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# Abstract

**Background** Patients with familial adenomatous polyposis (FAP) have a lifetime risk of developing duodenal adenomas approaching 100%, and the relative risk for duodenal cancer compared with the general population is high. We conducted a retrospective study to investigate the progression of non-ampullary duodenal adenomas (NADAs) and risk factors for advanced lesions in patients with FAP.

**Methods** Of 248 patients with 139 pedigrees at 2 institutes, we assessed 151 patients with 100 pedigrees with a pathogenic germline variant in the *adenomatous polyposis coli* gene, excluding mosaic variants. We evaluated the prevalence of NADAs in patients with FAP, the progression of these adenomas to advanced adenoma during the observation period, and the risk factors for the lifetime development of high-grade dysplasia (HGD), large ( $\geq$  10 mm) duodenal adenomas, and Spiegelman stage IV.

**Results** During the median observation period of 7 years, the incidences of patients with NADAs, with more than 20 polyps, with polyps ≥ 10 mm, with HGD, and with stage IV at the last esophagogastroduodenoscopy were increased 1.6-fold, 1.7-fold, 5-fold, 22-fold, and 9-fold, respectively. Intramucosal cancer occurred in three patients (2%), but no patients developed invasive cancer during the observation period because we performed endoscopic intervention for advanced adenomas. Stage progression was observed in 71% of 113 patients. Stage IV was more common in women, patients with a history of colectomy, and those with a 3' side mutation in their *adenomatous polyposis coli* gene.

**Conclusions** NADAs in patients with FAP frequently become exacerbated. Our findings suggest that patients with FAP who develop duodenal adenomas should be surveyed to prevent the development of duodenal cancer.

**Keywords** Familial adenomatous polyposis, Non-ampullary duodenal adenoma, Advanced adenoma, Endoscopic surveillance, Progression

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## Background

Familial adenomatous polyposis (FAP) is an autosomal dominant disorder caused by germline mutations in the *adenomatous polyposis coli* (*APC*) gene. It is characterized by multiple adenomas throughout the colon and rectum and results in colorectal cancer (CRC) in almost 100% of patients if left untreated.

FAP is associated with multiple lesions other than those in the large intestine, and the lifetime risk of developing duodenal adenomas approaches 100% [1–6]. Unlike adenomas in the large intestine, duodenal adenomas do not necessarily become cancerous. Nevertheless, the risk of duodenal cancer in patients with FAP is 250.0 to 330.8-fold higher than that of the general population [7, 8]. Therefore, patients with FAP constitute a high-risk group for duodenal cancer. The lifetime risk of developing duodenal cancer ranges from 3 to 5% [9–11] and is higher in other reports [1, 4, 12, 13]. Duodenal cancer is also the third leading cause of death in patients with FAP, accounting for the deaths of approximately 3% of these patients [14, 15]. Although half of duodenal cancer occur in the ampulla and periampullary region, we should consider ampullary and non-ampullary duodenal adenomas (NADAs) separately because ampullary adenoma rarely progresses, and when it does, it is slow [16]. Although a therapeutic strategy for ampullary adenoma has almost been established, no strategy for NADAs has been established.

CRC that occurs in patients with FAP can be managed by prophylactic total colectomy or intensive endoscopic intervention [17], leading to reduced mortality; thus, the prognosis of FAP has dramatically improved. With respect to duodenal cancer, the severity of NADAs is classified by Spigelman score. The risk of developing duodenal cancer is stratified according to its stage [18]. Because polyps larger than 10 mm and with highgrade dysplasia (HGD) are generally called "advanced adenoma" and are considered a precursor of cancer, we performed endoscopic interventions for these advanced adenomas. However, the extent to which NADAs worsen to advanced adenoma is unclear. Therefore, we proposed intensive downstaging polypectomy for all detected NADAs in our prospective study [19]. By examining the changes that occur in NADAs over time, we may more effectively stratify the risk of developing advanced duodenal adenomas. Therefore, we conducted a retrospective study to investigate the progression of NADAs and risk factors for advanced lesions in patients with FAP.

# Methods

## Patients

Patients with FAP who visited the Osaka International Cancer Institute and Ishikawa Gastrointestinal Clinic from January 1998 to September 2018 were retrospectively assessed for enrollment in this study. The diagnostic criteria for FAP in this study were (i) the presence of either  $\geq$ 100 adenomas in the colon and rectum or 10 to 99 adenomas with a family history of FAP and (ii) a pathogenic germline variant in APC. Patients with mosaic variants in APC were excluded. Esophagogastroduodenoscopy (EGD) was generally recommended for all patients with FAP. Patients who had not undergone EGD at our institutions and patients with a history of duodenal surgery, such as gastric surgery or pancreatoduodenectomy, were not enrolled in this retrospective study because we could not ascertain the progression of the duodenal neoplasms in such patients. Most of the cases in this study were also analyzed in the study by Shimamoto et al. (Genotype-phenotype correlation for life-threatening complications in patients with familial adenomatous polyposis, accepted in Cancer Science).

# Extraction of clinical data and definition of measured variables

We collected clinical data from the patients' medical and endoscopic records. The genetic information used in this study comprised *APC* pathogenic variant and pedigree data aggregated at the Medical Research Support Co., Ltd. (Osaka, Japan), a data center operated by academic doctors. The genetic information was aggregated and controlled at the center in accordance with strict guidelines. All patients received genetic counseling and provided informed consent for *APC* genetic testing.

The observation period was defined as the duration from the day on which the first EGD was performed to the day on which the last EGD was performed according to the patient's medical record or until immediately before the treatment intervention, if any. We generally performed annual surveillance EGD and targeted biopsies of polyps. We considered treatment interventions for patients with an advanced duodenal adenoma, namely>10 mm or HGD, defined as lesions with severe atypia according to the previous criteria, because advanced adenoma is generally considered a precursor of invasive cancer in colorectal polyp management, and we needed to prevent the development of duodenal cancer to secure the patients' safety.

Based on the number of colorectal adenomas, severe FAP was defined as the inability to visualize a patient's normal mucosa macroscopically because of the profusion of colorectal adenomas, and sparse FAP was considered typical FAP (which is not severe FAP). Attenuated FAP was defined as the presence of 10 to 99 colorectal adenomas. Patients were considered to have *Helicobacter pylori* infection when a positive result was obtained by a serological test, urea breath test, rapid urease test, or histological examination, and when they had a history of eradication therapy for *H. pylori*.

## Endoscopic system and settings

The endoscopic system consisted of a video processor (CV-260, CV-260SL, or CV-290; Olympus Co., Tokyo, Japan and VP-4450HD or VP-7000; Fujifilm Co., Tokyo, Japan) and a light source (CLV-260, CLV-260SL, or CLV-290; Olympus Co. and LL-4450, XL-4450, LL-7000, or BL-7000; Fujifilm Co.). A videoendoscope (GIF-H260Z, Q240Z, H290Z, PCF-Q260JI, or PCF-H290TL/I; Olympus Co. and EG-L590ZW, EG-L600ZW, or EGL600ZW7; Fujifilm Co.) was also used. Observations were performed by white light imaging and non-magnifying narrow-band imaging and/or with chromoendoscopy using indigo carmine with or without magnification.

## **Histological examination**

Lesions thought to be neoplastic or cancerous (welldemarcated lesions, large lesions, lesions with a depression at the center, and lesions with an irregular shape) were selected for biopsy. Biopsy tissue samples or endoscopic resection samples were collected and diagnosed. In patients who underwent intervention for duodenal neoplasms without prior biopsy, we collected information obtained by the following intervention. According to the Japanese classification of colorectal carcinoma, a histopathological examination was performed by two pathologists with expertise in gastrointestinal pathology. According to the Japanese classification, noninvasive cancer was evaluated as severe atypia by the Spigelman classification, which corresponds to Western standards. Because the grading of dysplasia according to the Vienna classification changed from mild/moderate/severe to low-grade/high-grade in 2000, our patients straddled the two eras. Our pathologists followed the mild/moderate/severe grading classification for a while, and then we adopted the original Spigelman stage (SS), using mild/ moderate/severe in the analysis. We thus refer to severe atypia as "HGD" in this study. The most severe histopathological diagnosis during the observation period was adopted as the final histopathological diagnosis.

#### Spigelman classification

When NADAs were found, they were scored using the Spigelman classification to indicate the severity of duodenal polyposis [18]. The Spigelman classification includes the number of polyps, maximum diameter, tissue structure, and degree of atypia. Stages 0 to IV were determined by the total score obtained using the abovementioned criteria. When a detected duodenal polyp was not histopathologically diagnosed as an adenoma, it was not considered to be a NADA.

## Statistical analysis

We evaluated the prevalence of NADAs in patients with FAP, the progression of these adenomas during the

observation period, the risk factors for the lifetime development of advanced adenoma, and SS IV until the end of the observation period.

The prevalence of NADAs in patients with FAP was indicated by the number and percentage of patients with NADAs. The progression of NADAs during the observation period was assessed with the SS, which includes the number of adenomas, size of adenomas, and development of severe dysplasia. Finally, we evaluated the lifetime risk of developing an advanced duodenal adenoma, and SS IV, which were reported to be risk factors for developing duodenal cancer [18, 20].

For the analysis of SS progression, patients for whom the observation period was <5 years without SS progression were excluded because the observation period was considered too short. We then examined the progression of patients with SS 0 to III at the first EGD because there was no room for progression for patients with stage IV at the first EGD. We also examined the progression of patients with SS 0 and I at the first EGD, who were recommended to undergo follow-up at 5-year intervals according to the guidelines. Furthermore, we examined the occurrence of HGD according to the SS at the first EGD.

HGD, a tumor size  $\geq 10$  mm, the surveillance period, a history of colon cancer, age, the SS at the first EGD, classic FAP, and the location of several mutations in APC were evaluated as risk factors for disease progression because they were relevant to the severity of duodenal polyposis and reported to be risk factors for duodenal cancer [2-6, 21-23]. We also investigated whether APC mutations at codons 1250 to 1464, which are common in patients with severe FAP, were associated with the incidence and severity of NADAs [23]. Because we used only the protein truncation test for mutation detection in APC in patients whose first endoscopy procedure was performed before 2009, we could not obtain detailed information on the gene mutation site in some cases. We excluded those cases from the analysis of risk factors for disease progression.

Statistical analyses were performed using R software version 3.3.0 (R Foundation for Statistical Computing, Vienna, Austria; http://cran.r-project.org/). The data were analyzed using the  $\chi^2$  test and Kruskal–Wallis test. Statistical significance was set at P<0.05.

#### Results

#### Patient characteristics at first endoscopy

Among 248 patients from 139 pedigrees diagnosed with FAP from January 1998 to September 2019 at our institutes, 50 patients who had not undergone EGD at our institutes and 4 patients with a history of duodenal surgery were excluded. *APC* genetic testing was performed for 90% of the remaining 194 patients from 110



Fig. 1 Flowchart of patient selection. EGD, esophagogastroduodenoscopy; APC, adenomatous polyposis coli gene

Characteristics	n = 151
Age at first endoscopy, years	30 (16–76)
Sex, male	65 (43)
Body mass index, kg/m <sup>2</sup>	21.0 (14.7–37.2)
Alcohol intake	
Past/current drinker	74 (49)
Never-drinker	77 (51)
Smoking	
Past/current smoker	36 (24)
Never-smoker	115 (76)
Family history of FAP	
Positive	130 (86)
Negative	8 (5)
Unknown	13 (9)
Helicobacter pylori status	
Positive	29 (19)
Negative	115 (76)
Unknown	7 (5)
Age at diagnosis of FAP, years	26 (13–64)
History	
Colorectal cancer	11 (7)
Colorectal surgery	49 (32)
FAP classification	
Classic FAP	138 (91)
Attenuated FAP	13 (9)
Ampullary adenoma	
Positive	22 (15)
Negative	130 (86)
Data are presented as median (range) or n (%).	
FAP, familial adenomatous polyposis	

Table 1	Patients	baseline	characteristics	

pedigrees, among whom 151 patients from 100 pedigrees were mutation-positive (Fig. 1). Table 1 shows the patients' background characteristics. Female patients were predominant (57%), most patients had a family history of FAP, one-third had a history of colectomy, and a few (7%) had a history of CRC. An ampullary adenoma was found in 22 patients (15%), including 1 patient with a history of papillectomy.

# Progression and risk factors for the progression of duodenal adenoma

After excluding 5 patients who underwent only one EGD procedure, we evaluated the remaining 146 patients who underwent two or more EGD procedures. EGD was performed approximately once per year. The median age (range) at the start and end of the observation period was 30 (16–76) and 39 (18–80) years, respectively. The median observation period (range) was 7 (0–19) years and the median number of EGD procedures (range) during the observation period was 7 (2–26).

The prevalence of NADAs at the start of the observation period was 52% (79/151), and histologically, all were tubular adenomas. The prevalence of NADAs at the end of the observation period was 85% (124/146) and this increased 1.6-fold during the observation period. The incidence of patients with more than 20 polyps increased 1.7-fold, with polyps≥10 mm increased 5-fold, with HGD increased 22-fold, and with stage IV at the last EGD increased 9-fold (Table 2; Fig. 2). 30% (21/70) of patients who did not have adenomas at the first EGD had developed at least one NADA by the last EGD. Intramucosal

 Table 2
 Duodenal adenoma distribution according to

Spigelman score at the first and last endoscopies

Spigelman score	First	Last	P-
	endoscopy	endoscopy	value
	(n=151)	(n=146)	
	n (%)	n (%)	
Spigelman score			
Number of polyps			
0 points	72 (48)	22 (15)	< 0.001
1 point: 1–4	19 (13)	28 (19)	
2 points: 5–20	18 (12)	26 (18)	
3 points: >20	42 (28)	70 (48)	
Polyp size			
0 points	72 (48)	22 (15)	< 0.001
1 point: 1–4 mm	29 (19)	42 (29)	
2 points: 5–10 mm	43 (28)	45 (31)	
3 points: >10 mm	7 (5)	37 (25)	
Histology			
0 points	72 (48)	22 (15)	< 0.001
1 point: tubular	79 (52)	124 (85)	
2 points: tubulovillous	0 (0)	0 (0)	
3 points: villous	0 (0)	0 (0)	
Dysplasia			
0 points	72 (48)	22 (15)	< 0.001
1 point: mild	47 (31)	42 (29)	
2 points: moderate	30 (20)	50 (34)	
3 points: severe	2 (1)	32 (22)	
Spigelman stage			
0	72 (48)	22 (15)	< 0.001
l: 1–4 points	11 (7)	4 (3)	
ll: 5–6 points	22 (15)	40 (27)	
III: 7–8 points	43 (28)	54 (37)	
IV: 9–12 points	3 (2)	26 (18)	

cancer occurred in three patients (2%), but no invasive cancer developed during the observation period because we performed endoscopic interventions for advanced adenomas before developing invasive cancer.

Stage progression was observed in 71% of 113 patients with stage 0 to III at the first EGD during the median observation period (2–19) of 9 years, excluding 4 patients with stage IV at the beginning and 29 patients with an observation period of  $\leq$ 5 years without stage progression (Table 3). In addition, stage progression was recognized in 83% of 72 patients with stage 0 to I at the first EGD during the median observation period (0–19) of 10 years, excluding 9 patients with a short observation period of  $\leq$ 5 years (Table 4). Stage progression among patients with stage 0 to III and stage 0 to I was observed significantly more often in those with classic FAP and a history of colectomy. Among patients with stage 0 to I, those with progression during the observation period were younger at the time of FAP diagnosis (Tables 3 and 4).

HGD developed at the last EGD in 17.1% (12/70) of patients with stage 0 at the first EGD, 9% (1/11) of those with stage I, 25% (5/20) of those with stage II, 29% (12/41)

of those with stage III, and 50% (2/4) of those with stage IV (Fig. 3).

# Risk factors for the lifetime development of each event until the end of the observation period

A duodenal polyp $\geq$ 10 mm developed in 25% of patients and was significantly more common in patients with a history of CRC, a history of colectomy, and a 3' side mutation (Table 5). HGD occurred in 22% of patients and was significantly more common in patients with a history of colectomy (Table 6). Progression to stage IV was observed in 18% (27/146) of patients. One patient exhibited improvement from stage IV to stage III. Stage IV was significantly more common in women and patients with a history of colectomy, and it was marginally more common in those with a 3' side mutation (Table 7).

## Discussion

In this study, we observed the progression of NADAs to advanced adenoma in patients with FAP. NADAs frequently become exacerbated, which may explain the need for regular surveillance and early intervention to prevent major morbidity. Studies have examined the timing and strategies of endoscopic treatment for NADAs in patients with FAP [9–11]; however, a sufficient consensus has not been achieved, and optimal management criteria have not been established. Our results provide important information regarding the management of NADAs in patients with FAP.

The reported prevalence of NADAs in patients with FAP ranges from 30 to 90% but increases after the age of 40 years [24, 25], and the SS also worsens with age [10, 12, 26, 27]. In the present study, the prevalence of NADAs at the start of the observation period was 52%. However, the numbers of patients with NADAs, more than 20 polyps, polyps>10 mm, HGD, and stage IV at the final EGD were substantially increased during the median 7-year observation period. Although 15% of patients did not develop NADAs during the observation period, the risk of developing NADAs increased over time, as previously reported, and even patients with early-stage NADAs showed frequent progression. Stage progression was observed in 71% of patients, and the same tendency was observed in patients with stage 0 to I at the first EGD. Because the incidence of HGD at the last EGD was higher among patients with a more advanced stage at the first EGD, it is desirable to prevent stage progression. Furthermore, the incidence of HGD in patients with stage 0 at the first EGD was 17% at the last EGD. Therefore, it might be better to consider surveillance EGD for all patients with FAP.

No patients in the present study developed invasive cancer during the study period because we recommended treatment intervention before the development



Fig. 2 Distribution of duodenal adenomas according to Spigelman score at the first and last EGD procedures. (a) Number of adenomas. (b) Size of adenomas. (c) Dysplasia. (d) Stage. EGD: esophagogastroduodenoscopy

of invasive cancer. Groves et al. [2] reported that SS IV was a risk factor for developing duodenal cancer at the time of the first endoscopy, and was a risk factor in patients who developed duodenal cancer. In other studies, 63% of patients with stage IV disease at the time of the first EGD developed HGD [10, 28], and  $a \ge 10$ -mm adenoma was shown to be a risk factor for HGD [6]. Therefore, advanced duodenal adenomas and SS IV have been cited as the main risk factors for the development of duodenal cancer, and can be considered surrogate measurement indices for the development of invasive cancer in this study. Although the usefulness of SS is controversial, there is currently no alternative measurement [20]. In addition, Thiruvengadam et al. reported that polyp size and HGD were associated with cancer development [20]. Therefore, our assessment using advanced adenoma and SS simultaneously is reasonable.

In the present study, we found no correlation between the development of severe duodenal adenomatosis or duodenal cancer and the specific location of the mutation in *APC*, although the risk factors for advanced duodenal adenoma and SS IV were classic FAP, a history of CRC, a history of colectomy, and a 3' side mutation. Therefore, it might be better to perform more intensive surveillance for patients with these risk factors instead. In the current study, no patients developed invasive cancer with annual surveillance; thus, annual surveillance may be sufficient, even for high-risk patients. However, for low-risk patients such as those with attenuated FAP and small polyps, the interval of surveillance endoscopy can be prolonged.

In the current study, 18% of the patients progressed to stage IV. Surgical intervention should be considered for such patients because of the risk of the development of duodenal cancer. Because surgical intervention (typically pancreatoduodenectomy) for the treatment of duodenal cancer or stage IV disease can be extremely invasive and is accompanied by severe sequelae, it should be avoided if possible. This is especially true for patients who have undergone total colectomy, which is a standard intervention for classic FAP. Although endoscopic intervention has not been actively recommended because of the high risk of adverse events and frequent recurrences, we have adopted safer endoscopic interventions for the management of duodenal neoplasms in patients with FAP [29]. We also recommended treatment intervention for patients with an advanced duodenal adenoma in the present study, and no invasive cancer developed in our cohort. This indicates that early intensive endoscopic management can avoid development of duodenal cancer or progression to SS IV in patients with FAP [10, 29-31].

Three main limitations of this study should be considered. First, this was a retrospective study involving a small number of patients in a small number of facilities. Therefore, the study findings may not be highly generalizable. However, we enrolled all consecutive patients who met the inclusion criteria, and the ability to collect detailed, high-quality endoscopic data from two linked

## Table 3 Progression of patients with stage 0 to III (n = 113)

	Progression (n = 80)	No progression (n = 33)	P-value
Sex			
Male	34 (43)	12 (36)	0.55
Female	46 (58)	21 (64)	
<b>Body mass index</b> , kg/m <sup>2</sup>	20.7 (14.7–36.2)	21.9 (15.6–27.3)	0.80
Alcohol intake			
Past/current drinker	39 (49)	13 (39)	0.36
Never-drinker	41 (51)	20 (61)	
Smoking			
Past/current smoker	19 (24)	7 (21)	0.96
Never-smoker	61 (76)	26 (79)	
Family history of FAP			
Positive	69 (86)	29 (88)	0.90
Negative	6 (8)	1 (3)	
Unknown	5 (6)	3 (9)	
Helicobacter pylori status			
Positive	12 (15)	8 (24)	0.69
Negative	66 (83)	25 (76)	
Unknown	2 (3)	0 (0)	
History of colorectal cancer			
Positive	6 (8)	1 (3)	0.64
Negative	74 (93)	32 (97)	
Type of polyposis			
Classic FAP	77 (96)	23 (70)	< 0.001
Attenuated FAP	3 (4)	10 (30)	
Age at diagnosis of FAP, years	24 (13–49)	27 (16–64)	0.08
Colectomy			
Positive	33 (41)	5 (15)	0.01
Negative	47 (59)	28 (85)	
Duodenal ampullary adenoma			
Positive	14 (18)	4 (12)	0.67
Negative	66 (83)	29 (88)	
Site of APC mutation (mutation at codons 279–1309)*			
Mutation within the region	41 (51)	20 (60)	0.21
Mutation outside the region	33 (41)	8 (24)	
Site of APC mutation*			
3': distal mutation group from codon 1051	23 (29)	14 (42)	0.08
5': proximal mutation group from codon 1051	51 (64)	14 (42)	
Site of APC mutation*			
3': distal mutation group (exons 10–15)	44 (55)	21 (64)	0.22
5': proximal mutation group (exons 1–9)	30 (38)	7 (21)	
Site of APC mutation (mutation at codons 1401–1580)*			
Mutation within the region	4 (5)	0 (0)	0.49
Mutation outside the region	70 (88)	28 (85)	
Site of APC mutation (mutation at codons 1250–1460)*			
Mutation within the region	7 (9)	0 (0)	0.21
Mutation outside the region	67 (84)	28 (85)	

Data are presented as n (%) or median (range)

FAP, familial adenomatous polyposis; APC, adenomatous polyposis coli gene

\*Excluding patients with unknown mutation site

facilities was advantageous. In addition, we focused on patients who met the diagnostic criteria for FAP and were positive for the *APC* mutation; therefore, we were able to extract and examine more reliable data. In future studies, we hope that patient data will be collected at

multiple facilities using registries. In addition, we hope our data will contribute to a future meta-analysis, which can provide more robust information. Second, the observation period was too short to determine the real lifetime risk of duodenal neoplasms. This might have caused

## **Table 4** Progression of patients with stage 0 to I (n = 72)

	Progression (n = 60)	No progression (n = 12)	P-value
Sex			
Male	25 (42)	4 (33)	0.83
Female	35 (58)	8 (67)	
<b>Body mass index</b> , kg/m <sup>2</sup>	20.7 (14.7–36.2)	23.1 (19.1–27.3)	0.09
Alcohol intake			
Past/current drinker	30 (50)	6 (50)	0.75
Never-drinker	30 (50)	6 (50)	
Smoking			
Past/current smoker	17 (28)	1 (8)	0.27
Never-smoker	43 (72)	11 (92)	
Family history of FAP			
Positive	52 (87)	9 (75)	0.38
Negative	4 (7)	0 (0)	
Unknown	4 (7)	3 (25)	
Helicobacter pylori status			
Positive	8 (13)	4 (33)	0.45
Negative	50 (83)	8 (67)	
Unknown	2 (3)	0 (0)	
History of colorectal cancer			
Positive	6 (10)	0 (0)	0.57
Negative	54 (90)	12 (100)	
Type of polyposis			
Classic FAP	58 (97)	3 (25)	< 0.001
Attenuated FAP	2 (3)	9 (75)	
Age at diagnosis of FAP, years	25.5 (13–49)	37.5 (30–64)	< 0.001
Colectomy			
Positive	25 (42)	1 (8)	0.06
Negative	35 (58)	11 (92)	
Duodenal ampullary adenoma			
Positive	11 (18)	0 (0)	0.24
Negative	49 (82)	12 (100)	
Site of APC mutation (mutation at codons 279–1309)*			
Mutation within the region	27 (45)	8 (67)	0.07
Mutation outside the region	27 (45)	1 (8)	
Site of APC mutation*			
3': distal mutation group from codon 1051 5': proximal mutation group from codon 1051	17 (28) 37 (62)	4 (7) 5 (8)	0.70
Site of APC mutation*			
3': distal mutation group (exons 10–15)	27 (45)	7 (58)	0.23
5': proximal mutation group (exons 1–9)	27 (45)	2 (17)	0.20
Site of APC mutation (mutation at codons 1401–1580)*			
Mutation within the region	2 (3)	0 (0)	0.66
Mutation outside the region	52 (87)	9 (75)	
Site of APC mutation (mutation at codons 1250–1460)*			
Mutation within the region	3 (5)	0 (0)	0.90
Mutation outside the region	51 (85)	9 (75)	

Data are presented as n (%) or median (range)

FAP, familial adenomatous polyposis; APC, adenomatous polyposis coli gene

\*Excluding patients with unknown mutation site

an underestimation of the risk of duodenal neoplasms; therefore, the study findings should be interpreted as at least a progression of duodenal manifestations in most of the patients with FAP. Finally, our study did not include ampullary adenoma to clarify the progression of NADAs separate from ampullary adenoma. Because half of duodenal cancers were reported to develop from ampullary adenomas, we should consider them as well as NADAs for the long-term prognosis of patients with FAP. However, the treatment strategies for ampullary adenoma and



Fig. 3 Incidence of HGD according to Spigelman stage at the first esophagogastroduodenoscopy procedure. HGD, high-grade dysplasia

NADAs are different, and we needed to investigate them separately.

# Conclusions

NADAs in patients with FAP frequently become exacerbated, and all patients with FAP should be surveyed to prevent the development of duodenal cancer. An early, minimally invasive therapeutic intervention should be considered to avoid disease progression.

# **Table 5** Risk factors for development of $\geq$ 10-mm adenoma (n = 146)

	≥10 mm (n=37)	< 10 mm (n = 1069)	P-value
Sex		· · ·	
Male	16 (43)	47 (43)	0.99
Female	21 (57)	62 (57)	
<b>Body mass index</b> , kg/m <sup>2</sup>	20.9 (14.7-30.1)	21.0 (15.6–37.2)	0.80
Alcohol intake			
Past/current drinker	21 (57)	51 (47)	0.29
Never-drinker	16 (43)	58 (53)	
Smoking			
Past/current smoker	8 (22)	28 (26)	0.78
Never-smoker	29 (78)	81 (74)	
Family history of FAP			
Positive	32 (86)	94 (86)	0.87
Negative	1 (3)	7 (6)	
Unknown	4 (10)	8 (7)	
Helicobacter pylori status			
Positive	6 (16)	22 (20)	0.63
Negative	28 (76)	84 (77)	
Unknown	3 (8)	3 (3)	
History of colorectal cancer			
Positive	6 (16)	4 (4)	0.03
Negative	31 (84)	105 (96)	
Type of polyposis			
	37 (100)	96 (88)	0.06
Attenuated FAP	0(0)	13 (12)	
Age at diagnosis of FAP, years	29 (15–59)	25 (13–64)	0.22
Colectomy			
Positive	23 (62)	24 (22)	< 0.001
Negative	14 (38)	85 (78)	
Duodenal ampullary adenoma		/	
Positive	6 (16)	15 (14)	0.92
	31 (84)	94 (86)	
Site of APC mutation (mutation at codons 279–1309)*	0.6 (70)	52 (10)	
Mutation within the region	26 (70)	53 (49)	0.08
	9 (24)	45 (39)	
Site of APC mulation"	15 (40)	22 (20)	0.21
3 : distal mutation group from codon 1051	15 (40)	32 (29)	0.31
Site of APC mutation*	20 (34)	04 (59)	
2/ distel mutation	20 (70)		0.00
3 : distal mutation group (exons 10–15)	29 (78)	57 (52)	0.02
5 : proximal mutation group (exons 1–9)	6 (16)	39 (36)	
Site of APC mutation (mutation at codons 1401–1580)*	2 (5)		
Mutation within the region	2 (5)	4 (4)	0.92
withation outside the region	33 (89)	92 (84)	
Sile OIAPC Initiation (Initiation at Codons 1250–1460)*	2 (0)	7 (7)	0.00
iviulation within the region	3 (8) 22 (94)	/ (b) 90 (92)	0.90
mutation outside the region	32 (00)	09 (02)	

Data are presented as n (%) or median (range)

FAP, familial adenomatous polyposis; APC, adenomatous polyposis coli gene

\*Excluding patients with unknown mutation site

# **Table 6** Risk factors for development of HGD (n = 146)

	HGD (n = 32)	No HGD (n = 114)	P-value
Sex			
Male	11 (34)	52 (46)	0.26
Female	21 (66)	62 (54)	
<b>Body mass index</b> , kg/m <sup>2</sup>	21.6 (14.7–30.1)	20.9 (16.8–27.5)	0.89
Alcohol intake			
Past/current drinker	15 (47)	57 (50)	0.75
Never-drinker	17 (53)	57 (50)	0.75
Smoking	x		
Past/current smoker	5 (16)	31 (27)	0.27
Never-smoker	27 (84)	83 (73)	0.27
Family history of FAP			
Positive	30 (93)	96 (84)	0.66
Negative	1 (3)	7 (6)	0.00
Unknown	1 (3)	11 (10)	
Helicobacter pylori status			
Positive	5 (16)	23 (20)	0.92
Negative	26 (81)	86 (75)	0.02
Unknown	1 (3)	5 (4)	
History of colorectal cancer			
Positive	4 (12)	6 (5)	0.30
Negative	28 (88)	108 (95)	0.00
Type of polyposis			
Classic FAP	30 (94)	103 (90)	0.81
Attenuated FAP	2 (6)	11 (10)	0.01
Age at diagnosis of FAP years	26 (15-59)	26 (13–64)	0.59
Colectomy	20(13/33)	20(13/01)	0.07
Positivo	17 (53)	30 (26)	0.004
Negative	15 (47)	84 (74)	0.004
Adenoma size of $> 10 \text{ mm}$			
	4 (12)	3 (3)	0.07
Negative	28 (88)	111 (97)	0.07
Duodenal ampullary adenoma	20 (00)		
Positivo	7 (22)	14 (12)	0.20
Negative	25 (78)	100 (88)	0.20
Site of APC mutation (mutation at codons 270, 1300)*	23 (70)	100 (00)	
Mutation within the region	19 (56)	61 (54)	0.77
Mutation outside the region	13 (41)	39 (34)	0.77
Site of APC mutation*	13(11)	33 (34)	
2': dictal mutation group from codop 1051	10 (21)	27 (22)	0.62
5': provimal mutation group from codon 1051	21 (66)	63 (55)	0.05
Site of APC mutation*	21 (00)	05 (55)	
2' distal mutation	22 (60)	64 (E6)	0.60
5': provimal mutation group (exons 10–15)	9 (28)	36 (32)	0.02
Site of APC mutation (mutation at codons 1/01–1580)*	5 (20)	56 (32)	
Mutation within the region	2 (6)	4 (4)	0.04
Mutation outside the region	∠ (0) 29 (90)	96 (84)	0.94
Site of APC mutation (mutation at codons 1250, 1460)*	27 (30)	20 (01)	
Mutation within the ragion	4 (12)	7 (6)	0.51
Mutation outside the region	4 (1 <i>5)</i> 27 (84)	/ (U) Q3 (82)	U.5 I
	2/ (04)	93 (02)	

Data are presented as n (%) or median (range)

 $\mathsf{HGD}, \mathsf{high}\text{-}\mathsf{grade}\,\mathsf{dysplasia}; \mathsf{FAP}, \mathsf{familial}\,\mathsf{adenomatous}\,\mathsf{polyposis}; \mathsf{APC}, \mathsf{adenomatous}\,\mathsf{polyposis}\,\mathsf{coli}\,\mathsf{gene}$ 

\*Excluding patients with unknown mutation site

# **Table 7** Risk factors for development of stage IV (n = 146)

	Stage IV (+) (n = 27)	Stage IV (–) (n=119)	P-value
Sex			
Male	9 (33)	54 (45)	0.35
Female	18 (67)	65 (55)	
Body mass index, kg/m <sup>2</sup>	20.7 (14.7–30.1)	21.0 (16.8–27.5)	0.51
Alcohol intake			
Past/current drinker	13 (48)	59 (50)	0.94
Never-drinker	14 (52)	60 (50)	
Smoking			
Past/current smoker	5 (19)	31 (26)	0.57
Never-smoker	22 (81)	88 (74)	
Family history of FAP			
Positive	24 (89)	102 (86)	0.98
Negative	1 (4)	7 (6)	
Unknown	2 (7)	10 (8)	
Helicobacter pylori status			
Positive	6 (22)	22 (18)	0.91
Negative	20 (74)	92 (77)	
Unknown	1 (4)	5 (4)	
History of colorectal cancer			
Positive	3 (11)	7 (6)	0.58
Negative	24 (89)	112 (94)	
Type of polyposis			
Classic FAP	26 (96)	107 (90)	0.50
Attenuated FAP	1 (4)	12 (10)	
Age at diagnosis of FAP, years	25 (15–53)	26 (14–64)	0.88
Colectomy			
Positive	14 (52)	33 (28)	0.02
Negative	13 (48)	86 (72)	
Adenoma size of ≥ 10 mm			
Positive	3 (11)	4 (3)	0.22
Negative	24 (89)	115 (97)	
Duodenal ampullary adenoma			
Positive	5 (19)	16 (13)	0.71
Negative	22 (81)	103 (87)	
Site of APC mutation (mutation at codons 279–1309)*			
Mutation within the region	17 (63)	62 (52)	0.52
Mutation outside the region	8 (30)	44 (37)	
Site of APC mutation*			
3': distal mutation group from codon 1051	7 (26)	40 (34)	0.50
5': proximal mutation group from codon 1051	18 (67)	66 (55)	
Site of APC mutation*			
3': distal mutation group (exons 10–15)	20 (74)	66 (55)	0.15
5': proximal mutation group (exons 1–9)	5 (19)	40 (34)	
Site of APC mutation (mutation at codons 1401–1580)*			
Mutation within the region	1 (4)	5 (4)	0.71
Mutation outside the region	24 (89)	101 (85)	
Site of APC mutation (mutation at codons 1250–1460)*		- *	
Mutation within the region	3 (11)	8 (7)	0.75
Mutation outside the region	22 (81)	98 (82)	

Data are presented as n (%) or median (range)

FAP, familial adenomatous polyposis; APC, adenomatous polyposis coli gene

\*Excluding patients with unknown mutation site

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### Author contributions

Conception and design: HN, YT, HI; Data Acquisition: HN, YS, SI, HY, YE, RI, TT, TY, HI; Drafting of the article: HN, YT; Critical revision for important intellectual content: HI, MM; Final approval of the article: All authors.

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#### Data Availability

The following data will be made available with publication: complete deidentified patient data set (contact Yoji Takeuchi; e-mail, yoji.endoscopy@ oici.jp). The following supporting documents will be made available with publication: analytic/statistical code (contact Yoji Takeuchi; e-mail, yoji. endoscopy@oici.jp). These data will be made available for any purpose to researchers whose proposed use of the data has been approved.

## Declarations

## Ethics approval and consent to participate

This study was approved by the Institutional Review Board of Osaka International Cancer Institute (2018–18229 and 2019–19006). Informed consent was obtained in the form of an opt-out on the institutional websites. The study was performed in accordance with the guidelines of the World Medical Association's Declaration of Helsinki. All authors consent to publish this manuscript.

#### **Consent for publication**

Not applicable.

#### **Competing interests**

Yoji Takeuchi has received honoraria for lectures from Olympus, Boston Scientific Japan, Daiichi-Sankyo, Miyarisan Pharmaceutical, Asuka Pharmaceutical, AstraZeneca, EA Pharma, Zeria Pharmaceutical, Fujifilm, Kaneka Medix, and Kyorin Pharmaceutical. The other authors declare that they have no competing interests.

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