% (6/26), consisting entirely of partial responses. Of note, all six patients with objective responses lacked peritoneal disease, and four of six had

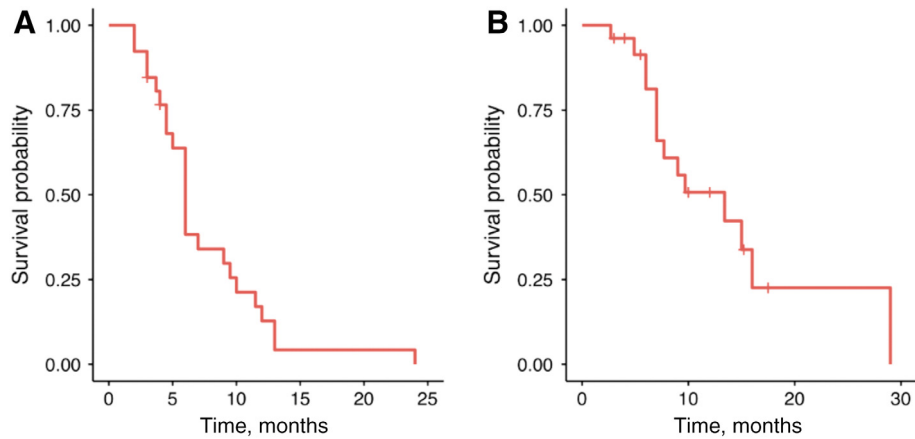


Figure 1. Median progression-free survival was 6 months (A), and median overall survival was 13.4 months (B) across the 26 evaluable patients.

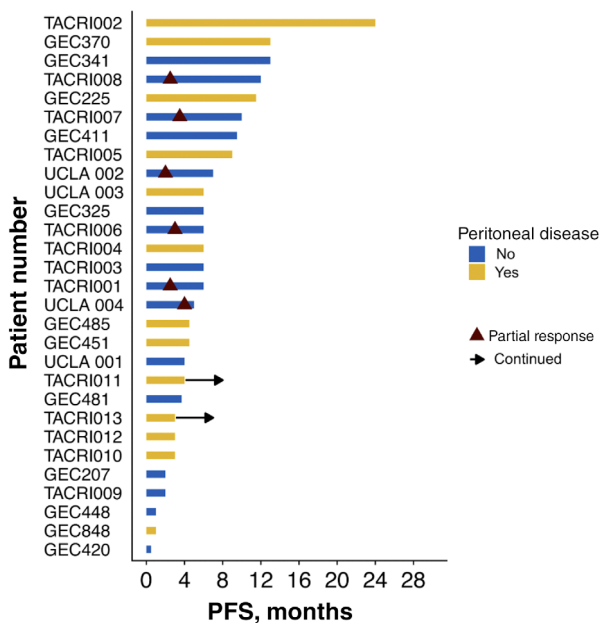


Figure 2. Swimmer plot of progression free survival by patient demonstrating RECIST responses only occurring in patients without peritoneal involvement. Abbreviation: PFS, progression-free survival.

proximal disease (Fig. 2). Six- and 12-month OS were 90% ($n = 18/20$) and 41% ($n = 7/17$), respectively. Toxicities were largely grade 1–2, with only 6.9% developing grade 3–4 adverse events (all grade 3 fatigue). Fatigue (75.6%), diarrhea (31%), anemia (24.1%), and neutropenia (13.8%) were the most common adverse events, and there were no toxic deaths (Table 3). Adverse events potentially associated with the VEGF pathway were uncommon, with 6.9% (2/29) of patients developing grade 1–2 hypertension and no bleeding events. Rates of proteinuria were not well captured in this retrospective cohort; however, there were no reported cases of nephrotic syndrome.

Univariate analysis identified a significantly better median PFS of 7.5 versus 5 months in intestinal versus diffuse type histology ($n = 12/24$; $p = .02$; 95% confidence interval [CI], 0.1–0.9; Fig. 3F) and inferior median PFS in

patients with predominantly signet cells ($n = 3$; $p = .05$; 95% CI, 1.1–57.3; Fig. 3G). No significant PFS differences were seen when stratified by age, location, HER2 status, PD-L1 expression, or ascites or peritoneal disease. Patients with first-line therapy responses of 6 or more months had a median OS of 15 versus 7 months compared to those with shorter first-line responses (online supplemental Fig. 1D). There was no significant association of patient age, presence of signet cells, ascites, or peritoneal disease with median overall survival.

All patients had molecular characterization, with 28 of 29 (96.6%) patients undergoing genomic profiling. All patients with available results (next-generation sequencing and/or IHC) were microsatellite stable, and 20% (4/20 tested) were PD-L1 positive. Molecular testing results and impact on FOLFIRI-ram PFS and OS are shown in Table 4. Among included genes, there were no statistically significant associations between individual gene alteration or pathway and clinical outcomes upon univariate analysis (data not shown). However, upon multivariate pathway analysis, including alterations in Wnt, FGF, DDR, or MAPK, the “wildtype Wnt” (i.e., no Wnt pathway genomic alterations) covariate suggested a median OS benefit ($p = .01$; data not shown).

DISCUSSION

Here we provide the first report of 2L FOLFIRI-ram in patients with advanced GEA treated with 1L platinum/5FU-based chemotherapy. Although limited by its retrospective nature, our analyses suggest FOLFIRI-ram is a safe and non-neurotoxic 2L regimen with favorable activity.

Since the 2014 approval of paclitaxel with ram, there have been no significant advancements in the 2L treatment of GEA, including HER2-directed therapies and immunotherapy [7, 22, 23]. Outside molecular subgroups such as microsatellite instability high, pac + ram remains the standard 2L therapy. Early support for 2L FOLFIRI was observed in the phase III epirubicin, cisplatin, and capecitabine (ECX) versus FOLFIRI 1L trial, in which 48% of patients receiving first-line ECX received 2L FOLFIRI. Among the 2L FOLFIRI

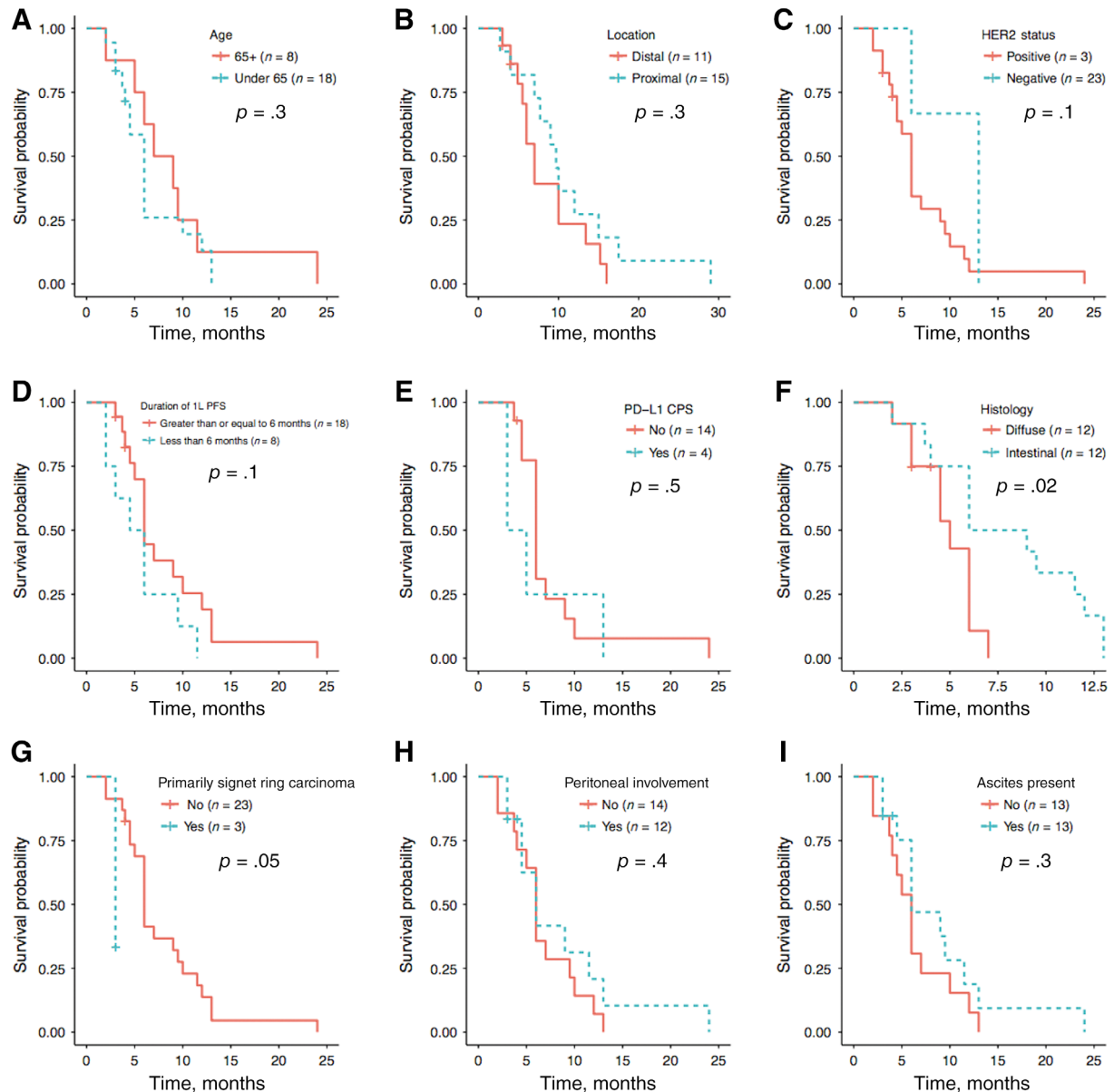


Figure 3. Patients 65 and older ($n = 8/26$) had an 8-month median progression-free survival (mPFS) versus 6 months in younger patients ($p = .3$; 95% confidence interval [CI], 0.6–3.7) (A). Patients with proximal gastroesophageal adenocarcinoma (GEA; $n = 11/26$) trended towards mPFS of 9.7 versus 7 months ($p = .3$; 95% CI, 0.3–1.5) (B). HER2 positive patients ($n = 3/26$) had an mPFS of 13 versus 6 months, although this was not statistically significant ($p = .1$, 95% CI, 0.1–1.5) (C). Those with first-line therapy mPFS <6 months ($n = 8/26$) had a second-line mPFS of 5.3 months versus 6 months for those with a 6 month or greater first-line PFS ($p = .1$, 95% CI, 0.8–4.9) (D). PD-L1 negative patients ($n = 14/18$) tested by immunohistochemistry had a nonsignificant mPFS benefit of 6 versus 4 months ($p = .5$; 95% CI, 0.5–4.8) (E). Median progression-free survival was 7.5 versus 5 months in patients with intestinal ($n = 12/24$) versus diffuse type histology ($n = 12/24$; $p = .02$; 95% CI, 0.1–0.9) (F). Similarly, 3 of 24 patients with known histology had predominantly signet cell carcinoma, and this portended a 3- versus 6-month mPFS ($p = .05$, 95% CI, 1.1–57.3) (G). Median PFS was 6 months in patients with peritoneal involvement ($n = 12/26$; $p = .4$; 95% CI, 0.3–1.7) (H) and ascites ($n = 13/26$; $p = .3$, 95% CI, 0.3–1.5) (I).

Abbreviations: 1L, first-line; CPS, combined positive score; PD-L1, programmed death ligand 1; PFS, progression-free survival.

patients, the overall response rate (ORR) was 13.7%, although second-line PFS2 was not reported [14]. Pilot support for irinotecan with ramucirumab was also shown in a small phase Ib ($n = 6$) Japanese trial [16]. Phase III trials utilizing paclitaxel monotherapy control arms reliably demonstrate ORR of 13%–20%, with PFS ranging from 3–4 months [7, 22, 23]. Although we cannot directly compare our retrospective data to prospective clinical trial data, our

ORR of 23% and median PFS of 6 months compare favorably with 2L GEA trials. Landmark survival analyses, specifically 6- and 12-month OS rates, are meaningful endpoints, and the 6- and 12-month OS rates of 90% and 41% observed in our cohort are well aligned with the RAINBOW results (72% and 40%, respectively) [7].

Treatment-associated toxicities impact the quality of life among patients with GEA, and the balance of efficacy

Table 4. Genomic features among evaluable GEA patients who received 2L FOLFIRI-ram

Genomic feature	No. altered samples (n = 26), n (%)	mPFS (HR), mo
HER2 NGS amplification		
Copy number ≥ 8	2/25 (6.9)	9.5 (0.14–2.6)
PD-L1 IHC		
CPS >1	4/18 (22.2)	4 (0.46–4.8)
Oncogenic pathways		
FGF altered	2 (8)	4.9 (0.58–12.0)
MAPK altered	18 (72)	7.8 (0.5–2)
DDR altered	7 (28)	9.5 (0.2–1.4)
Wnt/catenin altered	16 (64)	6 (0.33–1.9)

All samples were sequenced using the FoundationOne comprehensive genomic profiling assay. Sequencing data is missing from one patient.

Abbreviations: CPS, combined positive score; DDR, DNA damage response; FGF, fibroblast growth factor; GEA, gastroesophageal adenocarcinoma; HR, hazard ratio; IHC, immunohistochemistry; mPFS, median progression-free survival; NGS, next-generation sequencing; PD-L1, programmed cell death ligand 1; ram, ramucirumab.

and tolerance is particularly important in later lines of therapy. The adverse-event related treatment discontinuation rate was 12% for pac + ram with grade 3 events in 47% of patients [7]. We observed a favorable toxicity profile with FOLFIRI-ram with grade 3 adverse events in 7%, although 58% of our patients required dose reduction during therapy. Admittedly, these comparisons should be considered exploratory, and the ongoing phase II RAMIRIS trial (NCT03081143) will provide key data.

The clinicopathologic features among our patients are representative of U.S. patients with GEA. Consistent with prior reports, we observed lesser benefit among patients with diffuse type histology and signet ring cell features [7, 24]. The exact biologic rationale for this observation is unknown, and predictive and prognostic genomic biomarkers are essential to improving outcomes in GEA. Molecularly defined subsets including *HER2* amplification, *EGFR* amplification, *MET* amplification, and microsatellite instable, PD-L1 positive, and EBV-positive tumors have met with variable success when matched to targeted and/or immune therapies [4, 9, 18, 25–31]. To date, there have been no biomarkers clearly associated with ram in GEA, although high VEGF-D (>115 pg/mL) may identify a group with improved outcomes in the RAISE trial [32]. In the present cohort, exploratory analyses failed to identify any genomic biomarkers differing significantly between responders and nonresponders (Table 1). Interestingly, 64% of evaluable patients harbored Wnt-pathway gene alterations, and multivariate analysis suggests absence of Wnt alterations was associated with benefit, consistent with limited prior reports [28].

Notably, 45% of the patients here were enrolled in first-line clinical trials, including 7 of 13 (54%) who received first-line antiangiogenic therapy. In the exploratory analysis from the 1L phase III RAINFALL trial (cisplatin and 5-fluorouraci + ramucirumab/placebo), there

was a suggestion that more favorable OS was achieved in patients receiving 1L ram and subsequent ram-containing therapy [33]. Within our limited dataset, there was no improved PFS or OS among patients receiving 1L therapy with antiangiogenic agents followed by FOLFIRI-ram (data not shown). In fact, there was a trend toward inferior outcomes, although selection bias (more aggressive biology in 7/13 1L trial patients on antiangiogenic therapy) may confound this very small sample size.

There are several potential factors limiting the generalizability of our results, including the retrospective nature of our study. All patients received therapy at high-volume GEA centers with multidisciplinary approaches, trial availability, and aggressive supportive care, which may contribute to improved outcomes [34]. Nearly all patients (89.6%) had an Eastern Cooperative Oncology Group performance score of 0–1 at the initiation of 2L FOLFIRI-ram, and 45% were treated on a 1L clinical trial. However, poor prognostic features were well represented, with over 40% of patients with peritoneal disease, diffuse-type disease, and/or ascites and 75.9% with poorly differentiated tumors. Indeed, in this cohort, diffuse-type tumors had worse outcomes than nondiffuse type, confirming the negative prognostic connotation of diffuse-type disease. The retrospective nature of our findings requires cautious interpretation, although the 1L features and outcomes are well aligned with prospective data, providing some validation for our findings.

CONCLUSION

We provide initial support for the safety and efficacy of 2L FOLFIRI-ram after progression on 1L platinum/5FU in patients with GEA. To our knowledge, this represents the only existing data demonstrating favorable activity and safety in comparison with standard second-line pac + ram. Our preliminary analysis highlights the importance of the ongoing phase II RAMIRIS trial examining FOLFIRI-ram versus pac + ram in 2L GEA (NCT03081143). FOLFIRI-ram may warrant consideration for inclusion as an alternate regimen in consensus guidelines for GEA.

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DISCLOSURES

Samuel J. Klempner: Foundation Medicine, Inc. (H); Astellas Pharma, Lilly Oncology (C/A); Merck, Leap Therapeutics, Incyte (institutional RF); **Zev A. Wainberg:** Genentech/Roche, Amgen, Eli Lilly & Co., Five Prime, EMD Serono, Bayer (RF); **Daniel V.T. Catenacci:** Lilly Oncology, Amgen, Taiho, Five Prime, Merck, Bristol-Myers Squibb, Astellas, Roche/Genentech, Foundation One, Guardant Health (C/A, SAB, H). The other authors indicated no financial relationships.

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See <http://www.TheOncologist.com> for supplemental material available online.

For Further Reading:

Benjamin A. Weinberg, Joanne Xiu, Jimmy J. Hwang et al. Immuno-Oncology Biomarkers for Gastric and Gastroesophageal Junction Adenocarcinoma: Why PD-L1 Testing May Not Be Enough. *The Oncologist* 2018;23:1171–1177.

Implications for Practice:

Pembrolizumab is approved by the U.S. Food and Drug Administration for patients with refractory gastric and gastroesophageal cancers if the tumor and adjacent tissue stain positively for the programmed cell death ligand 1 (PD-L1) protein by companion diagnostic testing. Tumor mutational load, microsatellite instability (MSI), and alternative PD-L1 testing thresholds may serve as predictive biomarkers for response to immune checkpoint inhibition, and standard PD-L1 testing will not identify all patients who may benefit from this therapy.