

[CASE REPORT]

Attenuated Familial Adenomatous Polyposis

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Abstract:

A 36-year-old man was diagnosed with multiple gastric polyps by esophagogastroduodenoscopy. Subsequent colonoscopy identified two tubular adenomas, and computed tomography revealed subcutaneous tumors. Based on these findings, we suspected that gastric polyposis was associated with the *APC* gene, either attenuated familial adenomatous polyposis (AFAP) or gastric adenocarcinoma and proximal polyposis of the stomach (GAPPS). A genetic analysis demonstrated that he had a frameshift variant at codon 1928 of *APC*, suggesting AFAP. In this era of less *Helicobacter pylori* infection and frequent use of proton pump inhibitors, diagnoses of AFAP and GAPPS should be considered in patients with prominent gastric fundic gland polyposis.

Key words: attenuated familial adenomatous polyposis, gastric adenocarcinoma and proximal polyposis of the stomach, desmoid tumor, *APC*-associated polyposis

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Introduction

Fundic gland polyps are benign gastric epithelial lesions consisting of the hyperplastic expansion of the deep epithelial compartment of the mucosa (1). They are often detected at esophagogastroduodenoscopy (EGD) and regarded as characteristic of *Helicobacter pylori* uninfected gastric mucosa (2-4). Sporadic fundic gland polyposis may occur in *H. pylori* uninfected gastric mucosa and/or individuals with long-term use of proton pump inhibitors (PPI) (5).

Fundic gland polyposis occurs as a counterpart to *APC*associated gastric polyposis. The underlying conditions of *APC*-associated gastric polyposis include classical familial adenomatous polyposis (FAP), attenuated FAP (AFAP) and gastric adenocarcinoma and proximal polyposis of the stomach (GAPPS). Genotype-phenotype correlations between *APC*-associated gastric fundic gland polyposis and the *APC* gene have been reported (6-10).

We herein report a case of AFAP, in which the *APC* gene was diagnostic of the disease.

Case Report

A 36-year-old man was referred to our hospital for a detailed examination of multiple gastric polyps. He had a history of a back desmoid tumor at three years old, and his family history was remarkable for gastric cancer in his father but negative for colorectal cancer. There was no family history of cancer or gastrointestinal polyposis. A physical examination and laboratory data showed no abnormalities.

EGD showed multiple polyps in the fundus and body in non-atrophied background mucosa. The polyps were characterized by a sessile or semi-pedunculated configuration and were the same color as the background mucosa (Fig. 1). Biopsy specimens obtained from the polyps revealed dilated fundic glands with no cellular atypia and a partial increase in the density of the glands (Fig. 2). No abnormalities were found in the esophagus and duodenum. Two small polyps were found in the transverse colon during subsequent colonoscopy. Both polyps were removed by cold snare polypectomy and diagnosed as tubular adenomas (Fig. 3). Capsule endoscopy showed no abnormalities in the small in-

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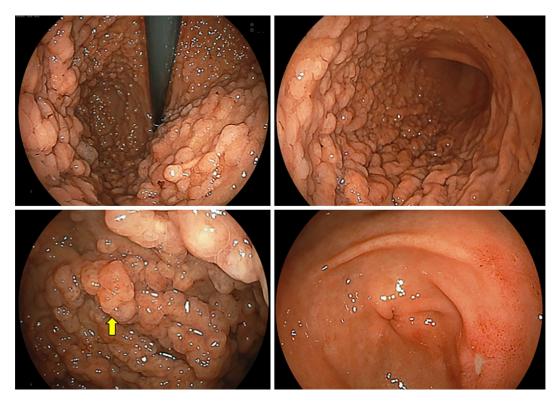


Figure 1. EGD identified no atrophy of the gastric mucosa but detected multiple polyps in the fundus and body that were the same color as the background mucosa, some of which were slightly reddish (yellow arrow).

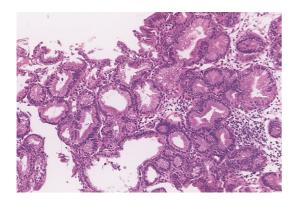


Figure 2. A biopsy of the reddish lesion revealed dilated fundic glands with no cellular atypia but a slight increase in the density of the glands.

testine. Contrast-enhanced computed tomography (CT) of the neck, chest, abdomen, and pelvis showed scattered elevated soft tissue density in the patient's back, suggestive of desmoid tumors (Fig. 4). There were no signs of tumors in the thyroid, pancreas, liver, or adrenal glands.

Based on the presence of multiple fundic gland polyps, desmoid tumors, and a small number of colonic adenomas, we considered a diagnosis of *APC*-associated gastric polyposis, specifically AFAP and GAPPS, as differential diagnoses. We therefore screened the patient's *APC* gene after obtaining his written informed consent. As a result, a frameshift variant of codon 1928 (c.5282del; Gln>Arg) was found. Based on this result, we diagnosed the patient with

AFAP. He has been under observation with annual EGD, colonoscopy (CS) and CT.

Discussion

To date, three categories of *APC*-associated fundic gland polyposis have been reported: classical FAP, AFAP and GAPPS (6). Spirio et al. reported AFAP as a family with few colorectal polyps and an older age at the onset of colon cancer than classical FAP, despite germline variants of *APC* (11). The reason the phenotype of AFAP is milder than that of FAP is not fully understood. It has been hypothesized that, in AFAP, mutant APC proteins with a mutation on the 5' side of the *APC* gene can degrade β -catenin. In contrast, the variant APC proteins with 3' variants found in AFAP families are partially functional, so the accumulation of intracellular β -catenin may be milder than in FAP. It has also been presumed that those 3' and 5' variants of *APC* are associated with a less dominant negative effect, which may be responsible for the phenotypic differences (12).

Burt et al. reported that the median number of colorectal polyps in AFAP was 25, with a range of 0 to 470 (13). However, the number of colorectal polyps in patients with FAP varies greatly with the colonoscopic procedure. It therefore seems possible that the actual number of colorectal polyps in patients diagnosed with AFAP may be greater than reported. However, as found in our patient, some cases of AFAP have an extraordinarily small number of colorectal polyps and desmoid tumors, most of which had 3' variants

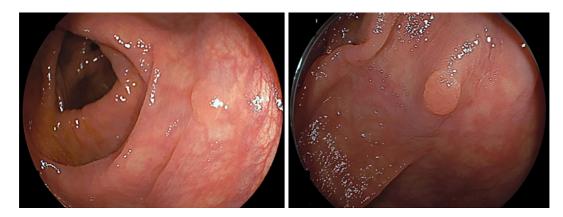


Figure 3. Colonoscopy identified two small polyps in the transverse colon. Cold snare polypectomy was performed, and both lesions were identified histologically as tubular adenomas.

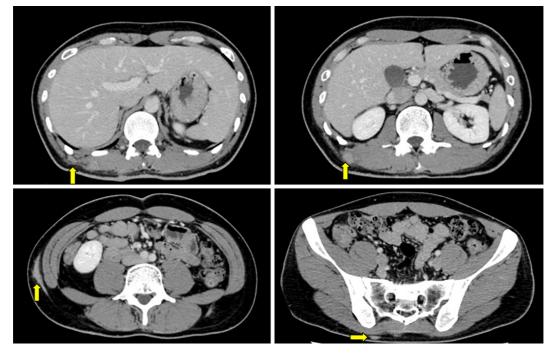


Figure 4. Contrast-enhanced computed tomography (CT) showed scattered, elevated masses with soft tissue density in the patient's back, suggestive of desmoid tumors (yellow arrows).

of *APC* located distal to codon 1595 (14-16). In fact, our patient carried a variant at codon 1928. Prominent gastric fundic gland polyposis with very few colorectal adenomas may therefore be a specific clinical manifestation of AFAP due to 3' variants of *APC*.

GAPPS is another condition that should be distinguished from AFAP. It is an autosomal dominant genetic disorder with variants in promoter 1B of the *APC* gene that cause gastric cancer to arise from fundic gland polyposis (10, 17, 18). The diagnostic criteria for GAPPS are dense gastric fundic gland polyposis without polyposis in the duodenum and colon, more than 30 gastric polyps in first-degree relatives, and pathologically verified dysplasia in fundic gland polyposis (9). In addition, Worthley et al. and McDuffie et al. reported that patients genetically diagnosed with GAPPS had 0-16 colorectal adenomas, suggesting difficulty distinguishing AFAP from GAPPS by clinical manifestations alone (10, 19).

Neither the Japanese nor American guidelines (20, 21) include any specific comments on *APC* gene testing for the differentiation of AFAP and GAPPS. However, we believe that AFAP and GAPPS should be differentiated genetically because of differences in the risk of gastric cancer (21, 22). Because GAPPS is characterized by variants in the promoter region of *APC*, routine gene testing may fail to identify variants in *APC* in patients with the disease. Patients with profuse gastric polyposis without apparent *APC* mutations are therefore appropriate candidates for the investigation of the *APC* promoter region.

Conclusion

Our experience suggests that, in addition to FAP, AFAP

should be considered seriously in patients with profuse gastric polyposis. Clinical guidelines for the clear distinction of AFAP from GAPPS must be established, because both conditions are characterized by no or very few colorectal adenomas.

As this paper is a single case report, it has not been approved by the Ethics Committee. Written informed consent was obtained from the patient for the publication of this case report and any accompanying images.

The authors state that they have no Conflict of Interest (COI).

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