Review

Gastric adenocarcinoma of the fundic gland: A review of clinicopathological characteristics, treatment and prognosis

rare tumors

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Abstract

Gastric adenocarcinoma of the fundic gland is a rare, well-differentiated gastric cancer entity, and very few patients transition to poorly differentiated tubular adenocarcinoma during progression. Gastric adenocarcinoma of the fundic gland originates from the mucosa of the gastric fundic gland, usually without chronic gastritis or intestinal metaplasia. Histo-logically, the tumor cells are closely arranged to form anastomosing tubular glands, and more than 95% of tumor cells differentiate towards chief cells. Most gastric adenocarcinoma of the fundic gland cases are characterized by submucosal involvement, but the tumor volume is usually small, with lymphatic and vascular invasion rarely observed. Therefore, endoscopic submucosal dissection can be an ideal treatment, leading to a favorable prognosis, and recurrence and metastasis of the disease are uncommon.

Keywords

Gastric adenocarcinoma of the fundic gland, chief cells, histopathology, treatment, prognosis

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Introduction

Morphologically, gastric cancer features a wide range of morphological heterogeneity, resulting in a variety of different histological subtypes.¹ According to the Lauren classification, gastric cancer is classified into intestinal and diffuse types²; according to the Nakamura classification, gastric cancer is grouped into differentiated and undifferentiated types.³ Based on molecular and genetic profiling, gastric cancer is divided into four genetic subtypes: EpsteinBarr virus (EBV)infected tumors, microsatellite instability (MSI) tumors, genomically stable tumors, and chromosomally unstable tumors.⁴ With the development of immunohistochemistry (IHC), it has been confirmed that intestinal gastric cancer contains a gastric phenotype. The gastric phenotypes include foveolar type, pyloric gland type, and fundic gland type.

In 2007, gastric adenocarcinoma of the fundic gland (GA-FG) was first reported as a novel tumor entity by Tsukamoto et al.⁵ According to tumor cell differentiation,

GA-FG can be divided into three subtypes: chief cellpredominant type, parietal cell-predominant type, and mixed type. The GA-FG chief cell-predominant (GA-FG-CCP) type accounts for approximately 99% of all GA-FG

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cases.⁶ With the increasing number of reported GA-FG cases, the purpose of this review is to summarize the clinicopathological characteristics, endoscopic appearance, pathogenesis, molecular phenotype, and prognosis of GA-FG to increase understanding of clinicians and pathologists regarding this type of disease.

Clinicopathological characteristics

GA-FG is a rare subtype of gastric cancer with welldifferentiated tissue and clear boundaries, accounting for approximately 1% of early gastric adenocarcinomas.⁷ Most lesions are located in the middle and upper third of the stomach.⁸ According to previous studies, the ratio of male to female patients with GA-FG is approximately 1.4⁹; however, based on our analysis, the ratio is approximately 2.2. The GA-FG tumor volume is usually small, approximately 1 cm.¹⁰ Most GA-FG patients do not have intestinal metaplasia, atrophic glands, or Helicobacter pylori (H. pylori) infection.^{8,11} However, in our study, according to the available data, 19 of 67 GA-FG patients had H. pvloripositive infection, and 11 of 48 GA-FGnegative patients experienced H. pylori eradication therapy. Therefore, we think that the H. pylori infection should be given more attention as a factor in GA-FG pathogenesis. GA-FG is a welldifferentiated and low malignant potential tumor. Lymphatic and vascular invasion rarely occurs.^{8,11,12} according to the Japanese classification of gastric carcinoma. Our current analysis shows that only 10 (6.3%) of 159 GA-FG cases with available relevant data had lymphatic invasion. and only 8 (5.0%) of 160 GA-FG cases had vascular invasion, which is similar to previous studies. Yasuhiro Okunura¹³ reported a rare case of GA-FG. This was the first report of GA-FG progressing to aggressive cancer with lymph node metastasis. More detailed clinical features are presented in Table 1.

Endoscopic imaging features

Endoscopically, most GA-FG lesions are located in the middle/upper third of the stomach. According to a previous review, the proportion of GA-FG lesions located in the middle/upper third of the stomach is up to 98% among the total GA-FG cases.⁶ In our current analysis, the proportion was approximately 97.4%, which is similar to the previous results. The number of cases located in the upper, middle, and lower third of the stomach was 156, 34, and 3, respectively. Macroscopically, according to the available data, 3 tumor lesions showed a protruded type (T0-I); 1 tumor lesion presented an ulcerative type (T2); and the number of cases of superficial and elevated (0-IIa), superficial and flat (0-IIb), superficial and depressed (0-IIc), and mixed type was 94, 14, 39, and 20, respectively. Narrow band imaging can enhance visualization of mucosal vessels and show

irregular patterns, which indicate heterogeneity of microvessels in GA-FG.^{7,14} The use of narrow band imaging and endoscopic ultrasound can improve the chance of complete resection through endoscopic mucosal resection (EMR) technology.

Histopathological features

The histopathological findings for GA-FG are shown in Figure 1. The tumor surface area is typically still covered with normal foveolar epithelium. Chronic inflammation. atrophic glands, and intestinal metaplasia are rarely observed in the surrounding mucosal background. Tumor lesions are composed of closely arranged and anastomosing glands, forming an "endless gland." Tumor cells are similar to fundic gland cells, appearing as well-differentiated, mildly heterogeneous columnar cells with moderately enlarged nuclei. In approximately 99% of cases, tumor cells differentiate into chief cells.⁶ In our study, according to the relevant data available, the number of GA-FG-CCP cases was 163, accounting for 96%. In addition, 2 cases appeared to exhibit the foveolar epithelium type, 1 case showed mixed type differentiation, and 3 cases differentiated towards the mucous neck cell type. GA-FG originates from the deep layer of the lamina propria. Although most tumor lesions show submucosal invasion, lymphatic and vascular invasion is very rare.^{7,8,15} According to our analysis, among 163 cases of GA-FG. 119 cases had submucosal infiltration (73%), and only two cases had subserosal infiltration. Although tumor submucosal involvement has been identified in most reports, thin wisps of muscularis mucosae with the absence of desmoplasia were identified throughout the le-sions. Therefore, Singhi et al.^{16,17} suggested that this change represents prolapse-type growth to the submucosa rather than the more typical submucosal infiltration seen with malignant tumors. Some researchers believe that it is excessive and exaggerated to name this disease "GA-FG," and this type of disease should be considered benign. They prefer to use "oxyntic gland polyps/adenomas" rather than "GA-FG."¹⁶ Although Yasuhiro Okumura reported a rare case of GA-FG that gradually transitioned into aggressive tubular adenocarcinoma with poorly differentiated components in the muscularis propria and subserosal layers, there were only two cases of subserosal infiltration in our analysis.^{13,18}

IHC was used to evaluate lineage differentiation in GA-FG. The main relevant cell differentiation markers are shown in Table 2. Mucin phenotypes of GA-FG were identified by detecting the expression of gastric phenotype markers, such as MUC5AC and MUC6, and intestinal phenotype markers, such as MUC2 and CD10.¹⁹ Pepsinogen-1 is the most specific marker of chief cell differentiation, and its immunohistochemical analysis will be necessary for diagnosis of GA-FG-CCP. Previously,

Author (Year)	No. of cases	Age or age range	Gender distribution (M/F)	Location of lesion (U/M/L)	H. pylori infection (+/—/post eradication)	Size of tumor or range (mm)	Macroscopic type (0-IIa/ IIb/IIc/mixed)	Invasion depth (M/SM1/ SM2/SS)	Lymphatic invasion (Present/ Absent)	Venous invasion (Present/ Absent)	Cell type
Fan ²¹	Ι	60	1/0	0/1/0	1/0/0	5	_	0/1/0/0			Chief cell
Li ⁸	8	48– 80	3/5	7/1/0	0/8/0	4–12	6/0/2/0	2/5/1/0	-	-	Chief cell
Chen	I	55	0/1	2/0/0	0/0/1	6 and 4	1/1/0/0	2/0/0/0	-	-	Chief cell
Takeshi ¹²	Ι	73	1/0	1/0/0	I/0/0	25	1/0/0/0	I/0/0/0	-	-	Foveolar epithelium
Yu ²⁷	I	77	0/1	0/0/1	0/0/1	6	1/0/0/0	0/1/0	-	-	Chief cell
Fumiaki ¹⁵	8	51– 83	4/4	6/2/0	0/4/4	4–14	6/2/0/0	5/3/0/0	-	-	Chief cell
Keita ²⁸	I	76	0/1		1/0/0	28	0/0/1/0	0/1/0/0	-	-	Chief cell
Atsushi ²⁹	Ι	70	1/0	0/1/0	_	23	—	0/1/0	-	—	Foveolar epithelium
Yasuhir ¹³	I.	55	1/0	I/0/0	_	47	Т2	0/0/0/1	+	+	Chief cell
Hitoshi ³⁰	I	78	1/0	_	0/0/1	4 and 7	T0-I	1/1/0/0	-	-	Chief cell
Shigeo ³¹	I.	58	I/0	I/0/0	0/0/1	7	1/0/0/0	0/1/0/0	-	-	Chief cell
Keitaro ³²	I.	87	I/0	0/0/1	1/0/0	3	T0-I	1/0/0/0	-	-	Mixed
Takashi ²⁶	26	49– 82	19/7	20/6/0	—	3–39	17/0/5/4	11/15/0	2/24	1/25	Chief cell
Yoriaki ³³	I	50	1/0	_	0/1/0	5	1/0/0/0	0/1/0		_	Chief cell
Gen ³⁴	4	42– 62	2/2	3/1/0	2/2/0	2–5	2/2/0/0	2/2/0	-	-	Chief cell
Keisuke ³⁵	I	62	I/0	0/1/0	0/1/0	—	0/0/1/0	0/1/0	-	+	Chief cell
Eleanor ₃₆	I	49	1/0	I/0/0	0/1/0	11	_	0/1/0	-	+	Chief cell
Takashi ²⁵	20	44– 85	16/4	14/6/0	13/5/2	—	12/5/3/0	—	—	—	Chief cell
Sato ³⁷	I.	77	0/1	I/0/0	0/1/0	10	1/0/0/0	0/1/0/0	-	-	Chief cell
Hee ³	I	69	1/0	1/0/0	—	25	_	0/1/0	-	-	Chief cell
Soichi ³⁸	I	66	1/0	1/0/0	0/1/0	—	0/1/0/0	1/0/0/0	-	-	Chief cell
Masaki ⁷	5	67– 78	3/2	5/0/0	0/5/0	5–13	4/1/0/0	0/2/3/0	1/4	-	Chief cell
Lee ³⁹	11	55– 82	8/3	8/3/0	_	3–18	T0-I:1; 5/1/4/ 0	_	-	-	Chief cell
Kato ⁴⁰	I	80	1/0	1/0/0	0/1/0	30	1/0/0/0	0/1/0/0	-	-	Chief cell
Ueyama ⁴¹	10	55– 78	6/4	6/4/0	*/6/1	3–31	*/1/3/0	5/4/1/0	1/9	-	Chief cell
Nomura ¹⁰	26	49– 79	22/4	23/3/0	_	3–85	15/0/5/6	0/20/6/0	3/23	2/24	—
Ueo ¹⁸	I	62	1/0	0/1/0	0/1/0	44		0/0/0/1	+	+	Chief cell
Ryoji ⁴²	3	56– 78	1/2	3/0/0	_	3–8	_	1/1/1/0	—	—	Chief cell
Hidaka ¹⁷	31	41– 79	21/10	28/3/0	_	3–85	14/0/10/7	0/31/0	1/30	1/30	Chief cell
Su park ⁴³	3	47– 76	3/0	1/1/1	—	12–36	0/0/0/3	1/1/1/0	-	-	Chief cell
Singhi ¹⁶	10	44– 79	4/6	10/0/0	_	20–80	_	10/0/0/0	-	-	Chief cell: 7; Mucous neck cell: 3
Fukatsu ⁴⁴	I.	55	1/0	1/0/0	_	5	_	0/1/0/0	-	-	Chief cell
Terada ²²	I	78	1/0	1/0/0	_	15	1/0/0/0		_		Chief cell

Table I. Characteristics of reported gastric adenocarcinoma of the fundic gland.

(continued)

Table I. (continued)

Author (Year)	No. of cases	Age or age range	Gender distribution (M/F)	Location of lesion (U/M/L)	H. pylori infection (+/—/post eradication)	Size of tumor or range (mm)	Macroscopic type (0-IIa/ IIb/IIc/mixed)	Invasion depth (M/SM1/ SM2/SS)	Lymphatic invasion (Present/ Absent)	Venous invasion (Present/ Absent)	Cell type
Ueyama ²⁰	10	42– 79	6/4	10/0/0		4–20	5/0/5/0	1/6/3/0	-	-	Chief cell
Total	195	41– 87	134/61	156/34/3	19/37/11	2–85	T0-l: 3; T2: 1; 94/14/39/ 20	44/119/2	10/159	8/160	Chief cell: 163; Foveolar epithelium: 2; Mixed: 1; Mucous neck cell: 3

---: Relevant data is not available; M: male; F: female; U: upper third of stomach; M: middle third of stomach; L: lower third of stomach; H. *pylori: Helicobacter pylori*; Type 0-I: protruded type; 0-Ila: superficial and elevated type; 0-Ilb: superficial and flat type; 0-Ilc: superficial and depressed type; T2: ulcerative; M: tumor confined to the mucosa; SMI: tumor confined to the submucosa, but depth is within 0.5 mm of the muscularis mucosae; SM2: tumor invasion is 0.5 mm or more deep into the muscularis mucosae; SS: tumor invades the subserosa; +: positive; -: negative; *: partial data is unknown.

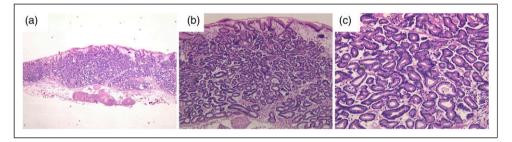


Figure I. Histopathological findings of representative GA-FG case. (A) In low-power view (×40), the tumor originated from the deep layer of the lamina propria and invaded submucosal layer. The surface was covered with non-atypical foveolar epithelium; (B) and (C): In high-power view (×100 and 200), the tumor comprised well-differentiated columnar cells mimicking the fundic gland cells with mild nuclear atypia.

Ueyama et al.²⁰ reported that in the tumor progression process, cell differentiation can change from the fundic gland type to the foveolar type in advanced GA-FG patients. GA-FG differentiation in several directions has more malignant potential than GA-FG differentiation into chief cells.¹²

Regarding expression of p53 protein and the Ki67 labeling index, none of the reported cases showed an overexpression of p53, and the Ki67 labeling index was very low (less than 5%).^{21,22}

Pathogenesis and molecular analysis

Although many studies have been performed to elucidate the pathogenesis of GA-FG, it is still unclear. *H. pylori* is a gram-negative, flagellated, acid-resistant bacterium that infects more than 50% of the world's population.²³ In etiology, *H. pylori* infection is associated with nonatrophic or atrophic gastritis, intestinal metaplasia,^{15,24} and peptic Table 2. Cell differentiation markers.

Marker	Cell type targeted
MUC2	Goblet cell
MUC5AC	Gastric foveolar epithelium
MUC6	Mucous neck cell and pyloric gland
CD10	Brush border
Pepsinogen-1	Chief cell
H+/K+-ATPase	Parietal cell

ulcers and is an increased risk factor for developing gastric adenocarcinoma and mucosa-associated lymphoid tissue lymphoma.²³

A recent study reported the effect of *H. pylori* infection in GA-FG. Compared with the *H. pylori*-negative group, a decreasing number of fundic glands and thinned foveolar epithelium covering tumor ducts were observed in the *H. pylori* eradication group.¹⁵ Takashi Chiba et al.²⁵ reported 20

cases of GA-FG, of which 15 were *H. pylori* positive or experienced *H. pylori* eradication, but almost all lesions occurred in nonatrophic gastric mucosa. Therefore, the association between *H. pylori* infection and GA-FG still needs to be further studied.

Some GA-FG cases showed distended fundic glands mixed with foveolar cells, similar to fundic gland polyps. Therefore, some hypotheses have been raised regarding whether the occurrence of GA-FG is related to the use of proton pump inhibitors. However, the associated research is limited, and the relationship between the occurrence of GA-FG and the use of proton pump inhibitors is not clear.

There are few studies on the molecular changes associated with GA-FG. Activation of the Wnt/β-catenin signaling pathway is considered to be related to GA-FG occurrence, but the detailed mechanism underlying the activation of this signaling pathway has not been elucidated. To elucidate the potential role of gene methylation in relation to the Wnt/β-catenin signaling pathway, Takashi Murakami et al.²⁶ performed β -catenin immunostaining and methylation-specific PCR detection of related genes, including SFRPs, APC, AXIN2, and MCC, in GA-FG-CCP. They found that the nuclear β -catenin labeling index was higher in GA-FG-CCP. SFRPs, APC, and AXIN2 are also more frequently methylated in GA-FG-CCP. A significant correlation was observed between nuclear β-catenin expression and SFRP1 methylation. In conclusion, the activation of this signaling pathway mediated by gene methylation may be related to the occurrence and development of GA-FG-CCP. In 2014, Ryosuke Nomura et al.¹⁰ detected the mutation status of GNAS and the Wnt/β-catenin signaling pathwayrelated genes CTNNB1/AXINS/APC in 26 cases of GA-FG. They found that nearly half of the cases harbored at least one mutation in CTNNB1/AXINS/APC, leading to persistent activation of the Wnt/B-catenin signaling pathway. Among 26 cases of GA-FG, only 5 cases contained GNAS mutations (R201 C). GNAS mutation may occur in a small proportion of GA-FG as an alternative mechanism to activate the Wnt/ β -catenin signaling pathway. Similarly, in 2013, Yasuhiro Hidaka et al.¹⁷ detected the mutation status of the CTNNB1/AXIN1/AXIN2/APC genes. The results showed that 14 (51.9%) of 27 GA-FG-CCP type lesions contained at least one gene mutation in CTNNB1/ AXIN1/AXIN2/APC. Among 27 cases of GA-FG, 17 cases presented low nuclear β -catenin expression, and 7 cases had high nuclear β -catenin expression. Therefore, as a unique nonaggressive tumor entity, GA-FG-CCP is characterized by accumulation of nuclear β -catenin and mutation of CTNNB1 or AXIN genes, indicating activation of the Wnt/ β -catenin pathway. However, in a recent paper on 8 cases of GA-FG,⁸ tumor nuclear β -catenin expression was negative in all cases, which may be attributed to the limited number of cases.

Treatment and prognosis

GA-FG is a well-differentiated, well-circumscribed, and low malignant potential gastric cancer entity that rarely exhibits lymphatic or vascular invasion. However, it is commonly observed in the submucosa. Endoscopic submucosal dissection (ESD) is considered the most effective regimen for restricted GA-FG. The present evidence shows that GA-FG has a favorable prognosis, and metastasis and recurrence rarely occur during follow-up. Li et al.⁸ performed a follow-up of 8 cases of GA-FG from 5 to 33 months. None of the patients had recurrence or metastasis. According to these characteristics, GA-FG confined to the submucosa can be treated by ESD. However, extended gastrectomy is needed for GA-FG that has transitioned into a more malignant tumor, disease with suspected lymph node metastasis, or multiple gastric cancer.

Conclusion

GA-FG is a well-differentiated, low malignant potential, gastric cancer entity originating from the gastric fundic gland. In the surrounding mucosal background, chronic gastritis, atrophic glands, and intestinal metaplasia cannot be observed. In more than 95% of GA-FG cases, the tumor cells differentiate towards chief cells. Tumor cells with mild atypia and enlarged nuclei are closely arranged to form anastomosing glands, the so-called "endless gland." Tumor lesions are very prone to involve the submucosa, despite their low malignancy and invasiveness. This phenomenon may not be malignant invasion of the tumor but instead may be a prolapse-type growth pattern into the submucosa. Although abnormal activation of the Wnt/β-catenin signaling pathway is related to GA-FG, there are few studies on the molecular changes in GA-FG and its pathogenesis, and further studies on GA-FG pathogenesis are necessary.

Author contributions

Guang Yang, Ru-yi Zheng, Xiang-yu Meng, and Cheng-ji Dong developed the main content of this manuscript. Guang Yang and Ru-yi Zheng were involved in writing the manuscript. Cheng-ji Dong and Xiang-yu Meng edited the language of revised manuscript. Guang Yang supervised the project and contributed to the revision of the final manuscript.

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Ethical approval

An approval of our Institutional Review Board or Ethics Committee was not needed to conduct this analysis.

Informed consent

Not applicable.

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