

From Antibody Drugs to RNA Interference and Nanotherapy: The Evolution and Integration of Targeted Strategies in Gastric Cancer

Songlin Wang^{1,*}, Yinhong Dong^{1,*}, Lan Ma^{2-4,*}, Jiangfeng Wu²⁻⁴, Daoguo Long¹, Zhao Mei^{5,6}, Chuanlin Jiang¹, Xiaoguang Lyu^{1,7}

¹Zhongxiang People's Hospital, Zhongxiang, People's Republic of China; ²Hubei Key Laboratory of Tumor Microenvironment and Immunotherapy, China Three Gorges University, Yichang, People's Republic of China; ³College of Basic Medical Science, China Three Gorges University, Yichang, People's Republic of China; ⁴Institute of Organ Fibrosis and Targeted Drug Delivery, China Three Gorges University, Yichang, People's Republic of China; ⁵Department of Pharmacy, The First College of Clinical Medical Science, China Three Gorges University, Yichang, People's Republic of China; ⁶Department of Pharmacy, Yichang Central People's Hospital, Yichang, People's Republic of China; ⁷Department of Gastroenterology, Renmin Hospital of Wuhan University, Wuhan, People's Republic of China

*These authors contributed equally to this work

Correspondence: Chuanlin Jiang; Xiaoguang Lyu, Email 13677265887@139.com; dawnlxg@whu.edu.cn

Abstract: Gastric cancer is a leading cause of cancer death worldwide. Targeted therapy is shifting from blocking signaling pathways to precise gene regulation. This review covers three key areas: antibody drugs, RNAi, and nanocarriers. Anti-HER2 agents, VEGFR2 inhibitors, Claudin 18.2-directed therapies and PD-1/PD-L1 inhibitors improve survival in biomarker-selected patients—but resistance and low response rates (ORR < 20% in unselected groups) limit their use. RNAi silences key oncogenes to reverse chemoresistance and reprogram the immunosuppressive tumor microenvironment, but it suffers from rapid degradation, poor endosomal escape, off-target effects, and weak tumor penetration. Nanocarrier systems—including lipid nanoparticles (LNPs), stimuli-responsive micelles, and mesoporous silica—combined with aptamers enable targeted delivery of therapeutic antibodies and siRNA, penetration across stromal barriers, and controlled, tumor microenvironment-triggered cargo release. Altogether, the future of gastric cancer therapy is moving beyond monotherapy to a five-part strategy: antibody targeting, RNA silencing, nanocarrier delivery, AI-guided decisions, and immune remodeling.

Keywords: gastric cancer, target therapy, RNA interference, siRNA, shRNA, nanoparticles

Gastric cancer remains a significant global health challenge—it is the fifth most commonly diagnosed cancer and the fourth leading cause of cancer-related death worldwide, with 684,000 lives lost in 2022 (GLOBOCAN).¹ In China, gastric cancer is the third in incidence and the second in mortality—accounting for 478,000 new cases and 373,000 deaths annually.^{2,3} Due to its asymptomatic nature in the early stage, 70% of patients are diagnosed at advanced stage, resulting in a five-year overall survival rate of less than 10%.⁴⁻⁷ These figures underscore an urgent need for more effective therapies to delay tumor progression and extend survival time.

Antibody-based drugs launched targeted therapy for gastric cancer. Trastuzumab validated HER2, ramucirumab enabled anti-angiogenesis, and immune checkpoint inhibitors achieved durable responses in biomarker-selected patients.⁸⁻¹¹ But resistance—due to target loss, pathway redundancy, or immune evasion—and low response rates in unselected patients remain key challenges. 10 RNA interference (RNAi) offers the capacity to silence pathogenic genes at the mRNA level.¹² In gastric cancer, siRNAs directed against HER2, c-MET, or KRAS mutants have shown promising results in preclinical studies. 13 The translational barrier, however, is delivery: naked nucleic acids are rapidly degraded, poorly internalized, and prone to off-target effects.^{12,13} It is here that nanotechnology becomes indispensable. Nanocarriers—such as lipid nanoparticles and polymeric micelles—protect siRNA from degradation, enhance cellular

uptake, and enable controlled release.^{14,15} They can also penetrate the dense gastric tumor stroma and respond to microenvironmental cues like low pH or elevated enzyme levels.^{14,16} Most promising is delivering antibodies and siRNA in the nanocarrier to combine extracellular targeting with intracellular gene silencing.^{13,17–19}

This review traces the evolution from antibody drugs to RNAi and nanotherapeutics in gastric cancer. We argue that monotherapy is inherently limited and that the future lies in synergistic integration: antibodies for targeting, RNAi for gene silencing, and nanotechnology for precision delivery.^{14,16}

Clinically Validated Molecular Targets in Gastric Cancer for Targeted Therapy and Immunotherapy (Table 1)

The Following are the Crucial Molecular Targets

Targeting the HER2 Receptor

HER2 (ErbB2), a ligand-independent receptor tyrosine kinase, preferentially heterodimerizes with HER3 or EGFR, triggering constitutive activation of the PI3K/AKT/mTOR and RAS/RAF/MEK/ERK pathways in the absence of ligand binding.^{37–41} These cascades promote cell survival, proliferation, and immune evasion through effectors including mTORC1, Elk-1, c-Myc, and STAT3.⁴²

PI3K/AKT/mTOR Pathway

When HER2 heterodimerizes with HER3, it recruits and phosphorylates the PI3K regulatory subunit, thereby activating the enzyme. Activated PI3K converts PIP2 to PIP3, which recruits and activates AKT (protein kinase B). AKT then promotes cell survival and suppresses apoptosis by phosphorylating key targets—including BAD and caspase-9—and enhancing pro-survival signaling.^{43–45} AKT also activates mTORC1, driving protein synthesis, cell growth, proliferation, and metabolic adaptation.^{46–48}

RAS/RAF/MEK/ERK Pathway

HER2 activation recruits Grb2-SOS complex, triggering RAS activation. Active RAS activates RAF (MAPKKK), which phosphorylates and activates MEK (a MAPKK); MEK then phosphorylates and activates ERK (a MAPK). Nuclear translocation of activated ERK leads to phosphorylation of transcription factors—including Elk-1, c-Fos, and c-Myc—driving G1-to-S-phase cell cycle progression, proliferation, differentiation, and motility.^{46,49} Both pathways also phosphorylate STAT3, thereby modulating JAK/STAT signaling to promote immune evasion and reshape the tumor microenvironment.⁵⁰

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Targeting the VEGF Receptor (VEGFR)

The VEGF family includes VEGF-A, VEGF-C, and VEGF-D. VEGF-A drives angiogenesis via VEGFR-1/VEGFR-2 (KDR); VEGF-C/VEGF-D drive lymphangiogenesis via VEGFR-3.¹⁰ In gastric cancer, VEGF-A and VEGFR-2 are overexpressed relative to adjacent normal tissue and correlate with microvessel density (MVD). This promotes leaky vasculature, fueling hypoxia, acidosis, invasion, and metastasis: high MVD supports tumor growth and spread—VEGF enhances permeability to facilitate intravasation; VEGF-C/VEGFR-3 drives lymphangiogenesis and lymph node metastasis. VEGF also induces an immunosuppressive microenvironment by impairing dendritic cell maturation and T-cell infiltration, expanding Tregs, and upregulating PD-L1—contributing to “cold tumor” resistance to checkpoint inhibitors.^{10,51,52} Consequently, targeting VEGF/VEGFR (eg, fruquintinib [FRUTIGA trial, HR 0.67] and surufatinib) improves survival in heavily pretreated Chinese patients.⁵³

Targeting the Claudin 18.2

CLDN18.2 is a tight junction protein normally confined to the apical surface of gastric epithelial cells, maintaining epithelial barrier integrity and polarity.^{30,54} In gastric cancer, loss of polarity exposes it across the entire tumor cell membrane—converting it from an immunologically “hidden” to a “conspicuous” target.^{5,55} Epigenetic derepression

Table I Gastric Cancer Targeted Drugs Currently Under Clinical Investigation

Target	Drugs/Treatment	MedianOS (m)	MedianPFS (m)	Grade≥3AEs	KeyTrials
HER2	Zanidatamab+chemotherapy. ²⁰	36.5	12.5	79% (diarrhea, neutropenia, thrombocytopenia)	Phase II trial (NCT03929666)
	Trastuzumab+chemotherapy. ^{21,22}	13.8~20.5	8.1	73% (neutropenia, hand-foot syndrome, anemia, emesis)	Phase III ToGA study
	T-DXd. ²³⁻²⁵	14.7	6-8	85.7% (neutropenia, anemia, anorexia, ILD thrombocytopenia)	Phase III ToGA study
	Ramucirumab+TAX ^{23,26}	11.4	4.4	92% (neutropenia, leukopenia, hypertension, weakness/fatigue, anemia)	Phase III RCT
VEGF/VEGFR	Apatinib+ Camrelizumab+SOX. ^{27,28}	18.0~19.6	7.8	86.5% (neutropenia, leukopenia, hypertension, hepatic dysfunction)	Phase II CAP 06 Research (Exclusively for the specific subtype characterized by AFP positivity)
	Fruquintinib+TAX. ²⁹	9.6	5.6	83.5% (neutropenia, leukopenia, anemia, hypertension, hand-foot syndrome)	FRUTIGA phase III trial
	AK104 (PD-1/CTLA-4) +Pulocimab +TAX	12.9	6.8	55.2% (neutropenia, leukopenia, hypertension)	AK109-201 phase II
	Ramucirumab monotherapy	5.2	2.1	74% (hypertension, abdominal pain, fatigue, anemia, ascites)	REGARD phase III trial
Claudin 18.2	Zolbetuximab+mFOLFOX6. ³⁰	18.2	10.6	87% (nausea, vomiting, neutropenia, anemia, thrombocytopenia, diarrhea)	SPOTLIGHT phase III trial
	Zolbetuximab+CAPOX. ^{30,31}	14.39	8.27	82.8% (nausea/vomiting, neutropenia, thrombocytopenia)	GLOW phase III trial
PD-1/PD-L1	Pembrolizumab+ Trastuzumab+chemotherapy. ³²	20.1	10	59% (neutropenia, anemia, nausea/vomiting, diarrhea)	KEYNOTE-811 phase III trial
	Sintilimab+chemotherapy. ³³	18.4	15.2	59.9% (neutropenia, leukopeni, anemia, thrombocytopenia)	ClinicalTrials.gov Identifier: NCT03745170
	Tislelizumab+CAPOX/FP. ³⁴	15.0	6.9	66.7% (neutropenia, thrombocytopenia, anemia, leukopenia)	RATIONALE-305 phase III Clinical Trial
	Cadonilimab+chemotherapy. ^{35,36}	15	7	65.1% (neutropenia, leukopeni, anemia, thrombocytopenia, hepatic dysfunction)	COMPASSION-15 phase III Clinical Trial

drives its overexpression: promoter hypomethylation lifts transcriptional silencing, enabling stable, high-level expression. This fuels oncogenesis via three key mechanisms: (1) PI3K/AKT activation → suppressed apoptosis and enhanced survival/proliferation; (2) ERK/MAPK-driven AP-1 activation → upregulation of cell-cycle genes; and (3) CLDN18-ARHGAP26 fusion (in ~15% of diffuse-type cases) → constitutive RhoA activation, cytoskeletal remodeling, increased motility, EMT, and invasion. CLDN18.2+ tumors also create an immunosuppressive microenvironment—recruiting Tregs, M2 macrophages, and MDSCs; impairing dendritic cell maturation and T-cell infiltration; and disrupting antigen presentation—enabling immune escape.^{56,57} Critically, its stable epigenetic expression—and selective pressure in metastatic niches—elevates CLDN18.2 in metastases, reinforcing its value as a metastatic target.^{54,58} Clinically, zolbetuximab + chemotherapy improved PFS (7.5 vs 5.3 mo) and OS (14.4 vs 12.2 mo) in SPOTLIGHT and GLOW trials, leading to FDA/NMPA approval in 2024.^{30,31} Claudin 18.2-directed CAR-T (CT053) achieved a 57.1% ORR in heavily pretreated patients.

Target of Immune Checkpoint Inhibitor Therapy

PD-1/PD-L1

In gastric adenocarcinoma, PD-L1 is upregulated by inflammatory cytokines—including IFN- γ and IL-6—via JAK/STAT and NF- κ B signaling, enabling immune evasion. Approximately 10% of cases are EBV-positive; these tumors consistently overexpress PD-L1 and show enhanced response to PD-1 blockade. PD-1 is expressed on T cells, B cells, and NK cells, whereas its ligand PD-L1 is upregulated on tumor cells and antigen-presenting cells (eg, macrophages, dendritic cells). PD-1/PD-L1 binding recruits SHP-1 and SHP-2 phosphatases, inhibiting TCR signaling—thereby suppressing T-cell proliferation, cytokine production (IFN- γ , IL-2), and effector function—and facilitating tumor immune escape.⁵⁹

Analysis of the Current Limitations in Targeted Therapy for Gastric Cancer

Dependence on Specific Target Expression

Targeted therapy requires uniform, high-level expression of tumor-specific markers—but many gastric cancers show low or heterogeneous expression. For example, only 7.3–20.2% of patients overexpress HER2, typically in focal, intensely stained regions.⁶⁰ Claudin 18.2 positivity varies widely (30–70%) depending on the IHC assay and cutoff: early studies using the 43–14A antibody reported >60% positivity, whereas the VENTANA SP97 assay (≥ 2 in $\geq 75\%$) yields 35–50% in both Western and Asian cohorts.⁶¹ This variability underscores the urgent need for standardized, companion diagnostic—approved testing.

Prominent Drug Resistance Challenges

Targeted therapies face primary and secondary resistance. Primary resistance arises from compensatory pathway activation (eg, MAPK, PI3K/AKT) or baseline mutations; secondary resistance stems from acquired target-gene mutations (eg, HER2 L755S/V777L), phenotypic shifts, epigenetic changes, or co-amplifications (MYC, CCNE1, CDK12).^{62–65} For example, trastuzumab resistance often involves HER2 alterations, IGF-1R upregulation, or bypass signaling via MET/FGFR. Apatinib may promote immune evasion by activating MAPK or inhibiting STAT3/PKM2.^{64–66} VEGF further contributes to a “cold tumor” microenvironment by impairing dendritic cell maturation and T-cell infiltration while expanding Tregs.⁶⁷

Abnormal Tumor Vasculature

Tumor blood vessels are chaotic, tortuous, leaky, and dysfunctional—causing poor drug delivery and an immunosuppressive microenvironment. Hypoxia from abnormal perfusion recruits immunosuppressive cells and factors, impairing immune cell infiltration and function. Anti-VEGF agents (eg, ramucirumab, apatinib) were initially intended to “starve” tumors. However, short-term, moderate VEGF inhibition could induce transient “vascular normalization”—a temporary restoration of vessel structure and function. Yet this window is narrow, dynamic, and highly variable across patients, making it hard to identify and easy to miss clinically. Moreover, tumors rapidly activate bypass pathways (eg, FGF, Angiopoietin) upon VEGF blockade, driving vascular regrowth and undermining combination therapy. This adaptation not only reduces efficacy but may also promote a more invasive phenotype.

Safety Concerns

Some targeted agents cause significant toxicities. Topoisomerase inhibitors like exatecan induce grade ≥ 3 neutropenia in 20–30% of patients.⁶⁸ CCR8 $\times$ CLDN18.2 bispecific antibodies (eg, LM-108) are still in early development; their long-term safety and efficacy remain undefined. Bemarituzumab causes vision impairment, dry eye, or corneal damage in $>25\%$ of patients—mostly reversible but potentially affecting adherence.⁶⁹ Long-term data on Claudin 18.2-targeted therapies are scarce, so the risks of chronic toxicities (eg, interstitial lung disease) are unclear.

Economic and Accessibility Barriers

High treatment costs limit the broad use of targeted therapies. Trastuzumab costs over 300,000 yuan annually; CAR-T therapy—typically priced above 1 million yuan per treatment—presents a substantially higher financial barrier, critically constraining patient access. So far, only one Claudin 18.2-targeted agent, zolbetuximab (not trastuzumab), has received global marketing approval; ADCs and CAR-T therapies remain under clinical investigation. First-line guidelines increasingly recommend “targeted therapy + immunotherapy + chemotherapy”—a regimen far costlier than chemotherapy alone. Biomarker testing (eg, for Claudin 18.2) adds further financial strain.

RNA Interference (RNAi) Technology Demonstrates Significant Potential in the Field of Gastric Cancer Therapy (Table 2); However, It Also Encounters Various Challenges

Key Advantages of RNAi-Based Therapeutic Approaches in Gastric Cancer Management

- 1) RNAi achieves high target specificity via Watson–Crick base pairing. It selectively silences oncogenic drivers such as NAT10, MALAT1, and UCA1—targets inaccessible to conventional small molecules or antibodies—while sparing normal tissues where the target is absent or barely expressed.¹⁰⁴ RNAi also precisely inhibits gastric cancer-linked genes such as AE1 and lncRNA CASC15 (lncRNA CASC15) without harming healthy gastric tissue. In a mouse gastric cancer model, AE1-targeted siRNA reduced tumor incidence from 62–70% to 15.8% ($p < 0.001$), with no pathology-confirmed damage to benign gastric mucosal lesions.¹⁰⁵
- 2) This technology effectively addresses drug resistance commonly encountered in conventional therapies. For example, knockdown of the lncRNA NCK-AS1 reverses cisplatin resistance and improves chemotherapeutic sensitivity.¹⁰⁶ Targeting the E3 ubiquitin ligase SYVN1 has also been shown to significantly inhibit gastric cancer cell (GC cell) proliferation.¹⁰⁷

Table 2 Recent Advances in Experimental Research on RNAi Therapy for Gastric Cancer

RNAi Therapies	Remarks
B7-H7 siRNA	Suppression of B7-H7 expression markedly enhances the chemosensitivity of MKN-45 GC cells to docetaxel, thereby improving therapeutic response and highlighting its pivotal regulatory role in drug resistance. ⁷⁰
CACNA2D1 siRNA	CACNA2D1 encodes the $\alpha 2\delta$ subunit of voltage-gated calcium channels. CACNA2D1 siRNA suppresses tumor growth, induces apoptosis, and inhibits migration and invasion in HGC-27 cells in mice. ⁷¹
ADAR1 siRNA	ADAR1 siRNA inhibits metastasis of GC cells and reverses cisplatin resistance by downregulating AZIN1, thereby suppressing EMT-related markers. ⁷²
Bcl2 siRNA	Bcl-2 siRNA selectively knocked down the Bcl-2 gene in a stomach cancer mouse model, leading to increased apoptosis and tumor remission. ⁷³
IQGAP3 siRNA	IQGAP3 siRNA markedly suppressed invasion and anchorage-independent growth in MKN-1 and TMK-1 GC cells. ⁷⁴
OSGIN2 siRNA	OSGIN2 siRNA inhibited tumor cell proliferation and induced cell cycle arrest. It was also associated with reduced infiltration of tumor-infiltrating lymphocytes (TILs) and impaired antitumor immune function. ⁷⁵

(Continued)

Table 2 (Continued).

RNAi Therapies	Remarks
EHMT2 siRNA	EHMT2 siRNA induced G1 phase cell cycle arrest and attenuated the proliferation of GC cells. ⁷⁶
EZH2 siRNA	EZH2 siRNA significantly inhibited tumorigenesis and suppressed proliferation, wound healing, migration, and colony formation in AGC and MGC-803 GC cell lines. ⁷⁷
CPLX1 siRNA CPLX1 shRNA	In vivo experiments demonstrated that RNAi-mediated silencing of CPLX1 markedly inhibited tumor progression in MKN-I GC cells. Consistently, CPLX1-specific shRNA significantly reduced the growth of tumors derived from MKN-I cells. Furthermore, treatment with CPLX1-targeting siRNA substantially enhanced responsiveness to fluorouracil across multiple GC cell lines, including KATOIII, OCUM1, MKN1, and AGS. ⁷⁸
STIM1 siRNA	In BGC-823 and SGC-7901 GC cells, the anticancer activity of DIM was counteracted by STIM1 siRNA knockdown via inhibition of p-AMPK-triggered ER stress. ⁷⁹
LRRC8A siRNA	LRRC8A siRNA suppressed proliferation and cell migration, and enhanced apoptosis in human MKN-74 and NUGC-4 GC cell lines. ⁸⁰
ILF2 siRNA	ILF2 siRNA significantly inhibited proliferation and colony formation in HGC-27 and AGS human GC cells. ⁸¹
TRAF6 shRNA	TRAF6 shRNA inhibited the nuclear translocation of NF- κ B p65, suppressed the proliferation of 5-FU-resistant GC cells, and blocked cell cycle progression in these cells. ⁸²
CLEC11A shRNA	Knockdown of CLEC11A using shRNA inhibited cell cycle progression, migration, and invasive capacity, and attenuated gastric cancer growth in vivo. Furthermore, it enhanced the recruitment of cytotoxic CD8+ T cells and helper CD4+ T cells into tumor tissues, while reducing the presence of M2 macrophages, myeloid-derived suppressor cells (MDSCs), and regulatory T cells (Tregs). ⁸³
PSMA3-ASI shRNA	PSMA3-ASI shRNA suppressed GC cell proliferation, migration, and invasion, enhanced apoptosis, and inhibited tumor growth both in vitro and in vivo by inducing oxidative stress and downregulating the Nrf2 signaling pathway. ⁸⁴
FAM117B shRNA	FAM117B shRNA promoted NRF2 ubiquitination, thereby accelerating NRF2 binding to KEAP1 and its subsequent degradation; this inhibition of the KEAP1/NRF2 pathway suppressed GC cell growth and enhanced chemosensitivity. ⁸⁵
VacA siRNA CagA siRNA	VacA siRNA and CagA siRNA silence the expression of <i>H. pylori</i> 's vacuolating cytotoxin A (VacA) and cytotoxin-associated gene A (CagA), thereby aiding gastric cancer treatment. ⁸⁶
HOXA10 shRNA METTL3 shRNA	HOXA10 shRNA and METTL3 shRNA reduced HOXA10 expression and inhibited EMT and GC metastasis by modulating the TGF β 2/Smad/METTL3 axis. ⁸⁷
ESM1 shRNA	ESM1 shRNA inhibited proliferation and migratory ability of HUVECs co-cultured with AGS cells and reduced tube formation. ⁸⁸
RAP2A siRNA	RAP2A siRNA inhibited DDP resistance in GC cells by modulating cellular proliferation, migration, and invasion, as well as apoptosis and DNA damage induced by DDP treatment. ⁸⁹
UCA1 siRNA	UCA1 siRNA enhanced cisplatin-induced apoptosis and reduced cisplatin resistance in human GC cells. ⁹⁰
MALAT1 siRNA MALAT1 shRNA	MALAT1 siRNA suppressed proliferation, migration, and invasion of GC cells, promoted apoptosis, and reduced chemotherapy-induced autophagy. Additionally, MALAT1 shRNA inhibited autophagy in GC cells by regulating ATG5 expression via its interaction with miR-30e. ^{91,92}
HAGLR siRNA	HAGLR siRNA inhibited GC proliferation, migration, and 5-FU resistance by upregulating miR-338-3p expression through sponging it in GC cells. ⁹³
SNHG16 siRNA PTBPI shRNA	SNHG16 and PTBPI silencing increased 5-FU sensitivity of GC cells by binding to miR-506-3p and inhibiting glucose uptake. ⁹⁴
ANXA1 siRNA	ANXA1 siRNA enhanced the sensitivity of oxaliplatin (OXA)-resistant GC cells to OXA and increased the apoptotic rate in AGS/OXA and HGC27/OXA cells via the PI3K/AKT/mTOR signaling pathway. ⁹⁵
TRIM14 shRNA	TRIM14 shRNA reduced tumor size and weight and inhibited migration and proliferation of GC cells in 5-FU-resistant GC mice. ⁹⁶
UCA1 siRNA EZH2 siRNA	UCA1 siRNA decreases GC cell proliferation and promotes cisplatin-induced apoptosis by downregulating EZH2 expression; conversely, EZH2 knockdown also enhances cisplatin-induced apoptosis in GC cells. ⁹⁷
SNHG16 siRNA	SNHG16 siRNA inhibited proliferation, cell cycle progression, migration, and invasion of GC cells, and promoted apoptosis. ⁹⁸
Si-circ_0081143	Si-circ_0081143 reduced drug resistance (DR), inhibited proliferation, migration, and invasion of DR-resistant GC cells by targeting miR-129-2-3p and suppressing its function. ⁹⁹
Circprx1 siRNA	Reversal of doxorubicin resistance in doxorubicin-resistant GC cells through targeting miR-3064-5p or regulating PTPN14. ¹⁰⁰

(Continued)

Table 2 (Continued).

RNAi Therapies	Remarks
URI siRNA	URI siRNA increased the sensitivity of GC cells to cisplatin by inhibiting proliferation, promoting apoptosis, and enhancing cisplatin-induced DNA damage; it also enhanced drug sensitivity via the ATM/CHK2 pathway. ¹⁰¹
DCLK1 shRNA SNHG1 shRNA	DCLK1 shRNA and SNHG1 shRNA inhibited the EMT process in GC cells through the DCLK1-mediated Notch1 signaling pathway and suppressed cell migration and invasion. ¹⁰²
E2F1 siRNA E2F1 shRNA	Silencing E2F1 decreased expression of drug resistance- and apoptosis- related proteins, enhanced the sensitivity of drug-resistant GC cells to paclitaxel combined with cisplatin, promoted apoptosis, and suppressed proliferation. ¹⁰³

- 3) RNAi technology demonstrates the capacity to modulate the tumor microenvironment (TME). Silencing of the NAT10 gene interrupts the CXCL2/STAT3 signaling axis, thereby suppressing M2 macrophage polarization—with the proportion of M2 macrophages in liver metastases declining from 42% to baseline levels—resulting in a 78% reduction in liver metastasis.¹⁰⁸ Furthermore, the novel delivery system TMAB3, which carries a RIG-I agonist RNA, enhances CD8⁺ T cell infiltration and effectively reverses the “cold tumor” phenotype, thereby augmenting anti-tumor immune responses.¹⁰⁹
- 4) The technology is associated with low toxicity and minimal side effects. For example, under physiological conditions, AE1 is exclusively expressed in red blood cells, and siRNA delivery vehicles are unable to cross the bone marrow-blood barrier (MBB).¹⁰⁵ Consequently, no hemolytic or hepatorenal toxicities were observed in preclinical animal studies.

Current Challenges and Limitations of RNAi Therapy

- 1) Instability of RNA molecules: Unprotected RNA is highly susceptible to degradation by endogenous nucleases in vivo, leading to a short half-life in blood and tissues. This instability prevents the maintenance of sustained therapeutic concentrations, thereby compromising treatment efficacy.^{110–112}
- 2) Suboptimal delivery efficiency: Conventional delivery systems—such as liposomes—exhibit limited targeting specificity. The lack of highly efficient and selective delivery platforms hinders the precise accumulation of RNAi molecules at tumor sites, thereby constraining their broad clinical applicability.¹⁰⁰ Consequently, numerous studies are currently focused on developing highly specific, tumor-targeted delivery systems—including aptamers and nanoparticles—to enhance the efficient delivery of siRNA and shRNA to tumor sites (Figure 1 and Table 3).^{111,113–115} Tumor-targeted lipid nanoparticles (such as Onpatro-like platforms) and GalNAc-conjugated siRNAs are currently undergoing phase I/II trials for solid tumors. Aptamer-siRNA chimeras specific to gastric cancer and pH-responsive nanoparticles have achieved over 50-fold tumor enrichment in orthotopic models.
- 3) Potential off-target effects: RNAi molecules may hybridize with non-target mRNA sequences, leading to unintended gene silencing and posing potential risks of adverse physiological effects.^{115,122}
- 4) Immune activation potential: Exogenous RNA can activate the innate immune system, inducing the production of inflammatory mediators that may compromise both the safety and tolerability of RNAi-based therapies.¹²²
- 5) Production complexity and cost constraints: The synthesis, purification, and quality control of RNAi therapeutics involve technically demanding processes, contributing to high manufacturing costs. These challenges hinder large-scale production and widespread clinical adoption. Moreover, complementary genetic testing (eg, assessment of NAT10 and KLF5 expression levels) imposes additional economic and therapeutic burdens on patients.
- 6) Inadequate long-term safety data: The majority of current evidence is derived from animal models, and comprehensive long-term safety evaluations in humans—particularly regarding immunogenicity and organ-specific accumulation—remain lacking.¹²² Furthermore, epigenetic targets such as NAT10 may disrupt normal RNA modification pathways in healthy cells, raising concerns about potential yet undefined biological risks.

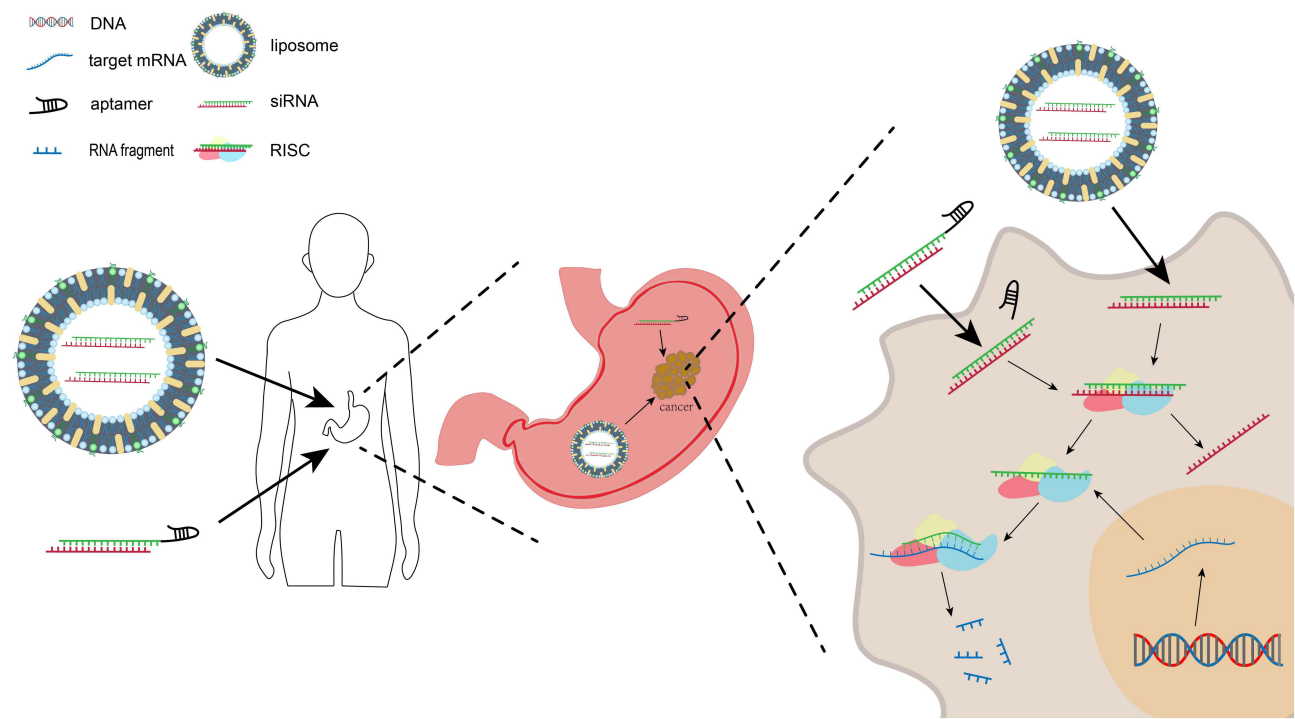


Figure 1 Targeted delivery systems, such as aptamers and nanoparticles, improve the efficient and precise delivery of siRNA to tumor sites, thereby enhancing therapeutic efficacy and enabling potent antitumor effects.

Nanoparticle-Based Therapeutics for Gastric Cancer: Advancing from Passive Targeting to Intelligent, Multimodal Synergy

Evolution of Targeting Strategies: From Active Targeting to Microenvironment-Responsive Delivery

Conventional passive targeting relies on the enhanced permeability and retention (EPR) effect. However, EPR effect is highly heterogeneous across and within gastric tumors, limiting its reliability. To improve targeting, nanoparticles are

Table 3 Recent Advances in Experimental Research on Targeted Delivery of siRNA and shRNA for RNAi Therapy in Gastric Cancer

Targeted Delivery of siRNA and shRNA	Remarks
Aptamer-siRNA Chimera/PEI/5-FU/Carbon Nanotube/Collagen Membranes	Aptamer-siRNA chimera specifically targets GC cells, enabling targeted delivery of 5-fluorouracil (5-FU) and silencing of drug-resistant-associated genes. The Chim/PEI/5-FU/CNT nanoparticles induce apoptosis and suppress invasion and proliferation in 5-FU-resistant GC cells. In vivo, the Chim/PEI/5-FU/CNT/collagen membrane significantly inhibits mitogen-activated protein kinase (MAPK) signaling and effectively suppresses peritoneal dissemination of 5-FU-resistant GC. ¹¹⁶
GADA/NAVI/siRNA	The GADA/NAVI/siRNA therapeutic regimen profoundly suppressed Bcl2 expression, thereby triggering robust apoptosis in cancer cells. This multimodal strategy has demonstrated superior efficacy in gastric cancer treatment compared to both free siRNA and NAVI monotherapy, highlighting its potential as a transformative approach in oncology. ¹¹⁷
PLGA-PEG(si-circRHBD1) NPs	PLGA-PEG(si-circRHBD1) NPs accurately delivered drugs to the tumor site in C57BL/6 GC mice bearing gastric cancer (GC), exhibiting significant inhibition of tumor growth without systemic toxicity. ¹¹⁸
Codelivery system of As2O3 and HER2-siRNA via RNPs	AH RNPs are capable of escaping from lysosomes to facilitate the cytosolic delivery of siRNA and As2O3, attributable to the pH-responsive dissolution of the CaP shell. These RNPs synergistically suppress gastric cancer growth and metastasis without inducing any evident adverse effects during treatment. ¹¹⁹
Bcl-xL shRNA and DOX co-delivery using AS1411 aptamer-conjugated NP	This system achieves precise targeting of GC by binding to nucleolin receptors, significantly reduces Bcl-xL protein expression, and demonstrates excellent tumoricidal efficacy with low cytotoxicity. ¹²⁰
Luc lipoplexes, composed of CLs containing Luc shRNA	Luc lipoplexes, composed of cationic liposomes (CLs) encapsulating Luc shRNA, can effectively treat peritoneal metastasis of gastric cancer without inducing adverse events. ¹²¹

surface-modified with ligands—such as folate, transferrin, antibodies, or aptamers—that bind receptors overexpressed on GC cells. For example, folate receptor alpha (FR α) is overexpressed in >60% of gastric cancers; folate-conjugated nanoparticles increase tumor accumulation 3-5-fold compared with non-targeted controls in mouse models.¹²³ Microenvironment-responsive nanoparticles offer greater precision: they release drug only in response to gastric tumor-specific cues—such as low pH (6.0–6.8), high GSH concentrations (2–10 mM), elevated ROS, or overexpression of matrix metalloproteinases MMP-2 and MMP-9—using cleavable linkers like hydrazone, disulfide, or thioacetal bonds. A pH/ROS dual-responsive docetaxel nanomicelle achieved 79.6% tumor growth inhibition in subcutaneous gastric cancer xenografts—significantly higher than that of free docetaxel.¹²⁴

Overcoming Chemotherapy Resistance: From Gene Silencing to Pathway Intervention

Chemotherapy resistance is a major cause of treatment failure in gastric cancer. Nanoparticles can deliver both chemotherapeutic drugs and gene therapeutics—such as siRNA, miRNA, or CRISPR-Cas systems—to reverse resistance at its source. Two key strategies are employed: (1) Gene silencing—target ABC transporters (eg, P-gp) or anti-apoptotic proteins (eg, Survivin, Bcl-2); and (2) Pathway inhibition—block pro-survival signals such as Notch, Wnt/ β -catenin, or NF- κ B. For example, liposomal nanoparticles co-delivering P-gp siRNA and paclitaxel reduced the IC₅₀ by ~80% in resistant SGC7901/VCR cells.¹²⁵ Similarly, SPIONs functionalized with the WSGC peptide inhibited Notch signaling, suppressing tumor growth by >60% and enhancing apoptosis in xenografts.¹²³

Emerging Physical Therapies for Cancer Treatment: Integrating Photothermal Therapy, Ferroptosis Induction, and Immunomodulation

Nanoparticles serve as programmable theranostic platforms—functioning either as photonic energy transducers or as precision inducers of regulated cell death.

Photothermal Therapy (PTT)

Carbon quantum dots, MXene, and polydopamine nanoparticles absorb near-infrared light (808 nm or 1064 nm) to generate heat, raising tumor temperature above 50 °C and killing cancer cells. Limitation: poor tissue penetration limits their use in deep gastric tumors or peritoneal metastases.^{126,127}

Ferroptosis Therapy

Nanoparticles deliver Fe²⁺/Fe³⁺ or inhibit GPX4, thereby inducing lipid peroxidation and ferroptosis. The biomimetic nanocrystal TRNC@P+L co-delivers chemotherapeutic drugs and ferroptosis inducers, showing synergistic efficacy in preclinical models of gastric cancer peritoneal metastasis.¹²⁸

Immunomodulatory Nanoparticles

In a gastric carcinoma mouse model, M2pep-Cs NPs/Plerixafor nanoparticles suppress tumor progression by concurrently blocking the CXCL12-CXCR4 axis and reprogramming tumor-associated macrophages (TAMs)—specifically shifting them from immunosuppressive M2 phenotype to immunostimulatory M1 phenotypes—which enhances T-cell infiltration, activation, and tumor antigen recognition.¹²⁹

Nanometer-Scale Chemotherapeutic Agents: Advancing Therapeutic Efficacy While Minimizing Systemic Toxicity

The real-world study of NPMP (paclitaxel nanomicelles) combined with immunotherapy and chemotherapy—though non-randomized and uncontrolled—achieved an objective response rate (ORR) of 42.4%, exceeding the typical ORR of 25–35% observed with conventional chemotherapy. Neutropenia rates did not increase significantly compared with historical controls, suggesting that the nanomicellar formulation improves the therapeutic index through tumor-selective drug accumulation.¹³⁰ SN BioScience has received FDA Orphan Drug Designation for SNB-101 in gastric cancer—the third such designation for this novel nanoparticle formulation of SN-38. As the world's first polymer-based nanoparticle formulation of the highly water-insoluble anticancer agent SN-38, SNB-101 is designed to enhance efficacy and reduce

toxicity. The designation addresses a critical unmet medical need in gastric cancer, where advanced disease has limited treatment options and a 5-year survival rate of only 36%.

Future Development of Gastric Cancer Treatment (Figure 2)

Future directions in gastric cancer treatment focus on four key areas: 1) Precision oncology: Multi-omics profiling (genomics, proteomics, metabolomics) identifies targetable molecular drivers; CRISPR/Cas9 and animal models validate function and therapeutic potential; resulting agents—small molecules, mAbs, or bispecific antibodies—are matched to patients' molecular profiles (eg, mutations, protein expression, immune contexture) for stratified therapy. 2) Next-generation ADCs: Optimized DAR, cleavable/conditionally active linkers, and novel payloads improve tumor selectivity and safety; rational combinations—eg, with chemo, immunomodulators, or dual-target inhibitors—overcome resistance and broaden indications. 3) CLDN18.2-directed CAR-T: Now in randomized clinical trials (eg, CT041-ST-01), this “living drug” bypasses vascular dependence, enabling direct tumor cell killing. It shows durable survival benefit after standard therapy failure, with manageable CRS and no neurotoxicity in trials. Limitations remain: strict CLDN18.2+ selection ($\geq 40\%$ tumor cells), complex manufacturing/logistics, high cost, and persistent challenges of solid tumors—including immunosuppressive microenvironments and target heterogeneity.

Systemic therapy for gastric cancer has evolved from sequential monotherapy to rational combinations of immune checkpoint inhibitors, targeted agents, and chemotherapy. Aptamer- and nanocarrier-based delivery platforms represent a paradigm shift—enabling spatiotemporally controlled treatment, not merely incremental improvement. Current standard therapies (eg, anti-HER2, CLDN18.2-targeted, and dual PD-1/CTLA-4 blockade) provide clinical benefit but suffer from narrow therapeutic indices due to on-target/off-tumor toxicity. Aptamers—synthetic, high-affinity, low-immunogenicity nucleic acid ligands—precisely target gastric cancer drivers. Nanocarriers exploit both passive (enhanced permeability and retention, EPR) and active (ligand-mediated) tumor accumulation and release their payloads selectively in response to the tumor microenvironment—within the tumor parenchyma or the immunosuppressive stroma.

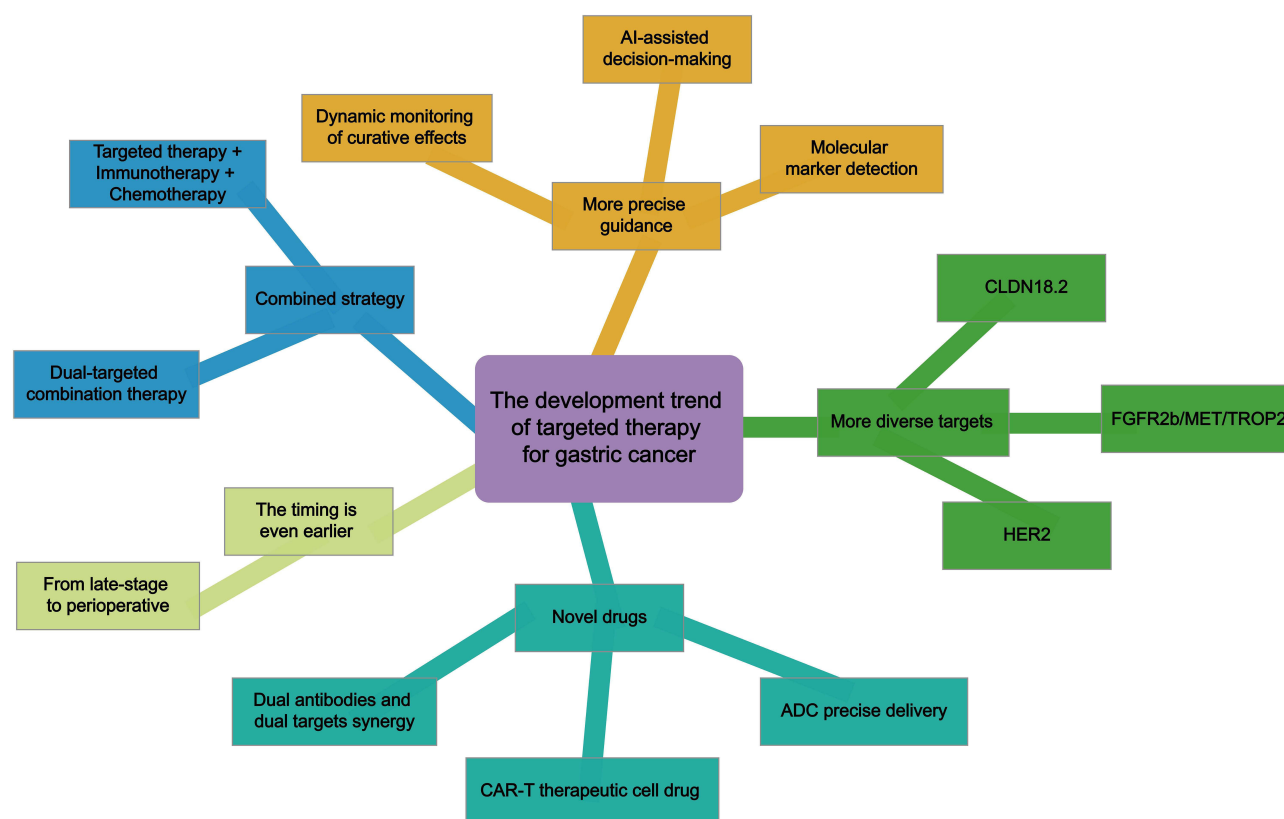


Figure 2 Trends in the Development of Targeted Therapy for Gastric Cancer.

Integrating these modalities enables intelligent nanosystems: either for localized intraperitoneal delivery or as multi-epitope tandem aptamer–drug conjugates (ADCs) that address gastric cancer’s core challenges—inter- and intra-tumoral heterogeneity and its strong propensity for peritoneal dissemination. This dual-strategy approach tackles three key clinical challenges in peritoneal metastases: (1) poor drug accumulation due to low peritoneal–plasma barrier permeability; (2) emergence of therapy-resistant clones under single-target treatment; and (3) systemic toxicities that limit dose intensity and duration. Clinical translation requires solving three hurdles: aptamer degradation by nucleases *in vivo*, RES-mediated nanoparticle clearance, and scalable GMP-compliant manufacturing with strict quality control. A phase I trial of this aptamer-nanocarrier platform in gastric cancer patients with peritoneal metastases is expected within five years. Primary endpoints include safety plus quantitative pharmacodynamic biomarkers—tumor-to-plasma drug ratio, target saturation kinetics, and spatially resolved intratumoral payload distribution. Critically, these mechanistic metrics—not just response rates—will determine whether the platform evolves from innovation to standard therapy.

The aptamer-nanodrug system’s key advantage in gastric cancer treatment is not antibody replacement, but programmable, precise targeting. Unlike antibodies, aptamers can both bind targets and execute logic operations—eg, releasing drugs in response to tumor microenvironment cues—via rational structural design. Yet current research focuses excessively on aptamer loading onto nanoparticles, overlooking a fundamental question: What target profiles do distinct gastric cancer molecular subtypes actually require? Future progress depends less on generating more aptamers and more on establishing a clinical-translational paradigm: begin with molecular subtyping, reverse-define target requirements, then custom-design the aptamer–nanoparticle system. Critically, this system must integrate into a closed loop with liquid biopsy-based early detection and post-surgical minimal residual disease monitoring—to translate precise targeting into real survival benefits.

Acknowledgments

This study was supported by the National Natural Science Foundation of China (Grant number [81670555]); State Key Laboratory of Medical Molecular Biology (Grant number [State Key Laboratory Special Fund 206024]); 2025 Yichang Medical and Health Research Project (No. A25-2-178); Chen Xiao-Ping Foundation for the Development of Science and Technology of Hubei Province (No. CXPJJH125001-2509).

Since English is not our native language, the article was refined with the assistance of Youdao Translate (Version: 11.0.0).

Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

Disclosure

The authors declare no conflicts of interest.

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