

Microsatellite instability and immunotherapy in gastric cancer: a narrative review

Yantao Duan, Dazhi Xu

Department of Gastric Surgery, Fudan University Shanghai Cancer Center, Shanghai, China

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Correspondence to: Dazhi Xu, PhD, MD. Department of Gastric Surgery, Fudan University Shanghai Cancer Center, 270 Dongan Road, Shanghai 200032, China. Email: xudzh@shca.org.cn.

Background and Objective: Immunotherapy is now a new treatment approach that is widespreadly accepted in the clinical practice, but the role in gastric cancer is still poorly elaborated. As mentioned in The Cancer Genome Atlas (TCGA) and Asian Cancer Research Group (ACRG) cancer classifications, the separate microsatellite instable (MSI) entirety is defined as a specific subtype of gastric cancer. In particular, the MSI hypermutator phenotype gastric cancer triggers immune response, allowing the specific molecular subset as a potential candidate for immunotherapy. The current review is to provide an updated overview of the available literature on the clinical implications of MSI in gastric cancer.

Methods: We performed an extensive literature research on PubMed regarding the molecular, immunotherapy, and prognostic characteristics of MSI in gastric cancer, selecting only English-language articles from January 11, 2017 to August 12, 2022.

Key Content and Findings: The complexity of gastric cancer puts forward higher demands for novel molecular-based individual therapeutics. MSI gastric cancers take up a relatively small patients population, and show distinctive clinicopathological features. MSI is significantly correlated with the response to immunotherapy in gastric cancer. The MSI cancers treated with immune checkpoint inhibitors show a favorable prognosis and MSI may serve as a biomarker for immunotherapy of gastric cancer.

Conclusions: MSI detection plays a promising role in guiding the immunotherapy of gastric cancer. However, this still needs to be further verified in larger prospective clinical trials.

Keywords: Microsatellite instability (MSI); mismatch repair genes system; immunotherapy; gastric cancer

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Introduction

Gastric cancer remains one of the most common cancer worldwide and is responsible for more than one million new cases and an estimated 769,000 deaths in 2020, ranking fifth in cancer incidence and fourth in cancer mortality (1). Surgical resection with adjuvant chemotherapy is widely acknowledged as an effective treatment for the early-stage gastric cancer. Surgery followed by chemotherapy may be an approach to treating localized gastric cancer, while in other parts of the world, neoadjuvant chemotherapy followed by surgery and adjuvant chemotherapy are also other approaches to gastric cancer treatment. In general, systemic therapy and chemoradiotherapy remain the standard first-line treatments in locally advanced or metastatic gastric cancer cases (2). Nevertheless, despite the advances of current comprehensive treatments, the 5-year survival rate remains very low, and a tremendous effort is still required for individualized treatment of gastric cancer and the improvement of clinical prognosis.

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Table 1 Search strategy summary

Items	Specification
Date of search	From January 11, 2017 to August 12, 2022
Databases and other sources searched	PubMed
Search terms used	Microsatellite instability, immunotherapy, and gastric cancer
Timeframe	Up to August 12, 2022
Inclusion and exclusion criteria	We only included studies published in English language
Selection process	The selection process was conducted independently by the authors

Gastric cancer is highly heterogeneous, and The Cancer Genome Atlas (TCGA) Research Network has classified gastric cancer into four subtypes based on a comprehensive molecular characterization: chromosome instability (CIN), microsatellite instability (MSI), genomically stable (GS), and Epstein-Barr virus (EBV) infected type (3). In addition, the Asian Cancer Research Group (ACRG) also proposes a novel gastric cancer molecular classification, identifying four molecular subtypes: MSI type, microsatellite stable with epithelial to mesenchymal transition features (MSS/ EMT), MSS with tumor protein p53 (TP53)-active (MSS/ TP53⁺) type and MSS with TP53-inactive (MSS/TP53⁻) type (4). Interestingly, both TCGA and ACRG researches distinguish the MSI entity as a separate and well-defined gastric cancer subgroup.

Tumor molecular classifications based on the comprehensive molecular profiles have more clinical significance in predicting treatment efficacy than traditional Lauren's classification and World Health Organization (WHO) classification, especially for immunotherapy (3,5). Over the past few years, immunotherapy has received great attention in cancer treatment, and has been proven to greatly improve the therapeutic effect and survival of gastric cancer. Nevertheless, the widespread clinical application of immunotherapy has been limited owing to the relative poor efficacy and low clinical response rates (2). Therefore, the specific biomarkers are desperately required to discriminate responders from non-responders. An increasing evidence suggests that MSI status is associated with the response to immunotherapy in advanced gastric cancer (6,7). In addition, a relative new development in the workflow of all newly diagnosed gastric cancer patients is the recommendation of MSI detection in National Comprehensive Cancer Network (NCCN) clinical guidelines (8). Meanwhile, the application of immune checkpoint blockade have been granted the US Food and Drug Administration (FDA) approval

for deficient mismatch repair (dMMR) and MSI-high (MSI-H) solid tumors (9). MSI may serve as a biomarker of immunotherapy for gastric cancer. However, the correlation between MSI and clinicopathological features in gastric cancer is still unclear.

In this review, we analyze the current evidence about MSI-H gastric cancer from a clinical perspective, focusing on the molecular and pathological features, prognostic values, and the future perspectives for immunotherapeutic applications in the MSI-H gastric cancer subgroup. We now present the subsequent article according to the Narrative Review reporting checklist (available at https://pcm. amegroups.com/article/view/10.21037/pcm-22-48/rc).

Methods

In this narrative review, we focused on MSI and immunotherapy in gastric cancer. We performed a systematic literature research on PubMed from January 11, 2017 to August 12, 2022, using the keywords: "MSI", "immunotherapy", and "gastric cancer". We selected articles published in the English language (*Table 1*).

MSI and MMR system

Those tumor individuals with MSI-H or dMMR status exhibit a favorable response to immunotherapy (10). Microsatellites are short tandem repeat DNA sequences of mononucleotide, or higher-order nucleotide repeats, which are located throughout the entire human genome (11). MSI refers to a hypermutator phenotype that happens at genomic microsatellites coupled with a dMMR machinery (12). Due to the repeated structures, microsatellites are specially vulnerable to replication errors that are usually repaired by the MMR machinery (13). MMR machinery was initially discovered as a specific somatic instability in Lynch



Figure 1 Functional schema of MMR system. MMR system ensures the genome stability through the identification of MSH2/ MSH6 complex to the genetic mismatches, and further repair of MLH1/PMS2/1 complex to the mismatched sites. Conversely, the dMMR machinery leads to the defect of repairing the DNA mismatches, leading to accumulation of abnormal mutations. IR, ionizing radiation; MSH2/6, mutS homolog 2/6; MLH1, mutL homolog 1; PMS2/1, PMS1 homolog 2/1; MMR, mismatch repair; dMMR, deficient MMR.

syndrome and was subsequently identified as a microsatellite (14,15). MMR is an extremely conserved cellular process, including a specific set of MMR genes. Under the normal DNA replication conditions, DNA mismatch sites are originally recruited and combined with mutS homolog (MSH)2/MSH6 heterodimers, in turn, mutL homolog 1 (MLH1)/PMS1 homolog 2 (PMS2) complex is responsible for the precise excision and synthesis of a corrected strand to replace the mismatched sites. Both abnormal expressions and defects in MMR elements lead to deficiency of MMR system and subsequent failed repair of mismatched DNA sites; this phenotype is known as MSI (*Figure 1*) (16).

Mutations in MMR genes result in an accumulation of DNA replication errors, which further leads to MSI (12). Indeed, the abnormal MMR genes are proved to be the transforming events to determine the tumor progression (17). Due to the large amount of microsatellites and spread over the entire genome, the MSI will lead to the dysfunction of multiple genes in numerous signaling pathways associated with tumors, further leading to the development of MSI-H tumors (18). In addition, MSI-H cancers harbor a maximum of 1,000-fold elevated mutation frequencies of missense mutations compared with MSS malignancies (19,20). Mutations occur frequently in DNA repair genes *MRE11A* and *bRAD50*, kinase genes *BRAF* and *PIK3CA*, and MMR genes *MSH3* and *MSH6*, leading to the abnormal cellular functions and signaling pathways (13).

MSI gastric cancer

Evidences in the literatures (3,21) reported the MSI-H rates in gastric cancer patients (about 5-22%) generally depended on the geographical differences, the different tumor stages and the approaches utilized to analyze the MSI status (22). MSI gastric cancer is related to an older age, tumoral location, lack of lymph node metastases and lower malignant potential (23). Additionally, MSI gastric cancers are more inclined to be diagnosed at an earlier tumor stage and categorized as the intestinal Lauren's histological type (4,23,24). Since MSI mainly occurs at the early stage of tumorigenesis, some studies have demonstrated the occurrence of MSI in most gastric precancerous lesions, demonstrating that MSI may be served as an early event during gastric tumorigenesis (25,26). A previous multinational meta-analysis showed that MSI status could be served as a robust prognostic marker in patients with

resectable primary gastric cancer (27). Compared with the MSS/MSI-low (MSI-L) subgroup, MSI-H status group demonstrated a better outcome in surgery-alone group [hazard ratio (HR), 0.35; 95% confidence interval (CI): 0.11 to 1.11; P=0.08] and a worse survival prognosis in chemotherapy + surgery treatment group (HR, 2.22; 95% CI: 1.02 to 4.85; P=0.04] in MAGIC trial (28). In addition, the methylation accumulation of MMR genes during gastric cancer progression has been also reported. Epigenetic silence of *bMLH1* caused by promoter hypermethylation accounts for primary reason of dMMR during progression of gastric cancer, however, the mutations in *bMLH1* and bMSH2 genes are less common (29-31). bMLH1 gene mutations and methylation are mainly correlated with immunohistochemistry (IHC) loss of MLH1 and PMS2 proteins. Over 50% of MSI-H gastric cancer subgroup contain hypermethylation of *bMLH1* promoter, while mutations in *bMLH1* are present in approximately 15% of MSI-H gastric cancers (17).

Studies have explored the molecular features of MSI gastric cancer, and identified a series of distinctive changed genes in the unique molecular subtype (3,4). Thirty-seven genes are significantly mutated in MSI gastric cancer according to the whole-genome analysis of TCGA dataset. These altered genes participate in a series of cellular life processes including the regulation of cell cycle, DNA integrity maintenance, chromatin remodeling, cell death, transcription regulation, apoptosis regulation and cell signal transduction. MSI gastric cancer also shows increased expressions of multiple mitotic network moleculars (3,4). KRAS mutation is significantly associated with MSI status (32-34). A recent research conducted on 595 gastric cancer patients, examining the KRAS mutation in 14.9% of MSI cases, and 1.2% of MSS. And so beyond that, patients with both KRAS mutation and MSI demonstrated a better prognosis compared with KRAS mutation and MSS patients (33,35). Furthermore, another large multicenter study was performed to detect the KRAS mutation and MMR status in locally advanced resectable gastric cancer, and the observations also confirmed the significant association between the KRAS mutation and dMMR status (36).

MSI detection

Cancers harboring a dMMR mechanism are frequently hypermutated in monomorphic microsatellites that are extremely inclined to mismatch errors. The context is defined as MSI, which can be usually detected by IHC and two other molecular examinations, including conventional MSI polymerase chain reaction (PCR) and the new next-generation sequencing (NGS) approaches (37,38).

IHC for MMR proteins is used as the first-line approach for the MSI detection due to the convenience of testing and less measurement criteria of tumor tissue compared with other molecular tests (37). MLH1, MSH6, and PMS2 antibodies for the determination of MMR proteins are commonly used, and the explanation of the findings is dependent upon the biological roles of the complex created by the detected genes (39). In fact, the alterations in the MMR genes are also in charge of protein degradation of specific complex. Mutations in MSH2 are generally correlated to the IHC deficiency in MSH2 and MSH6 proteins (37,40). Therefore, IHC detection leads to imperfect MMR genes testing and requires further detection of MMR genetic analysis.

PCR amplification with specific primers for microsatellite repeats results in a unique magnification curve. According to the length of nucleotide repeats in tumor and adjacent normal mucosa, MSI can be evaluated as a "shift" in the phenograms of one or more microsatellites (41). Usually, two mononucleotides (BAT25 and BAT26) and three dinucleotides (D5S346, D2S123, and D17S250) are routine testing sites in Bethesda panel for the MSI molecular detection (16,42). These regions are amplified using fluorescent multiplex PCR and the amplification products are further detected by the following capillary electrophoresis (43). If two or more loci (or >30% of loci) are found, the tumor is regarded as MSI-H; if only one locus (or in 10-30% of loci) is detected, the tumor is defined as MSI-L; MSS, indicating none of the markers (or <10% of loci) with instability (13,44). MSI-H and dMMR are highly concordant in many cancers, and usually these two terms can be used in place of each other (45). The variability of MSI-H frequency varies in different tumor types, among which uterine corpus cancer, colon cancer, and gastric cancer rank in the top three malignancies (46,47).

In 2014, NGS with whole genome sequencing was firstly suggested as an alternative tool for the verification of the MSI phenotypes (48,49). The superiority of NGS analysis for MSI assessment is that it is not tumortype-specific, and it does not need the matched normal specimen. Moreover, NGS-based method covers a wider range of microsatellite sites, allowing it not restricted to the conventional microsatellite loci detected by PCR/IHCbased approaches (50). However, the high expenses for NGS and the expertise required to analyze the NGS data limit the widespread application of NGS in the routine clinical diagnosis.

Immunotherapy for MSI gastric cancer

Previous clinical trials manifested that dMMR or MSI status were obviously associated with the response to the immune checkpoint inhibitors (ICIs), independent of the sites of tumor origin (47,51,52). MSI-H status has been put forward as a positive indicator of ICIs efficacy in advanced cancer patients. Evidence for the application of immunotherapy in MSI-H gastric cancer comes from the specific hypermutated phenotype in the subgroup (7). Meanwhile, MSI-H gastric cancers are able to express plentiful peptides that can trigger tumor infiltrating lymphocytes (TIL) recruitment and activation (27,53). The effector T cells in tumor microenvironment (TME) and T-cell exhaustion status are proved to be significantly associated with the response to pembrolizumab in MSI gastric cancer, and this means that both quantity and functional status of TIL in the TME are indispensable (27). In the KEYNOTE-012 trial (54), 22% of programmed death-ligand 1 (PD-L1)positive advanced gastric cancer patients obtained an overall response. Further genome analysis showed MSI-H in 17% of all enrolled patients, and half of the MSI-H patients had a partial response. In addition, MSI-H tumors demonstrated responses to ICIs independent of the PD-L1 expression (55,56). These observations from all above studies proposed evidences for potential application of MSI as a predictor for immunotherapy (57,58). In another encouraging KEYNOTE-059 clinical trial, the therapeutic effect of pembrolizumab was also evaluated in gastric/ gastroesophageal junction cancer patients. Of note, patients with MSI demonstrated an objective response rate (ORR) of 57.1%, while the patients with MSS exhibited a lower ORR (9%) (59). Among the MSI-H patients recruited in KEYNOTE-059, KEYNOTE-061, and KEYNOTE-062 clinical trials, higher response rates to pembrolizumab immunotherapy were 57.1%, 46.7%, and 57.1%, respectively. However, the median overall survival (OS) was not reached (NR) for pembrolizumab among MSI-H patients. Based on these findings, FDA approved the approval for pembrolizumab application in PD-L1 positive metastatic gastric cancer patients and unresectable dMMR/MSI solid tumor patients, independent of the primary cancer (58). In addition, the phase II KEYNOTE-158 trial also validated a significant curative effect of pembrolizumab for gastric cancer

individuals with nonresponsive in the traditional standard medications. MSI-H gastric cancer patients presented an ORR of 46% and a PFS of 11 months (60). Another clinical trial worthy of being analyzed is CHECKMATE-032 trial, which was performed to explore the efficacy and safety of nivolumab in PD-L1 unselected metastatic gastric cancer patients. The 12-month OS rates of MSI-H patients were 57% in the NIVO3 group, 50% in the NIVO1-plus-IPI3 group, and 50% in the NIVO3-plus-IPI1 group, while the OS rates of the patients with non-MSI-H patients were 33%, 32%, and 23%, respectively (61). Subgroup analysis revealed that MSI patients achieved a better median OS compared with the MSS entirety.

As to post-hoc analyses of MSI-H predictive effect in randomized controlled trials (RCTs), including the KEYNOTE-061, KEYNOTE-062, JAVELIN Gastric 100 (62), and CHECKMATE-649 phase III trials, a meta-analysis was performed and covered a total of 2,545 advanced gastric cancer patients. In total, 4.8% of the recruited participants displayed MSI-H status, and demonstrated an HR for OS benefit of 0.34 when applied with anti-programmed death 1 (PD-1) drugs versus chemotherapy. These results promote a better efficacy of pembrolizumab than the chemotherapy in support of the median OS (63).

Although these findings acquired in aforementioned studies are tremendously encouraging, due to the relatively few recruited participants and the low phenotypic frequency of MSI subtype gastric cancer, the application of ICIs in MSI patients is obviously not as good as the colorectal cancer trials findings (64). Some representative clinical trials on ICIs for MSI gastric cancer are listed below (*Table 2*). Nevertheless, immunotherapy opens up a new way of cancer treatment and improves the therapeutic effect of MSI gastric cancer, emphasizing a strong theoretical support for the management of ICIs in MSI subtype (*Figure 2*).

Conclusions

Although the efficacy of surgical treatment and targeted therapy have been greatly improved, gastric cancer remains one of the most important global disease burdens. The complexity of gastric cancer put forward higher demands for novel molecular-based individual therapeutics. The comprehensive classification of gastric cancer into four well-defined molecular subtypes, laying the foundation for propose innovative treatment strategies for the patients

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Reference	ClinicalTrials.gov number	Phase	Tumor type	Treatment settings [number of participants]	Results
Kwon e <i>t al.</i> (27)	NCT02589496	II	MSI-H gastric cancer	Pembrolizumab [61]	ORR of 55.6% and DCR of 88.9%
Muro <i>et al.</i> (54)	KEYNOTE-012 (NCT01848834)	lb	PD-L1 ⁺ advanced gastric cancer	Pembrolizumab [39]	MSI gastric cancer ORR 50%
Fuchs <i>et al.</i> (59)	KEYNOTE-059 (NCT02335411)	II	G/GEJ cancer	Pembrolizumab [259]	MSI gastric cancer ORR of 57.1%
Shitara <i>et al.</i> (65)	KEYNOTE-061 (NCT02370498)	III	G/GEJ cancer	Pembrolizumab [296]	MSI gastric cancer ORR of 46.7%
Shitara <i>et al.</i> (66)	KEYNOTE-062 (NCT02494583)	III	G/GEJ cancer	Pembrolizumab [256]; pembrolizumab plus chemotherapy [257]; chemotherapy [250]	MSI gastric cancer ORR of 57.1%
Marabelle <i>et al.</i> (60)	KEYNOTE-158 (NCT02628067)	II	Nonresponsive gastric cancer	Pembrolizumab [163]	MSI gastric cancer ORR of 45.8% PFS, 11.0 months
Janjigian <i>et al.</i> (61)	CHECKMATE-032 (NCT02267343)	1/11	PD-L1 unselected metastatic gastric cancer	Nivolumab 3 mg/kg [59]; nivolumab 1 mg/kg plus ipilimumab 3 mg/kg [49]; nivolumab 3 mg/kg plus ipilimumab 1 mg/kg [52]	MSI gastric cancer OS 15 months
Janjigian <i>et al.</i> (67)	CHECKMATE-649 (NCT02872116)	111	Gastric, gastroesophageal junction, or esophageal adenocarcinoma	Nivolumab plus chemotherapy [789]; chemotherapy [792]	Nivolumab plus chemotherapy OS (HR: 0.71) and PFS (HR: 0.68)

Table 2 Clinical trials on ICIs for MSI gastric cancer

ICIs, immune checkpoint inhibitors; MSI, microsatellite instability; MSI-H, MSI-high; ORR, objective response rate; DCR, disease control rate; PD-L1, programmed death-ligand 1; G/GEJ, gastric/gastroesophageal junction; PFS, progression-free survival; OS, overall survival; HR, hazard ratio.



Figure 2 Frameshift mutations generated by MMR system deficiency, produce neoantigens and trigger immune responses, resulting in an enhanced antitumor effect by ICIs. MMR, mismatch repair; ER, endoplasmic reticulum; PD-L1, programmed death-ligand 1; MHC, major histocompatibility complex; PD-1, programmed death 1; ICIs, immune checkpoint inhibitors.

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with specific molecular features. MSI gastric cancers take up a relatively small patients population, and show distinctive clinicopathological features. The favorable prognosis resulting from the MSI cancers treated with ICIs should be taken into account in the future clinical practice. Despite the retrospective feature of the clinical researches generally included for analysis, and the relative small number of MSI gastric cancer patients recruited in most clinical trials, the purpose of our current review is to illustrate the MSI entity as a specific subtype and these patients may be prone to immunotherapy. In conclusion, MSI detection has a promising role in guiding the immunotherapy for gastric cancer. However, they still need to be further verified in larger prospective trials for this specific clinical type.

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Footnote

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Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

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