

Treatment-related severe and fatal adverse events with molecular targeted agents in the treatment of advanced gastric cancer: a meta-analysis

Liang Wang
Yagang Liu
Wenyong Zhou
Wei Li

Department of General Surgery,
The Central Cangzhou Hospital,
Cangzhou, Hebei Province, People's
Republic of China

Aim: To perform a systematic review and meta-analysis of Phase III randomized controlled trials (RCTs) to determine the incidence and risk of severe adverse events (AEs) with molecular targeted agents (MTAs) in advanced/metastatic gastric cancer (GC) patients.

Methods: A comprehensive literature search for related trials published up to December 2015 was performed. Eligible studies were Phase III RCTs of advanced/metastatic GC patients assigned to MTAs or control group. Data were extracted by two authors for severe and fatal AEs (FAEs).

Results: A total of nine Phase III RCTs involved 4,934 GC patients were ultimately identified. The pooled results demonstrated that the addition of TAs to therapies in advanced GC significantly increased the risk of developing severe AEs (relative risk: 1.12, 95% confidence interval: 1.02–1.24, $P=0.02$), but not for FAEs (relative risk: 0.97, 95% confidence interval: 0.65–1.45, $P=0.88$). Additionally, the most common causes of FAEs with MTAs were infections (16.3%), gastrointestinal hemorrhage (8.2%), and arterial thromboembolic events (8.2%), respectively.

Conclusion: With available evidence, the use of TAs in GC patients was associated with an increased risk of severe AEs, but not for FAE. Clinicians should be aware of the risk of severe AEs with the administration of these drugs in these patients.

Keywords: advanced gastric cancer, molecular targeted agents, randomized, meta-analysis

Introduction

Gastric cancer (GC) is one of the most common malignant diseases worldwide, accounting for 8% (989,600 million) of the total new cancer cases and 10% (738,000) of the total cancer deaths in 2008.¹ Generally, GC is a heterogeneous disease, which usually includes different subgroups according to histological, anatomical, genomic, or molecular classifications.^{2,3} Regardless of these subtypes, the current treatment of GC is based on a multidisciplinary approach that combines gastrectomy, radiotherapy, and chemotherapy.⁴ Despite the advances in the treatment, nearly 50% of patients with locally advanced-stage GC relapse after gastrectomy.^{5,6} For such patients, palliative chemotherapy is the mainstay treatment to prolong the survival. Currently, combination chemotherapy based on 5-fluoropyrimidines/platinum, with possible addition of docetaxel in fit patients, represent the landmark of first-line treatment of advanced GC patients. However, the efficacy of first-line chemotherapy is modest, with a median survival 8–12 months,^{7–9} and most patients are nonresponders or eventually experience disease progression. Thus, it is clear that novel treatments are badly needed in advanced GC patients.

Correspondence: Liang Wang
Department of General Surgery, The
Central Cangzhou Hospital, Cangzhou,
Hebei Province 061000, People's
Republic of China
Tel +86 0317 207 5646
Fax +86 0317 207 5647
Email wangliang201601@sina.com

During the past decades, a better understanding of the molecular events involved in the tumorigenesis of GCs has led to development of new targeted agents.¹⁰ A recent meta-analysis conducted by Qi et al¹¹ demonstrated that the use of anti-vascular endothelial growth factor (anti-VEGF) agents provided a significant survival benefit in previously treated GC patients. Another meta-analysis conducted by Ciliberto et al¹² also showed that antiangiogenic agents (hazard ratio [HR]: 0.759; 95% confidence interval [CI]: 0.655–0.880; $P < 0.001$) and anti-HER-2 agents (HR: 0.823; 95% CI: 0.722–0.939; $P = 0.004$) significantly improve overall survival, while no benefit was found for anti-EGFR agents (HR: 1.077; 95% CI: 0.847–1.370; $P = 0.543$). To date, two molecular targeted agents (MTAs) targeting VEGF signal pathway, bevacizumab and ramucirumab,¹³ and one MTA-targeting EGFR signal pathway, trastuzumab,¹⁴ have been approved for use in advanced GC patients due to survival benefits. Therefore, it is anticipated that the use of these MTAs in GC would increase in the future. However, VEGF and EGFR play multiple roles in physiologic processes, and thus their inhibition could have potentially serious systemic consequences. To our best knowledge, there is no specific meta-analysis to assess the severe adverse events (AEs) and fatal adverse events (FAEs) associated with MTAs in GC. We, therefore, conduct this comprehensive meta-analysis of Phase III randomized controlled trials (RCTs) to assess the toxicities of MTAs in advanced GC patients.

Materials and methods

Study design

We performed this meta-analysis according to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses statements.¹⁵

Search strategy

In December 2015, an extensive search of the following databases was performed: Embase, Medline, the Cochrane Central Register of Controlled Trials, and the Cochrane Database of Systematic Reviews. The following keywords were used: “gastric neoplasms,” “gastric cancer,” “gastric carcinoma,” “sorafenib,” “unitinib,” “pazopanib,” “axitinib,” “cediranib,” “regorafenib,” “ramucirumab,” “vandetanib,” “bevacizumab,” “angiogenesis inhibitor,” “mTOR inhibitor,” “everolimus,” “cetuximab,” “panitumumab,” “lapatinib,” “trastuzumab,” “molecular targeted agents,” and “randomized controlled trials.” The language of publication and years were not limited.

Selection of trials

Clinical trials that met the following criteria were included: 1) Phase III RCTs in patients with pathologically confirmed GC; 2) participants assigned to treatment with or without MTAs; and 3) reported outcomes of interest (ie, severe AE and FAEs). We used the five-item Jadad scale including randomization, double-blinding, and withdrawals as previously described to approximately assess the quality of included trials.¹⁶

Data extraction

Two investigators independently performed data extraction. If reviewers suspected an overlap of cohorts in a report, they contacted the corresponding author for clarification; we excluded studies with a clear overlap. The following information was recorded for each study: first authors' name, year of publication, study period, median age, MTAs dosage, number of patients enrolled, and events of severe and FAEs. The primary end point of this study was FAE, which was defined by the National Cancer Institute's Common Terminology Criteria for Adverse Events as deaths occurring during a clinical trial as a result of exposure to an experimental drug. We did not include FAEs that were related to disease progression. The second end point of this study was severe AE, which was defined by the National Cancer Institute's Common Terminology Criteria for Adverse Events as Grade 3 or 4 toxicities occurring during a clinical trial as a result of exposure to an experimental drug.

Statistical analysis

Statistical analysis of severe and fatal AEs was calculated using comprehensive meta-analysis software version 2.0 (Biostat, Englewood, NJ, USA). Between-study heterogeneity was estimated using the χ^2 -based Q statistic.¹⁷ Heterogeneity was considered statistically significant when $P_{\text{heterogeneity}} < 0.05$ or $I^2 > 50\%$. We calculated the pooled relative risk (RR) and 95% CIs by using random-effect or fixed-effect models according to the heterogeneity of included studies. A statistical test with a P -value less than 0.05 was considered significant. $RR > 1$ indicates more toxicities in MTAs group, and vice versa. Finally, publication bias was evaluated through funnel plots and with Begg and Egger's tests.^{18,19}

Results

Search results

Our literature search revealed 252 potential relevant publications, and 21 reports were retrieved for full-text evaluation;

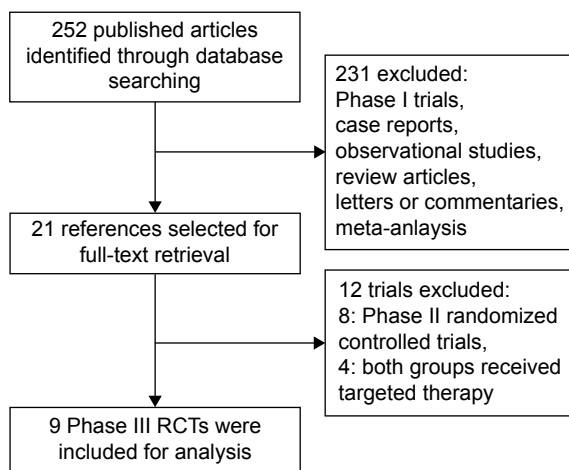


Figure 1 Studies eligible for inclusion in the meta-analysis.

Abbreviation: RCTs, randomized controlled trials.

the reasons for study exclusion are illustrated in Figure 1. Finally, a total of nine RCTs with 4,934 patients were included.^{20–28} The baseline characteristics of each trial are listed in Table 1. Five trials were performed in first-line settings and four in second-line settings (Table 1).^{20,22,23,25} The quality of each included study was approximately assessed according to Jadad scale; five trials were placebo-controlled, double-blinded randomized trials, and thus had Jadad score of 5.^{20,21,23–28} The other four trials had Jadad score of 3.

Table 1 Baseline characteristics of nine included trials

Study	Treatment line	Total (N)	Treatment arms	Number for analysis	Median age (years)	Median PFS	Median OS	Jadad score
Fuchs et al ²³ (2014)	Second-line	335	Ramucirumab 8 mg/kg Placebo	236 115	60 60	NR NR	5.2 3.8	5
Wilke et al ²⁰ (2014)	First-line	655	Ramucirumab 8 mg/kg + PTX Placebo + PTX	327 329	61 61	4.4 2.9	9.6 7.4	5
Shen et al ²¹ (2015)	Second-line	202	Bevacizumab 2.5 mg/kg/wk + capecitabine + DDP Placebo + capecitabine + DDP	100 102	54.2 55.5	6 6.3	11.4 10.5	5
Satoh et al ²² (2014)	Second-line	261	Lapatinib 1,500 mg qd + PTX PTX	132 129	61 62	5.4 4.4	11 8.9	3
Waddell et al ²⁴ (2013)	First-line	553	Panitumumab + EOC EOC	278 275	63 62	7.4 6	11.3 8.8	3
Ohtsu et al ²⁵ (2013)	Second-line	656	Everolimus 10 mg/d Placebo	439 217	62 62	1.7 1.4	5.4 4.3	5
Lordick et al ²⁶ (2013)	First-line	904	Cetuximab + capecitabine + DDP Capecitabine + DDP	455 449	60 59	4.4 5.6	9.4 10.7	3
Ohtsu et al ²⁷ (2011)	First-line	774	Bevacizumab 2.5 mg/kg/wk + capecitabine + DDP Placebo + capecitabine + DDP	386 381	58 59	6.7 5.3	12.1 10.1	5
Bang et al ²⁸ (2010)	First-line	594	Trastuzumab + chemotherapy Chemotherapy	294 290	59.4 58.5	6.7 5.5	13.8 11.1	3

Abbreviations: PTX, paclitaxel; DDP, cisplatin; EOC, epirubicin plus oxaliplatin plus capecitabine; PFS, progression-free survival; OS, overall survival; NR, not reported.

Incidence of severe and fatal AEs

A total of 2,647 patients from nine treatment arms receiving MTAs were available for severe AEs incidence analysis. Using a random-effects model, the summary incidence of severe AEs was 72.5% (95% CI: 66.4%–77.8%). As for FAEs, a total of 2,647 patients from nine treatment arms were included, and the pooled incidence was 2.2% (95% CI: 1.6%–2.9%) using a fixed-effects model ($I^2=48%$, $P=0.051$).

RR of severe and fatal AEs

A meta-analysis of RR for severe and fatal AEs attributable to MTAs compared with control was performed. The pooled results showed that the use of MTAs significantly increased the risk of severe AEs (RR: 1.12, 95% CI: 1.02–1.24, $P=0.02$; Figure 2), but not FAEs when compared with controls (RR: 0.97, 95% CI: 0.65–1.45, $P=0.88$; Figure 3) using a fixed-effects model.

Specific FAEs

Individual specified and nonspecified causes of FAEs are listed in Table 2. There were 49 FAEs on the treatment arms and 46 FAEs on the controlled arms; 42.9% and 50% of them were nonspecified etiology, respectively. For those specified FAEs in this study, the most common causes of FAEs with MTAs

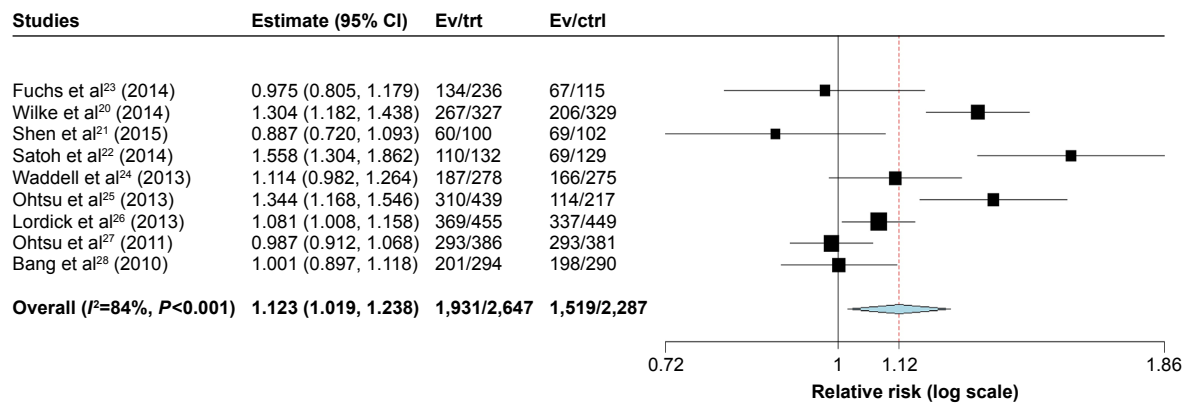


Figure 2 RR of severe AEs (95% CI) associated with therapies with or without MTAs. **Abbreviations:** RR, relative risk; AEs, adverse events; MTAs, molecular targeted agents; CI, confidence interval; Ev, events; trt, treatments; ctrl, control.

were infections (16.3%), gastrointestinal hemorrhage (8.2%), and arterial thromboembolic events (8.2%), respectively. Additionally, the specified FAEs of eight infections were pneumonitis (three), *Candida* sepsis (one), *Klebsiella* sepsis (one), sepsis (two), and neutropenic sepsis (one), respectively.

Publication bias

We used Begg’s funnel plot and Egger’s test to assess the publication bias. The Begg’s funnel plots did not show any evidence of publication bias ($P=0.89$ for severe AEs and $P=0.54$ for FAEs, respectively). Additionally, Egger’s test also did not suggest any evidence of publication bias ($P=0.56$ for severe AEs and $P=0.30$ for FAEs, respectively).

Discussion

During the past decades, the introduction of biological agents targeting specific growth and survival pathways, such as EGFR, PI3K/Akt/mTOR pathway, and angiogenesis through the VEGF signaling cascade, seems to be the most promising

strategy to improve outcome of advanced GC patients. Trastuzumab in combination with chemotherapy has been approved by US Food and Drug Administration (FDA) as first-line treatment for patients with HER2-positive advanced gastric or gastroesophageal junction cancer due to its survival benefit when compared to chemotherapy alone.²⁸ More recently, ramucirumab, a monoclonal antibody VEGFR-2 antagonist, in combination with paclitaxel also significantly increased overall survival in previously treated patients with advanced GC compared with paclitaxel, which led to its approval for use in second-line treatment for advanced GC.²⁹ However, no clear survival benefit was experienced with agents targeting EGFR (cetuximab and panitumumab), VEGF-A (bevacizumab), or mTOR (everolimus). Because of the wide use of MTAs in GC patients, concerns have arisen regarding the risk of severe and fatal AEs with these drugs. Indeed, several previous meta-analyses have been performed to assess the severe and fatal toxicities associated with these MTAs. For example, a previous meta-analysis conducted by Ranpura et al³⁰ showed that the addition of bevacizumab

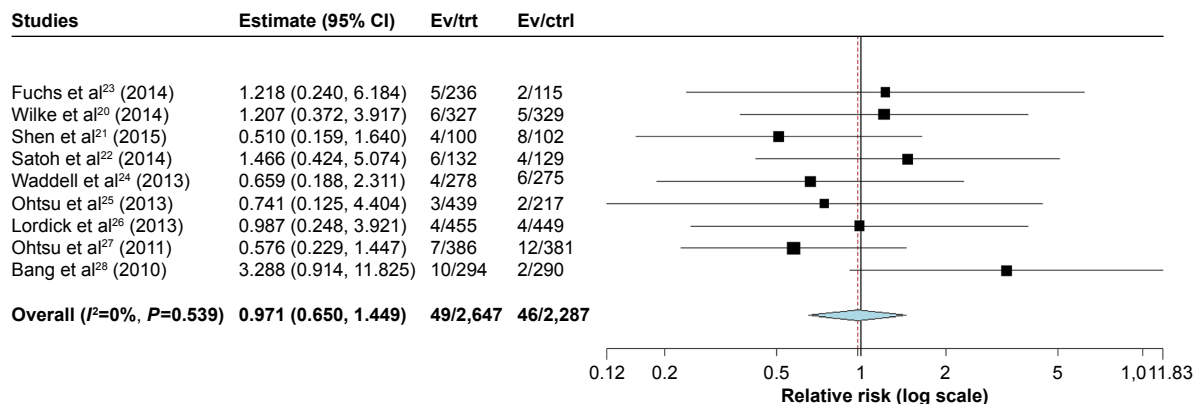


Figure 3 RR of FAEs (95% CI) associated with therapies with or without MTAs. **Abbreviations:** RR, relative risk; FAEs, fatal adverse events; MTAs, molecular targeted agents; CI, confidence interval; Ev, events; trt, treatments; ctrl, control.

Table 2 FAEs by specific type

	Events on MTAs groups	Events on control arms
Unspecified	21	23
Specified		
Infections	8	7
Hemorrhage	4	2
Arterial thromboembolic events	4	2
Pulmonary embolism	2	2
GI perforation	2	1
Sudden death	3	2
Diarrhea	1	1
Malabsorption	2	–
Renal failure	–	2
Cardiac failure	1	1
Cardiac arrest	1	–
Subileus	–	1
Multiorgan failure	–	1
Dehydration	–	1
Overall	49	46

Abbreviations: FAEs, fatal adverse events; GI, gastrointestinal; MTAs, molecular targeted agents.

to chemotherapy significantly increased treatment-related mortality (RR: 1.33, 95% CI: 1.02–1.73, $P=0.04$), and two later meta-analyses also demonstrated that the use of VEGF receptor tyrosine kinase inhibitors was associated with increased risk of FAEs.^{29–32} Increased risk of severe and fatal AEs associated with anti-EGFR agent cetuximab has also been observed in colorectal cancer patients.³³ Additionally, two recent meta-analyses have demonstrated that the use of mTOR inhibitors significantly increase the risk of FAEs.^{34,35} To the best of our knowledge, there is limited data specifically focusing on the severe and fatal AEs related to MTAs in advanced GC patients. Therefore, we conducted this meta-analysis of Phase III RCTs with available toxicity data of MTAs in advanced GC patients.

Our study includes a total of nine Phase III RCTs involving 4,934 GC patients. The summary incidence of severe and fatal AEs with MTAs was 72.5% and 2.2%, respectively. We also find that the addition of MTAs to therapies in advanced GC significantly increased the risk of developing of severe AEs, but not for FAEs. Additionally, the most common causes of FAEs with MTAs were infections (16.3%), gastrointestinal hemorrhage (8.2%), and arterial thromboembolic events (8.2%), respectively. On the basis of our findings, we agree with the continued use of MTAs in GC patients owing to its survival benefits, but suggest close monitoring for treatment-related complications. In this study, we also find that infections are the most common FAEs associated with MTAs, which is consistent with previous studies.^{31–34} In a recent meta-analysis conducted by Qi et al,³⁶ the authors

found that the risk of developing an infection of any grade was 1.45-fold higher in patients treated with bevacizumab. More importantly, there was a 1.59-fold increase in the risk of high-grade infection associated with the use of bevacizumab. Another meta-analysis also demonstrated that the use of anti-EGFR MoAbs significantly increased the risk of developing severe infections (RR: 1.34, 95% CI: 1.10–1.62, $P=0.003$) in cancer patients. More importantly, the use of anti-EGFR MoAbs significantly increased the risk of severe sepsis in cancer patients (RR: 4.30, 95% CI: 1.80–10.27, $P=0.001$).³⁷ On the basis of these findings, clinicians should pay more attention to severe infections to reduce the risk of FAEs in advanced gastric patients. Before the initiation of MTAs in gastric patients, clinicians should fully treat patients with any active infection and must monitor patients during the course of MTAs treatment. However, patients with active or recently active infections are excluded from clinical trials; therefore, the true incidence of these infections could be widely under-reported. More trials focusing on this issue are still needed.

Several limitations exist in this analysis. First, although AEs are prospectively collected for each individual study, this analysis remains a retrospective research that is subject to the method deficiencies of the included trials. We minimized the likelihood of bias by strictly selecting Phase III RCTs with direct comparison with and without MTAs before the analysis. Second, we included patients treated with different targeted agents, which would increase the clinical heterogeneity among included trials, which also makes the interpretation of a meta-analysis more problematic, although we pooled subgroup analysis according to treatment line. Finally, in the meta-analysis of published studies, publication bias is important because trials with positive results are more likely to be published and trials with null results tend not to be published. Our research detects no publication bias using Begg and Egger tests for severe and fatal AEs.

Conclusion

In conclusion, this is the first meta-analysis that specifically assessed the severe and fatal toxicities of adding MTAs to therapies in the treatment of GC patients. The results of our study suggest that the addition of MTAs to therapies in GC significantly increases the risk of developing severe AEs, but not for FAEs. Additionally, the most common causes of FAEs with MTAs were infections, gastrointestinal hemorrhage, and arterial thromboembolic events, respectively.

Disclosure

The authors report no conflicts of interest in this work.

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