



2016 Gastric Cancer: Global view

Changing strategies for target therapy in gastric cancer

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Abstract

In spite of a worldwide decrease in the incidence of gastric cancer, this malignancy still remains one of the

leading causes of cancer mortality. Great efforts have been made to improve treatment outcomes in patients with metastatic gastric cancer, and the introduction of trastuzumab has greatly improved the overall survival. The trastuzumab treatment took its first step in opening the era of molecular targeted therapy, however several issues still need to be resolved to increase the efficacy of targeted therapy. Firstly, many patients with metastatic gastric cancer who receive trastuzumab in combination with chemotherapeutic agents develop resistance to the targeted therapy. Secondly, many clinical trials testing novel molecular targeted agents with demonstrated efficacy in other malignancies have failed to show benefit in patients with metastatic gastric cancer, suggesting the importance of the selection of appropriate indications according to molecular characteristics in application of targeted agents. Herein, we review the molecular targeted agents currently approved and in use, and clinical trials in patients with metastatic gastric cancer, and demonstrate the limitations and future direction in treatment of advanced gastric cancer.

Key words: Advanced gastric cancer; Target therapy; Chemotherapy; Strategy; Signal pathway

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Core tip: This review summarizes the development of molecular targeted therapeutic agents in advanced gastric cancer. Agents targeting angiogenesis as well as ERBB receptors and their downstream signaling pathways are introduced. Current efforts to overcome resistance to the human epidermal growth factor receptor 2-targeted agents are also presented from the ongoing clinical trials. Future direction of target therapy should be guided according to further clarification of the molecular mechanisms of gastric cancer and by exploring appropriate indications for application of molecular targeted therapy to improve its efficacy.

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INTRODUCTION

Although the incidence of gastric cancer has been declining worldwide, it is the fifth most frequently diagnosed cancer and the third leading cause of cancer mortality^[1]. Gastric cancer is frequently diagnosed at an advanced, incurable stage due to its asymptomatic feature at its early stages. Systemic chemotherapy is usually offered as treatment for patients with advanced incurable gastric cancer, but treatment outcomes are dismal, with a range of median survival of 6-11 mo^[2]. Recent advances in molecular targeted therapies have led to an improved prognosis in patients with advanced, unresectable gastric cancer. A monoclonal antibody interfering with the activation of human epidermal growth factor receptor 2 (HER2) was the first targeted agent to demonstrate significant survival benefit in the treatment of gastric cancer. Despite the proven survival advantage of the HER2-directed monoclonal antibody in patients with HER2-overexpressing advanced gastric cancer (AGC), several problems still remain to be solved^[3]. One of them is the emergence of gastric tumor cells resistant to treatment with the HER2 monoclonal antibody. In order to overcome resistance, a variety of investigational molecular targeted agents have been developed and some have shown encouraging results in clinical trials^[4,5].

On the other hand, several targeted therapies have been studied in patients with AGC, but few agents have been proven to be beneficial. This is, in part, thought to be attributed to the biological heterogeneity of gastric cancer, and, therefore, careful selection of patients may be a key factor in the successful target therapies in patients with AGC.

This article reviews the molecular targeted agents in clinical use, their limitations and potential strategies to overcome them, and introduces ongoing clinical trials as well as the future direction of target therapy in unresectable AGC.

AGENTS TARGETING ERBB FAMILY RECEPTORS

The ERBB family of receptors, receptor tyrosine kinases (RTKs), consists of four members, epidermal growth factor receptor (EGFR) and the EGFR-related receptors - HER2, HER3, and HER4. This family of receptors is transmembrane receptors consisting of an extracellular domain, a single hydrophobic transmembrane segment and an intracellular domain containing

a preserved tyrosine kinase residue (Figure 1).

EGFR is ubiquitously expressed in epithelial, mesenchymal and neuronal cells, and plays a role in development, proliferation and differentiation^[6]. The signaling through the EGFR is initiated with binding of the ligands to domain I and III of the extracellular domain, which subsequently induces formation of a heterodimer or homodimer between the receptor family members leading to autophosphorylation of the tyrosine kinase residues in the carboxy-terminus of the receptor protein. The autophosphorylated receptor subsequently activates a downstream signaling cascade through the RAS-RAF-mitogen activated protein kinase kinase (MEK)-mitogen-activated protein kinase (MAPKs) pathway. In addition to the RAS-RAF-MAPKs, several other pathways, such as the phosphatidylinositol 3-kinase (PI3Ks)-AKT or RAS-PLC γ -PKC are known to be activated by ERBB receptor signaling^[7-10] (Figure 2).

The activation process of ERBB signaling pathway ranges from the tumorigenesis such as cell division and migration to differentiation and apoptosis, depending on cellular context^[11]. ERBB receptors are associated with development and alteration of various types of cancer with several mechanisms. The best known example of the alteration is amplification of ERBB2 in a subset of breast cancers as well as in gastric, ovarian, and salivary cancers^[12-14]. In non-small cell lung cancers (NSCLC), mutations in the tyrosine-kinase domain of EGFR have been found in a subset of patients^[15-17]. With regard to tumorigenesis, ERBB receptors have been the candidates as targets for anti-cancer therapy. The ERBB receptors-targeted agents are summarized in Table 1.

Anti-EGFR targeting agents

Cetuximab: Cetuximab is a mouse/human chimeric monoclonal antibody that targets the EGFR. Treatment with cetuximab monotherapy in patients with AGC who had received prior chemotherapy showed minimal clinical activity in a phase 2 clinical trial^[18]. Another study in patients with AGC treated with cetuximab in combination with irinotecan as a second-line chemotherapy, revealed that combination therapy was effective in a subset of patients [median overall survival (OS) 5.5 mo, 95%CI: 3.6-7.3]^[19]. The controversy was terminated by a randomized, open-label phase 3 trial (EXPAND), which showed no benefit with the addition of cetuximab to combination chemotherapy. Patients diagnosed with advanced gastric or gastroesophageal junction cancer were randomized to receive capecitabine and cisplatin combination chemotherapy with or without cetuximab as a first-line chemotherapy. No significant difference in progression-free survival (PFS), the primary endpoint of the study, was shown in this study [4.4 mo vs 5.6 mo, hazard ratio (HR) = 1.09, 95%CI: 0.92-1.29, $P = 0.32$]^[20].

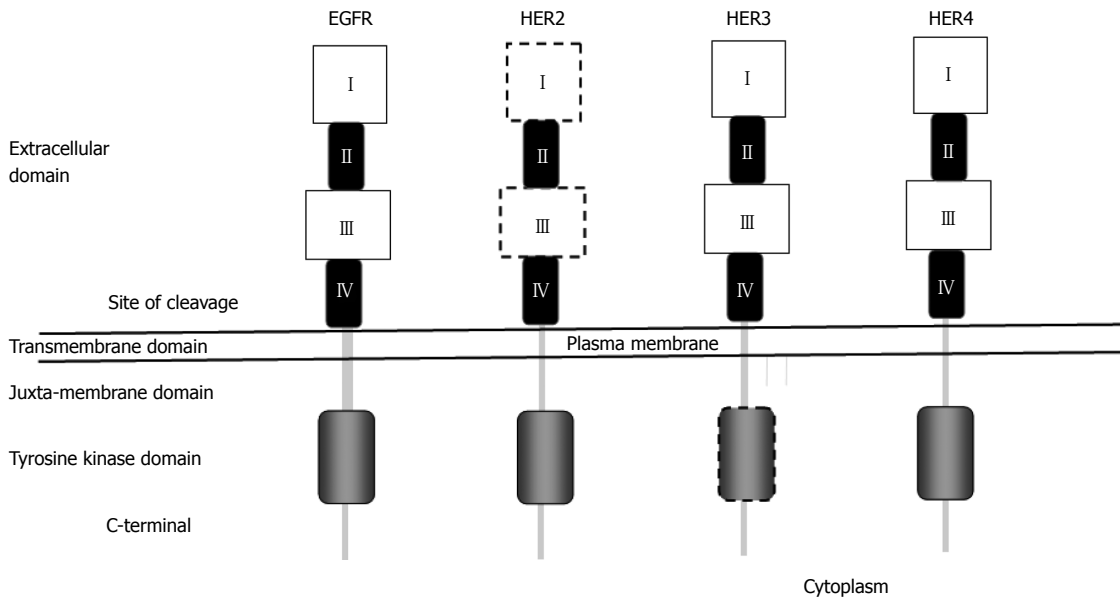


Figure 1 Structure of ERBB family of receptors. EGFR: Epidermal growth factor receptor; HER: Human EGFR.

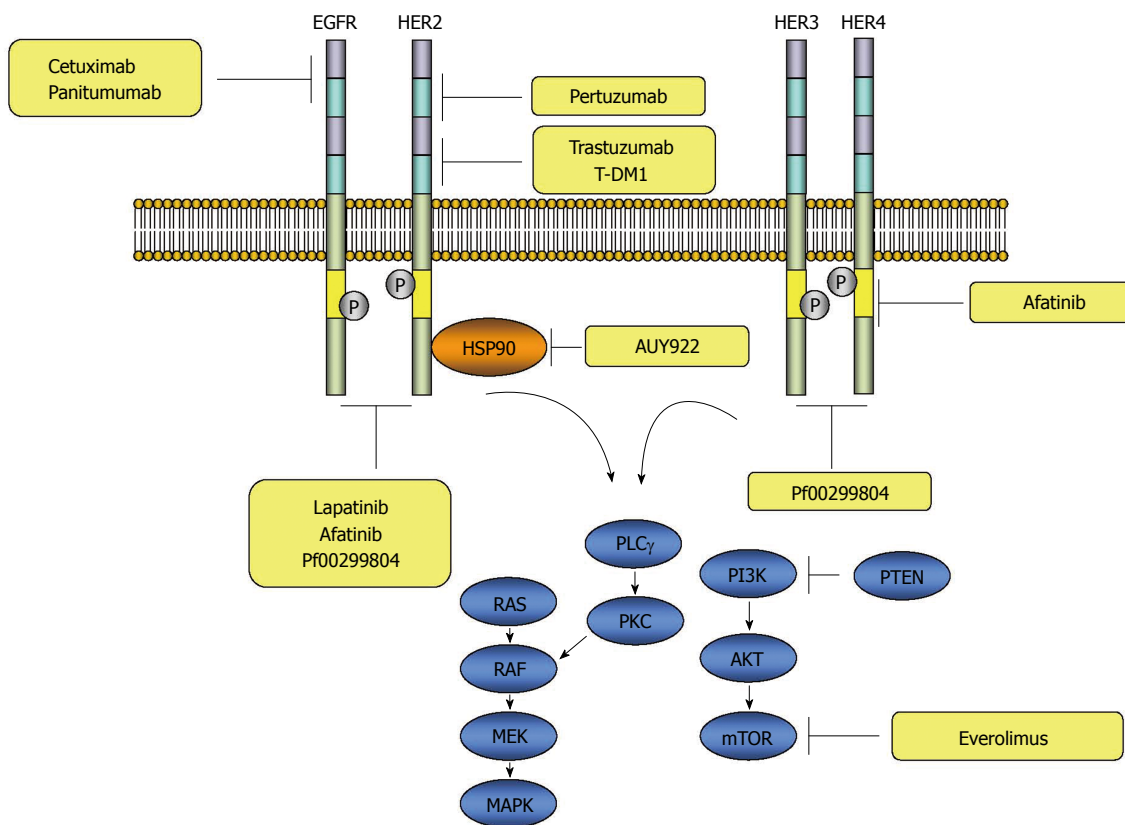


Figure 2 Signal transduction cascade through activation of ERBB receptors and schematic diagram of molecular targeting agents where they work. EGFR: Epidermal growth factor receptor; HER: Human EGFR; PLC- γ : Phospholipase C- γ ; PKC: Protein kinase C; MEK: Protein kinase kinase; MAPK: Mitogen-activated protein kinase; mTOR: Mammalian target of rapamycin; PI3K: Phosphatidylinositol 3-kinase.

Panitumumab: Panitumumab is a fully human immunoglobulin (Ig) G2 monoclonal antibody directed against the EGFR. After determination of the optimal dosage of panitumumab as 9 mg/kg when used in combination with epirubicin, oxaliplatin, and capecitabine in a dose-finding study^[21], a randomized,

open-label phase 3 trial was performed. Patients with previously untreated advanced esophagogastric adenocarcinoma were randomized to receive either epirubicin, oxaliplatin and capecitabine (EOC) or modified EOC plus panitumumab. The primary endpoint was OS, however, the addition of panitumumab did

Table 1 Clinical trials with agents targeting ERBB family receptors

Study	n	Design	Line	Treatment	Primary end point	Results	P value
EXPAND ^[19]	904	Phase 3, RCT	First	Cetuximab + XP vs Placebo + XP	PFS	HR = 1.09; 95%CI: 0.92-1.29	0.320
REAL3 ^[22]	553	Phase 3, RCT	First	Anitumumab + mEOC vs EOC	OS	HR = 1.37; 95%CI: 1.07-1.76	0.013
TOGA ^[3]	594	Phase 3, RCT	First	Trastuzumab + XP vs XP	OS	HR = 0.74; 95%CI: 0.60-0.91	0.005
HERBIS-1 ^[29]	56	Phase 2, non-RCT	First	Trastuzumab + S-1 + Cisplatin	RR	RR = 68%; 95%CI: 0.54-0.80	
TyTAN ^[38]	261	Phase 3, RCT	Salvage	Lapatinib + Paclitaxel vs Paclitaxel	OS	HR = 0.84; 95%CI: 0.64-1.11	0.350
LOGiC ^[39]	545	Phase 3, RCT	First	Lapatinib + CapeOx vs Placebo + CapeOx	OS	HR = 0.91; 95%CI: 0.73-1.12	0.104

RCT: Randomized controlled trial; XP: Capecitabine plus Cisplatin; PFS: Progression free survival; RAM: Ramucirumab; OS: Overall survival.

not increase OS with significantly better survival in the chemotherapy only group [11.3 mo vs 8.8 mo (panitumumab plus mEOC group), HR = 1.37, 95%CI: 1.07-1.76, $P = 0.013$]^[22].

HER2 targeted agents

HER2 is a transmembrane RTK, which belongs to the ERBB family of receptors. Like other HER family receptors, the HER2-neu receptor consists of an ectodomain, transmembrane domain, and endodomain. The ectodomain of the receptor has four domains, including two insulin-like growth factor-like ligand binding domains (I-III) and two cysteine-rich domains (II-IV) (Figure 1). Unlike the other family members of the ERBB, no ligands for HER2 have been identified. The ligand-independent transactivation of HER2 receptors through homo- or hetero-dimerization with other ERBB family members leads to activation of a downstream signaling cascade through the RAS-RAF-MEK-MAPKs or PI3Ks-AKT-mammalian target of rapamycin (mTOR) pathway^[23,24] (Figure 2).

Trastuzumab: Trastuzumab is a humanized monoclonal antibody directed against the HER2 receptor and exerts activity by binding to the domain IV of the extracellular domain^[25]. Several mechanisms by which trastuzumab inhibits activation of HER2 receptors include antibody-dependent cellular cytotoxicity (ADCC)^[26], inhibition of intracellular signal transduction, blocking proteolytic cleavage of the extracellular domain, reduction of tumor angiogenesis, and inhibition of recovery from DNA damage^[27]. Because HER2 was reported to be amplified in 13%-23% of all gastric cancers^[28], the agent targeting the HER2 receptor was introduced for the treatment of gastric cancer. Trastuzumab was the first molecule-targeted agent approved for the treatment of gastric cancer after the randomized, prospective, multicenter, phase 3 (ToGA) study. The significant survival benefit in patients overexpressing HER2 was demonstrated in ToGA study, in which patients with AGC were randomized to receive cytotoxic chemotherapy comprising fluoropyrimidine and cisplatin with or without trastuzumab (13.8 mo vs 11.1 mo, HR = 0.74, 95%CI: 0.60-0.91, $P = 0.0046$)^[3].

Another study with trastuzumab in combination

with S-1 and cisplatin reported favorable efficacy in a multicenter phase 2 trial. The HERBIS-1 study was designed for patients with HER2-positive AGC to receive S-1 and cisplatin in addition to trastuzumab, and reported a 68% response rate, 16 mo of OS, and 7.8 mo of PFS^[29].

Pertuzumab: Pertuzumab is a monoclonal antibody that interferes with dimerization by binding the domain II of the HER2 ectodomain^[30]. Based on a pre-clinical study, in which the anti-tumor activity of combination immunotherapy with pertuzumab and trastuzumab was proved to be superior to a monotherapy with either antibody in a HER2-positive human gastric cancer xenograft model^[31], and the CLEOPATRA study that demonstrated the superior OS as well as PFS in HER2-positive metastatic breast cancer patients treated with the combined pertuzumab and trastuzumab in addition to docetaxel compared with patients treated with placebo, trastuzumab and docetaxel^[32,33], a phase 2a trial was designed with combination of pertuzumab, trastuzumab and chemotherapy. The dose of pertuzumab used in a phase 3 study was determined in the phase 2a trial, and a pertuzumab dose of 840 mg every three weeks for six cycles in addition to trastuzumab, capecitabine and cisplatin, showed a 55% partial response rate in patients with HER2-positive AGC without prior chemotherapy^[34].

Lapatinib: Lapatinib is a small molecule tyrosine kinase inhibitor that simultaneously inhibits phosphorylation of both EGFR and HER and prevents activation of the downstream signaling cascade. A pre-clinical study demonstrated effectiveness of lapatinib against p96HER-2 expressing cells which were resistant to trastuzumab because p95HER2 is an amino terminally truncated receptor with preserved kinase activity that results in interruption of trastuzumab to bind the HER2 receptor^[35]. Lapatinib was proven to have clinical benefit in treatment of patients with HER2-positive metastatic breast cancer in terms of OS (HR = 0.76, 95%CI: 0.60-0.96) and PFS (HR = 0.61, 95%CI: 0.50-0.74) in a meta-analysis^[36].

In contrary to the proven benefit of lapatinib in HER2-positive metastatic breast cancer, the outcome

of patients with gastric cancer is poor in clinical trials. In a phase 2 study of lapatinib used as first-line treatment, the response rate was only 9% and median OS was 4.8 mo (95%CI: 3.2-7.4)^[37]. Two phase 3 trials on lapatinib also showed unsatisfactory results. In the TyTAN study, a combination of lapatinib and weekly paclitaxel was compared to weekly paclitaxel monotherapy as the second-line treatment in HER2-positive gastric cancer. No significant advantage in terms of OS (11 mo vs 8.9 mo, $P = 0.1044$) and PFS (5.4 mo vs 4.4 mo, $P = 0.2441$) was shown^[38]. The efficacy of lapatinib was also studied as a first-line treatment in the LOGiC phase 3 trial. Combination chemotherapy of capecitabine and oxaliplatin with lapatinib was compared to that without lapatinib in HER2-positive gastric cancer, and no significant benefit in survival was demonstrated (HR = 0.91, 95%CI: 0.73-1.12, $P = 0.35$) or PFS (HR = 0.86, 95%CI: 0.71-1.04, $P = 0.10$)^[39].

Strategies to overcome resistance to anti-HER2 in gastric cancer

Despite the proven efficacy of trastuzumab in the treatment of HER2-overexpressing gastric cancer, 12% of patients treated with chemotherapy plus trastuzumab were refractory to the therapy, and disease progression eventually documented in 7 mo from the initiation of the therapy^[3], suggested presence of primary resistance and development of acquired resistance against the antibody. A variety of mechanisms of acquired resistance to trastuzumab in gastric cancer has been proposed. These include: (1) dimerization or crosstalk of HER2 with other molecules such as HER3 and MET leading to subsequent activation of downstream signaling pathways such as PI3K pathway^[40]; (2) genetic alteration and subsequent aberrant activation of HER2 downstream signaling pathways^[40]; (3) epithelial-mesenchymal transition signaling^[41]; and (4) intra-tumoral heterogeneity of gastric cancer^[42,43]. To overcome these resistance-mediating mechanisms, a paradigm shift of concept for gastric cancer treatment is needed. Good candidate drugs used for cancers originating from other organs are not always good for gastric cancer due to the concept of cancer addiction difference, which means that different cancer cells use different mechanisms for carcinogenesis.

c-Met inhibitor

c-Met is a RTK that stimulates cell proliferation, survival and invasion/metastasis. Binding of hepatocyte growth factor (HGF) to its receptor, MET, initiates activation of downstream signal transduction pathways including MAPK cascades and the PI3K-Akt axis^[44]. It has been known that the Met/EGFR family receptors' crosstalk plays a role in the development of drug resistance, such as resistance to gefitinib and erlotinib in NSCLC^[45]. Furthermore, Met transcript and protein levels have

also been reported to be elevated in breast cancer cells overexpressing HER2 in response to treatment with HER2 inhibitor, suggesting that Met signaling compensates for HER2 inhibition^[46]. In gastric cancer, MET gene amplification and MET protein overexpression have been reported with a frequency of 10%-20% and 50%, respectively^[47,48]. Based on these findings, an open-label dose de-escalation phase 1b and double-blind randomized phase 2 trial were performed using rilotumumab, a fully humanized monoclonal IgG2 antibody against HGF. Patients received epirubicin, capecitabine and cisplatin with or without rilotumumab. Significantly improved PFS was reported in the rilotumumab group compared to the placebo group with metastatic AGC who had not received previous systemic therapy (5.7 mo vs 4.2 mo, HR = 0.60, 80%CI: 0.45-0.79, $P = 0.016$)^[4].

mTOR inhibitor

Aberrant activation of the HER2 signaling pathways, including PI3K/Akt/mTOR pathway, is known to be one of the mechanisms of trastuzumab resistance. As loss of function mutations in *PTEN* or activating mutations in *PIK3CA* is known to cause constitutive activation of the PI3K, use of PI3K inhibitors or mTOR inhibitors such as everolimus could overcome trastuzumab resistance in gastric cancer. The efficacy of everolimus was studied in a phase 2 trial, and the results showed that disease control rate, the median OS and the median PFS were 26%, 10.1 mo (96%CI: 6.5-12.1), 2.7 mo (95%CI: 1.6-3.0), respectively, in previously treated metastatic gastric cancer patients^[49]. Based on these results, a multicenter, double-blind, randomized, phase 3 trial was performed in previously treated AGC patients. Patients were assigned to receive either everolimus or placebo. The primary endpoint was OS. Although significant improvement in PFS (1.7 mo vs 1.4 mo, HR = 0.66, 95%CI: 0.56-0.78, $P < 0.0001$) was observed, this clinical trial failed to meet the primary objective, as there was no significant difference in OS (5.4 mo vs 4.3 mo, HR = 0.9, 95%CI: 0.75-1.08, $P = 0.124$)^[50].

Pan-HER inhibitor

Although afatinib, a tyrosine kinase inhibitor directed to multi-ERBB family receptors, inhibits multiple tyrosine kinase receptors of ERBB family, activation of HER3 is not blocked. HER3 is regarded as an important, intimate signaling partner in HER2-mediated tumorigenesis through the PI3K/Akt pathway and is one of the molecules responsible for resistance to HER2 targeted therapies^[40]. Indeed, it was reported that overexpression of HER3 was observed in trastuzumab-resistant HER2-positive breast carcinoma cell lines after long-term trastuzumab exposure^[51]. It was also reported that PI3K/Akt dependent up-regulation of HER3 mRNA and protein was observed after inhibition of the HER2 tyrosine kinase with lapatinib,

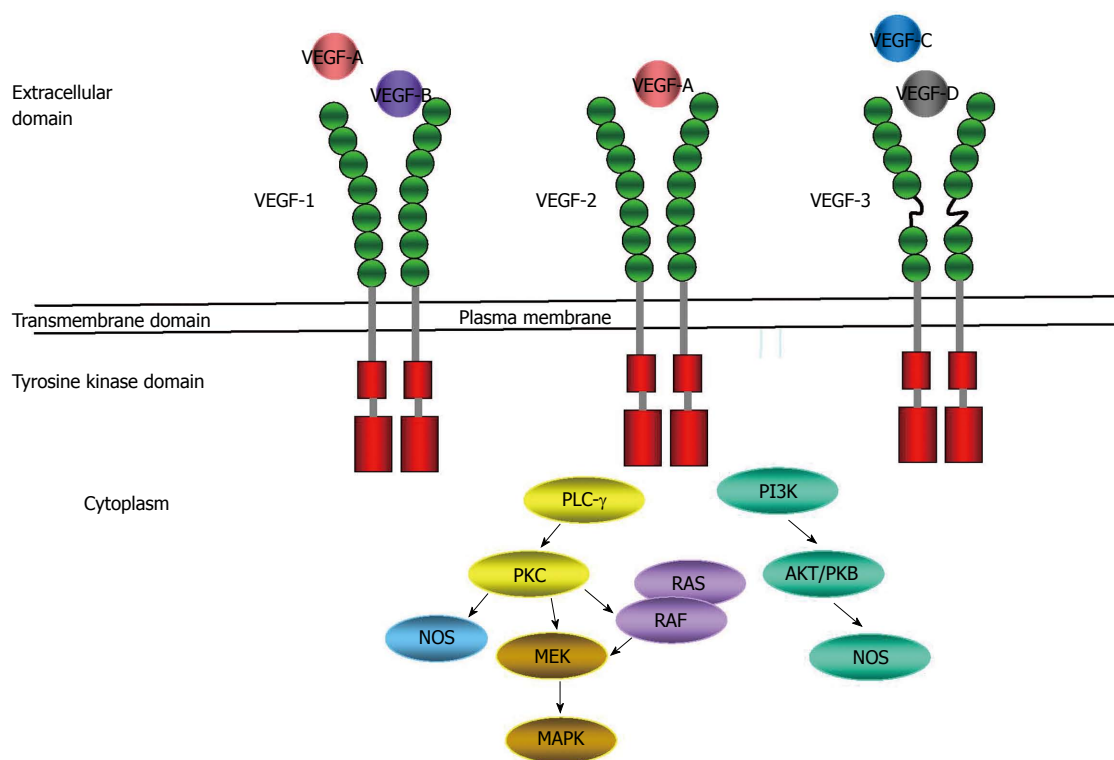


Figure 3 Vascular endothelial growth factor receptor and signal transduction. Binding of ligands to receptors leads to phosphorylation of the receptors by activation of receptor-kinase activity, which leads to subsequent signal transduction. VEGF: Vascular endothelial growth factor; PLC- γ : Phospholipase C- γ ; PKC: Protein kinase C; MEK: Protein kinase kinase; MAPK: Mitogen-activated protein kinase; PI3K: Phosphatidylinositol 3-kinase.

suggesting incomplete block of the PI3K pathway by HER2 inhibitors because of HER3-mediated compensation^[52]. These findings indicate that the combined targeting of HER2 and HER3 may be more effective in blocking HER2 downstream signaling activation. Indeed, a preclinical study with gastric cancer cell lines demonstrated the synergistic effects of a combination of the pan-HER inhibitor (PF00299804) and trastuzumab or chemotherapeutic agents^[53].

ANTI-ANGIOGENESIS

Angiogenesis is a multistep process of new vasculature formation from the pre-existing blood vessel. The vascular endothelial growth factor (VEGF)-mediated signaling is known to play an essential role in the angiogenesis and vascular permeability. In addition to these roles, it also contributes to the tumorigenesis, tumor migration and metastasis^[54,55]. VEGF consists of a large family of growth factors that include VEGFA, VEGFB, VEGFC, VEGFD, and placental growth factor. The classical VEGF receptors (VEGFRs) that mediate signaling are RTKs VEGFR1, VEGFR2, and VEGFR3 expressed by vascular and lymphatic endothelial cells^[54]. VEGFR2 is the predominant RTK that mediates VEGF signaling to induce angiogenesis. Despite higher affinity of VEGFR1 for binding to VEGF, the tyrosine phosphorylation of the receptor is weaker than VEGFR2. The downstream signaling by activation of VEGFR2 is mediated by several pathways, including

the phospholipase C- γ , protein kinase C, extracellular signal-related kinase, PI3Ks, and endothelial nitric oxide synthase pathways^[54,56,57] (Figure 3).

In patients with gastric cancer, high expression of VEGF is known to be associated with poor prognosis^[58]. Several clinical trials to evaluate the efficacy of anti-angiogenic agents have been carried out in patients with gastric cancer (Table 2).

Bevacizumab

Bevacizumab is a humanized monoclonal IgG1 directing against VEGF-A. A large randomized phase 3 trial, Avastin in Gastric Cancer Trial, evaluated the clinical benefit of the addition of bevacizumab to combination chemotherapy. Patients with previously untreated AGC were randomized to receive bevacizumab or placebo in combination with capecitabine and cisplatin. The median overall survival (OS) between two groups was not significantly different (12.1 mo vs 10.1 mo, HR = 0.87, 95%CI: 0.73-1.03, $P = 0.1002$) and this study failed to reach its primary objective^[59].

Ramucirumab

Ramucirumab is a fully humanized IgG1 monoclonal antibody directed to the extracellular VEGF-binding domain of VEGFR-2^[60]. An international, randomized, double-blinded, placebo-controlled, phase 3 trial was conducted in patients with AGC who had been previously treated with platinum or fluoropyrimidine-containing chemotherapy. Patients were randomly

Table 2 Clinical trials with angiogenesis targeting agents

Study	<i>n</i>	Design	Line	Treatment	Primary end point	Results	<i>P</i> value
AVAGAST ^[59]	774	Phase 3, RCT	First	Bevacizumab + XP vs Placebo + XP	OS	HR = 0.87; 95%CI: 0.73-1.03	0.1002
REGARD ^[61]	355	Phase 3, RCT	Second	RAM vs Placebo	OS	HR = 0.81; 95%CI: 0.68-0.96	0.0470
RAINBOW ^[62]	665	Phase 3, RCT	Second	RAM + paclitaxel vs Placebo + paclitaxel	OS	HR = 0.81; 95%CI: 0.68-0.96	0.0170
Qin <i>et al</i> ^[63]	270	Phase 3, RCT	Third	Apatinib vs placebo	OS	HR = 0.71; 95%CI: 0.54-0.94	0.0160

RCT: Randomized controlled trial; XP: Capecitabine plus Cisplatin; RAM: Ramucirumab; OS: Overall survival.

assigned to receive ramucirumab monotherapy or placebo. The significant benefit in terms of prolonged survival was demonstrated in this study (5.2 mo vs 3.8 mo, HR = 0.776, 95%CI: 0.603-0.998, *P* = 0.047), meeting its primary endpoint^[61]. Another more recent international phase 3 trial also evaluated the clinical advantage of ramucirumab in combination with chemotherapy. Patients with metastatic AGC which progressed despite first-line chemotherapy were randomized to receive ramucirumab plus paclitaxel or placebo plus paclitaxel. Significantly longer OS was observed in the ramucirumab plus paclitaxel group (9.6 mo vs 7.4 mo, HR = 0.807, 95%CI: 0.678-0.962, *P* = 0.017), satisfying the primary endpoint^[62]. The efficacy of ramucirumab as a first-line treatment in patients with AGC was also examined in a randomized, double-blinded, multicenter, phase 2 trial. Patients with previously untreated metastatic AGC were randomized to receive mFOLFOX6 plus ramucirumab or mFOLFOX6 plus placebo, and the primary endpoint was PFS. However, no significant improvement of PFS was observed by adding ramucirumab to mFOLFOX6 (6.4 mo vs 6.7 mo, HR = 0.98, 95%CI: 0.69-1.37, *P* = 0.89), and the study failed to meet its primary endpoint (NCT 01246960, Clinicaltrial.gov).

Apatinib

Apatinib is an oral small molecular inhibitor of VEGFR-2 tyrosine kinase. A multicenter, randomized, double-blind, placebo-controlled phase 3 trial to evaluate the survival benefit of apatinib in AGC patients with prior failure on second-line chemotherapy has been completed. Patients were randomly assigned to receive apatinib or placebo. Significantly improved OS was observed in patients treated with apatinib (6.5 mo vs 4.7 mo, HR = 0.71, 95%CI: 0.54-0.94, *P* < 0.016), meeting the primary objective^[63] (NCT01512745, ClinicalTrials.gov).

ONGOING CLINICAL TRIALS

Trastuzumab-emtansine

Trastuzumab-emtansine (T-DM1, Genetech/Roche, South San Francisco, CA, United States) is an antibody-drug conjugate comprising trastuzumab and DM1, a microtubule inhibitor (maytansine). After binding of T-DM1 to the HER2 receptor in HER2 expressing cells, internalization occurs, and the cytotoxic DM1 moiety

is released inside cells. T-DM1 also retains all the mechanisms of action of trastuzumab such as ADCC and inhibition of the downstream signaling pathway^[64]. The clinical benefit of T-DM1 was demonstrated in patients with HER2-positive metastatic breast cancer in phase 1 to 3 studies^[65-67]. T-DM1 was also demonstrated to be more effective than trastuzumab in xenograft gastric cancer models^[68]. The phase 2/3 clinical trial of T-DM1 is ongoing currently in patients with HER2-positive AGC who failed in the first-line therapy (MCT01641939; ClinicalTrials.gov).

Afatinib

Afatinib (GilotrifTM, Boehringer Ingelheim) is an irreversible inhibitor of the tyrosine kinases of ERBB1-2 and ERBB4 receptors. It is also reported to inhibit transphosphorylation of HER3^[69]. This oral treatment agent has antitumor activity against acquired mutations resistant to first-generation inhibitors in NSCLC^[70]. Clinical trials to examine the efficacy of this agent in NSCLC, breast cancer, and head and neck cancer are now underway^[69]. In gastric cancer, a phase 2 trial is ongoing in patients with metastatic HER2-positive gastric cancer resistant to trastuzumab (NCT01522768; ClinicalTrials.gov).

Heat shock protein 90 inhibitor

Heat shock protein 90 (HSP90) is a ubiquitously expressed chaperone involved in post-translational structural folding and protein stability. The structure of HSP90 consists of an NH₂-terminal region, middle region, and a COOH-terminal region, and inhibition at the NH₂-terminal ATP-binding site results in degradation of the client proteins through the ubiquitin proteasome pathway^[71]. NVP-AUY922 is part of the isoxazole HSP90-inhibitor family and inhibits ATPase activity. Using the gastric and breast cancer cell lines or xenograft models, AUY922 was demonstrated to have anti-proliferative activity in HER2-amplified cell lines and showed a synergistic effect with trastuzumab in trastuzumab-resistant models^[72,73]. Based on these preclinical studies, a clinical trial in gastric cancer is in progress (NCT01402421, ClinicalTrials.gov).

Pembrolizumab

Pembrolizumab is a humanized monoclonal IgG4 directed against programmed death-1 (PD-1), mainly expressed on the cell surface of regulatory T cells.

PD-1 receptor is an immune-checkpoint receptor engaged by two known ligands, PD-L1 and PD-L2. Engagement of one of these ligands to the receptor inhibits T cell activation and eventually leads to apoptosis, which results in a blunted immune response in the tumor microenvironment^[74-76]. Interruption of the engagement of the ligands to their receptors using the anti-PD-1 monoclonal antibody can reverse the inhibition of the immune response, and this approach has been successful in the treatment of many cancers. Evaluation of the efficacy of pembrolizumab in patients with AGC is now underway in an international, multicenter, open-label phase 2 trial with treatment naïve patients or patients who received at least two prior chemotherapies (NCT02335411, Clinicaltrial.gov).

BEYOND ONGOING TRIALS

Clinical trials in molecular targeted agents for patients with AGC are reviewed in this article. The introduction of trastuzumab for a combination immune-chemotherapy in patients with AGC has taken a step forward in improving treatment outcomes. However, several limitations have been suggested in application of molecular targeted therapy.

Although abundant data from clinical trials with immune-chemotherapy in AGC has been reported, the efficacy according to the combination of target agents and chemotherapeutic agents has been different. Monotherapy with ramucirumab or a combination therapy with paclitaxel showed encouraging results in the second-line treatment. However the application of the same agent in combination with mFOLOX in the front-line treatment failed to show significant benefit^[61,62]. On the other hand, lapatinib showed clinical advantage in combination with capecitabine or paclitaxel at the second-line or first-line treatment, respectively, in patients with breast cancer^[77-79]. In contrast to the results in breast cancer, no benefit was shown in AGC in combination with paclitaxel or capecitabine plus oxaliplatin^[38,39]. These different responses suggest the presence of different mechanisms of action by which the combination therapies exert their effects. Because signaling pathways activated by the ligand binding may be altered by different combinations of immune-chemotherapeutic agents, the possible molecular mechanisms responsible for resistance should be clarified for the right combination of therapeutic agents. Drug interaction is another possible reason. Lordick *et al.*^[20] described that one of the reasons for failure of the EXPAND trial may be related to the negative pharmacokinetic interaction between capecitabine and cetuximab, and suggested the importance of choice and schedule of fluoropyrimidine in combination with cetuximab.

The tumor component, such as molecular heterogeneity, is also an important factor to be considered. A recent study divided gastric cancer into four molecular classifications, Epstein-Barr virus positive tumor,

microsatellite unstable tumor, genomically stable tumors, and tumors with chromosomal instability^[80]. Each classification has distinct molecular features, and future clinical trials should be performed in homogeneously defined subtypes of patients to raise the quality and achieve improved outcomes.

Appropriate selection of patients is also thought to be crucial before planning the clinical trials. The importance of selecting patients is highlighted in the ToGA study, in which clinical benefit was proven in selected patients overexpressing HER2^[3]. In addition, expression of MET was reported to be a prognostic marker in AGC patients treated with rilotumumab, suggesting the importance of indication selection for molecular targeted therapy^[4].

CONCLUSION

Despite several limitations, molecular targeted therapy is now regarded as an essential component in the treatment of cancers. In gastric cancer, numerous targeting agents have been examined in clinical trials since the introduction of trastuzumab. However, few targeted agents have been successful in establishment as a standardized therapy^[3]. Resistance to trastuzumab is an emerging issue to be solved and a considerable number of preclinical studies and clinical studies are now underway to overcome this limitation. Selection of patients should always be taken into consideration when designing clinical trials given that the molecular characteristic of gastric cancer is heterogeneous. By selecting targeted agents on the basis of known molecular mechanisms, a more potent activity of the molecular target agents would be expected.

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