Prognostic relevance of FGFR2 expression in stage II/III gastric cancer with curative resection and S-1 chemotherapy

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Abstract. Curative gastrectomy and adjuvant chemotherapy using S-1 is a standard treatment for stage II/III gastric cancer in Japan. The purpose of the present study was to evaluate the prognostic relevance of fibroblast growth factor receptor (FGFR)2 expression in patients with stage II/III gastric cancer that underwent postoperative adjuvant chemotherapy with S-1. Formalin-fixed paraffin-embedded surgical specimens were retrospectively examined in 167 patients with stage II/III gastric cancer that underwent curative gastrectomy followed by adjuvant S1 chemotherapy. FGFR2 expression was measured using immunohistochemistry (IHC) staining. The IHC results for FGFR2 were as follows: Grade 1+, 32; grade 2+, 80; grade 3+, 55 patients. The FGFR2 expression level was not significantly associated with relapse-free or overall survival rates. However, in the diffuse type, the FGFR2 expression level tended to be negatively correlated with relapse-free survival. In particular, the proportion of patients who recurred >5 years following surgery was significantly larger in the FGFR2 grade 3+ group than in the grade 1+, 2+ group (4/22 vs. 1/35; P=0.047). The recurrent sites of long-term failure were mostly peritoneum among the diffuse type. To the best of our knowledge, the present study indicated for the first time that FGFR2 could predict long-term failure of adjuvant S-1 chemotherapy in curative advanced gastric cancer. There was no interaction between FGFR2 expression and patient survival outcomes in stage II/III gastric cancer. Patients with FGFR2 3+ in stage II/III gastric cancer should carefully be followed-up for >5 years after surgery.

Introduction

Gastric cancer is one of the leading causes of cancer related death worldwide (1). Advanced gastric cancer exhibits poor prognosis, and even optimal combination modalities of surgery, chemotherapy, and radiotherapy cannot attain satisfactory survival outcomes.

S-1 is a fluoropyrimidine containing compound consisting of a combination of tegafur, gimeracil, and oteracil potassium. A randomized phase III trial of adjuvant chemotherapy with S-1 (ACTS-GC) demonstrated that surgery with S-1 adjuvant chemotherapy surpassed surgery alone in stage II/III gastric cancer (2,3). Thereafter, curative gastrectomy and adjuvant chemotherapy using S-1 have become a standard treatment for stage II/III gastric cancer in Japan. However, in the subgroup analysis of ACTS-GC, the 5-year overall survival (OS) rate of patients with stage III disease was 50.2%, leaving room for improvement.

Molecular targeted therapy is an alternative therapeutic tool for advanced cancer, and the receptor tyrosine kinase (RTK) family is one such promising candidate target. The fibroblast growth factor (FGF) family which includes important regulatory factors of cell growth and differentiation, has been found to be involved in embryonic development, angiogenesis, and tumorigenesis (4,5). The FGF receptor (FGFR) family, one type of RTK family, is a transmembrane tyrosine kinase receptor involved in signaling via interaction with the FGF family. To date, four different members of the FGFR family, FGFR1, FGFR2, FGFR3 and FGFR4, have been cloned and characterized (6-8). FGFR2 gene amplification was initially found in a gastric cancer cell line originating from diffuse type gastric cancer (9). FGFR2 has been demonstrated to be a poor prognostic biomarker (10,11) and antibodies (12-14) or small molecule inhibitors (15-18) targeting FGFR2 can suppress gastric cancer progression in vivo and in vitro.

The human epidermal growth factor receptor (EGFR/HER) family is involved in complex and tightly controlled signaling pathways that regulate various cellular functions, such as cell proliferation, organ development, and organ repair. Among four members of the HER family (EGFR, HER2, HER3 and HER4), HER3 is indicated to play an important role in HER signaling and drug resistance (19). With dimerization with EGFR or HER2, HER3 signals through phosphatidylinositol 3-kinase pathway (20,21).

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We have previously analyzed various RTK expressions and reported that HER3 and EGFR were significant and marginally significant independent prognostic factors, respectively, in patients with stage II/III gastric cancer who underwent curative gastrectomy followed by adjuvant chemotherapy with S-1 (22). However, we have not included this critical RTK of FGFR2 in the study. We hypothesized that FGFR2 alone or that in combination with HER3 and EGFR plays an important role on survival outcomes of these patients. Therefore, we analyzed correlation between HER3 or EGFR expression and FGFR2 expression using the identical patients of the previous study, and evaluated the prognostic impact of FGFR2 expression in patients with stage II/III gastric cancer treated by curative gastrectomy followed by adjuvant chemotherapy with S-1.

Materials and methods

Patients. From January 2000 to December 2010, 172 patients with stage II/III advanced gastric cancer underwent curative surgery followed by adjuvant S-1 chemotherapy at Kitasato University Hospital. Of the 172 patients, informed consent to use specimens was obtained from 167 patients, for whom the median follow up term was 75 months (inter quartile range, 61-90 months). These patients were identical to the patients whom we had analyzed previously (22).

This study was conducted in accordance with the Declaration of Helsinki and was approved by the Research Ethics Committee of the Kitasato University School of Medicine (Sagamihara, Japan).

Surgery. Gastrectomy with D1 lymph node dissection was performed for clinical stage IA gastric cancer (n=25). Gastrectomy with D2 lymph node dissection was otherwise performed (n=142). The extent of lymph node dissection was determined according to the Japanese Classification of Gastric Carcinoma (second English edition) (23).

Postoperative chemotherapy of S-1. The dose of S-1 was based on body-surface area: <1.25 m² (80 mg daily); ≥1.25 m² but <1.50 m² (100 mg daily); ≥1.50 m² (120 mg daily). The adjuvant S-1 was administered for 4 weeks followed by 2 weeks rest. This 6-week cycle was repeated principally during the first year after surgery.

Immunohistochemistry (IHC) analysis. Tumor specimens used in this study were derived from routine formalin-fixed, paraffin embedded tissue samples obtained from resected gastric cancer specimens. Sections (3- μ m thick) were cut from the paraffin blocks and mounted on silanized slides. For IHC analysis of FGFR2, tumor tissue sections were deparaffinized in xylene and dehydrated with graded ethanol. After washing with distilled water, antigen unmasking was performed with a pressure cooker using citrate buffer. Endogenous peroxidase activity was blocked by incubation in 3% H₂O₂/methanol for 15 min, and nonspecific antibody binding was blocked by incubation with 1% diluted normal horse serum for 30 min. Sections were then incubated at room temperature for 60 min with the following antibodies: Mouse FGFR2 monoclonal antibody (M01, clone 1G3) (dilution of 1:800; Abnova, Heidelberg, Germany,). Immune complexes were detected with a Vectastain Elite ABC kit (Vector Laboratories, Inc., Burlingame, CA, USA) according to the manufacturer's instructions. These immune complexes were detected using the 3,3'-diaminobenzidine tetrahydrochloride substrate with/without nickel ammonium sulfate (DAB; Vector Laboratories, Inc.) as a chromogen for 5 min. Sections were counterstained with hematoxylin.

IHC staining of HER3 and EGFR was performed as previously reported (22).

Scoring system of IHC staining. To assess IHC staining, all slides were observed under a microscope. Investigators were blinded to the prognostic analysis data. We adopted almost the same scoring system for FGFR2 as was reported by Nagatsuma *et al* (24). We defined the scores for FGFR2 as follows: 0, staining reactivity in <50 % of cancer cells; 1+, cytoplasm and/or nuclear reactivity with faint staining in more than 50% of the cancer cells; 2+, cytoplasm and/or nuclear reactivity with strong staining in more than 50% of the cancer cells; 3+, cytoplasm and/or nuclear reactivity with strong staining in more than 50% of the cancer cells.

The scoring system for IHC staining of HER3 and EGFR was described before (22).

Statistical analysis. Relapse-free survivals (RFS) were measured from the dates of surgery to the dates of relapse of gastric cancers or dates of last follow-up. Patients who died from causes other than gastric cancer were regarded as censored at the time of death. Patients who were alive at the times of their last visits were also regarded as censored. Student's t-test was used to analyze continuous variables, and Chi square test or Fisher's exact test was used to analyze categorical variables. Survival curves were estimated using the Kaplan-Meier method, and the statistical significance of differences between survival curves was assessed using the log-rank test. Survival analyses were conducted using a Cox proportional hazards model. All calculations were performed using the JMP® 11.2 (SAS Institute Inc., Cary, NC, USA), and P<0.05 was considered to indicate a statistically significant difference.

Results

Patient characteristics and stage distributions. The characteristics of patients included in this study are listed in Table I. More than half of the patients had pT4a tumors. The FGFR2 IHC results were obtained in all 167 specimens as follows: FGFR2 grade 1+, 32 (19%); FGFR2 grade 2+, 80 (48%); FGFR2 grade 3+, 55 (33%). Representative examples of immunostaining for FGFR2 in diffuse type cancer (Fig. 1A) and intestinal type cancer (Fig. 1B).

Relations of FGFR2 expression to clinicopathological features and correlation between FGFR2 expression and HER3 or EGFR expression. We divided the study samples into two groups [low FGFR2 expression group: FGFR2 IHC grade 1+, 2+ (n=112); high FGFR2 expression group: FGFR2 IHC grade 3+ (n=55)]. The relations of IHC staining levels

Table I. Characteristics of the patients.

Category	n=167
Age (years)	
Median (range)	65 (35-83)
Sex, n (%)	
Male	117 (70)
Tumor stage (TNM 7th), n (%)	
T2	28 (17)
T3	34 (20)
T4a	103 (62)
T4b	2 (1)
Nodal stage (TNM 7th), n (%)	
NO	24 (14)
N1	43 (26)
N2	38 (23)
N3	62 (37)
TNM 7th stage, n (%)	
IIA	13 (8)
IIB	36 (22)
IIIA	43 (26)
IIIB	37 (22)
IIIC	38 (23)
Histological type, n (%)	
Diffuse	58 (35)
Intestinal	109 (65)
FGFR2 IHC status, n (%)	
0	0
1+	32 (19)
2+	80 (48)
3+	55 (33)
HER3 IHC status, n (%)	
0	69 (41)
1+	68 (41)
2+	30 (18)
3+	0
EGFR IHC status, n (%)	
0	0
1+	62 (37)
2+	57 (34)
3+	48 (29)

TNM, tumor node metastasis; IHC, immunohistochemistry; EGFR, epidermal growth factor receptor; HER3, human epidermal growth factor receptor 3.

to clinicopathological features were then examined. FGFR2 IHC staining level was significantly related to patients' age (P=0.043) and tended to be related to lymph node metastasis (P=0.099) (Table II). There was no correlation between FGFR2 expression and HER3 expression, whereas the FGFR2 expression level was correlated with the EGFR expression level (P=0.037; Fig. 2).

Table II. Relations of FGFR2 IHC staining levels to clinicopathological features.

Variable/category	FGFR2 $1+/2+$ (n=112)	FGFR2 3 + (n=55)	P-value
	11721 (II=112)	51 (II-55)	i varae
Age			0.043
<65	53	17	
≥65	59	38	
Sex			1.000
Male	78	39	
Female	34	16	
Serosal invasion			0.734
Absent	43	19	
Present	69	36	
Lymph node metastasis			0.099
Absent	20	4	
Present	92	51	
Cancer stage, n (%)			0.283
IIA/IIB	36	13	
IIIA/IIIB/IIIC	76	42	
Histologic type, n (%)			0.604
Diffuse	37	21	
Intestinal	75	34	

IHC, immunohistochemistry; FGFR, fibroblast growth factor receptor.

Effects of clinicopathological factors and FGFR2 expression on RFS and OS. Five-year RFS and OS were 67.7 and 70.1%, respectively. Elderly patients aged 65 or more had significantly worse RFS and OS than young patients aged less than 65 [hazard ratio (HR) 1.75, 95% confidence interval (CI) 1.02-3.10, P=0.042 and HR 2.20 95% CI 1.28-3.94, P=0.005], while female patients had significantly better OS than male patients (HR 0.47, 95% CI 0.24-0.85, P=0.012). Patients with TNM stage III had significantly worse RFS than those with TNM stage II (HR 2.68, 95% CI 1.38-5.87, P=0.003). On the other hand, there was no relation of FGFR2 expression status to RFS or OS in the study group (Table III, Fig. 3). However, when we restricted patients to those with diffuse type cancer, patients with high FGFR2 expression had slightly worse RFS than those with low FGFR2 expression (P=0.37; Fig. 3C). Moreover, when we combined FGFR2 and HER3 expression, prognostic impact of HER3 was augmented by FGFR2 expression with 5-year survival rate of 95 and 51% in patients with FGFR2/HER3 3+/0 and FGFR2/HER3 3+/1+ or 2+, respectively (P<0.001; Fig. 4A). Statistical significant differences were also found between the FGFR2/HER3 3+/0 group and the FGFR2/HER3 1+ or 2+/1+ or 2+ group and between the FGFR2/HER3 1+ or 2+/0 group and the FGFR2/HER3 3+/1+ or 2+ group with P values of 0.021 and 0.004, respectively. When we combined FGFR2 and EGFR expression, no correlation was found in RFS (Fig. 4B).

We further analyzed RFS using the combined expressions of the three biomarkers (Fig. 5). Patients with FGFR2 3+, HER3 1+/2+, and EGFR 3+ belonged to one of the worst



Figure 1. Immunohistochemical staining of FGFR2 expression in the stage II/III gastric cancer of (A) diffuse type and (B) intestinal type (magnification, x100). FGFR, fibroblast growth factor receptor.

		HER3							
	2+ 1+ 0				HE	R3			
	3+	9	26	20				2+/1+	0
FGFR2	2+	16	29	35			3+/2+	80	55
	1+	5	13	14] [FGFKZ	1+	18	14
	P=0.77 EGFR								P=0.84
			EGFR						
		3+	EGFR 2+	1+				EG	FR
	3+	3+ 19	EGFR 2+ 17	1+ 19				EG 3+	FR 2+/1+
FGFR2	3+ 2+	3+ 19 15	EGFR 2+ 17 30	1+ 19 35			3+/2+	EG 3+ 34	FR 2+/1+ 101
FGFR2	3+ 2+ 1+	3+ 19 15 14	EGFR 2+ 17 30 10	1+ 19 35 8		FGFR2	3+/2+	EG 3+ 34 14	FR 2+/1+ 101 18

Figure 2. Correlation between FGFR2 expression and HER3 or EGFR expression. There is no correlation between FGFR2 expression and HER3 expression, whereas the FGFR2 expression level was correlated with the EGFR expression level. FGFR, fibroblast growth factor receptor; HER, human epidermal growth factor receptor; EGFR, EGFR, epidermal growth factor receptor.

survival groups with 5 year relapse-free and OS rates of 53 and 59%, respectively. However, no significant difference was found between the FGFR2 3+, HER3 1+/2+, EGFR 3+ group, the FGFR2 1+, 2+, HER3 1+/2+, EGFR 3+ group, and the FGFR2 3+, HER3 1+/2+, EGFR 1+, 2+ group (P=0.82).

Timing of recurrence. Of the 165 patients, 57 recurred. Proportion or patients with recurrence was similar between the high FGFR2 expression group and the low FGFR2 expression group [40% (22/55) vs. 31% (35/112), P=0.27]. Of the 57 recurred patients, 5 recurred more than 5 years after surgery (Table IV). Proportion of patients who recurred more than 5 years after surgery was significantly larger in the high FGFR2 expression group than in the low FGFR2 expression group [18% (4/22) vs. 2.9% (1/35), P=0.047] (Fig. 6). All the 5 patients had tumors of diffuse type and serosal invasion (pT4a). Reciprocal relation was found between FGFR2 expression and EGFR expression. That is, 4 had tumors with FGFR2 grade 3+ and EGFR grade 1+, while the other had tumors with FGFR2 grade 3+ and EGFR grade 3+. One female patient with high FGFR2 expression recurred more than 10 years after surgery. She was 55 years old at the time of surgery and underwent total gastrectomy with D2 lymph node dissection. She had a tumor with serosal invasion (pT4a) but no lymph node metastasis at the time of surgery. At the 10th year check-up, CT scan revealed that she had para-aortic lymph node metastasis.

Discussion

The present study retrospectively evaluated the influence of FGFR2 expression on the survival outcomes of patients with stage II/III gastric cancer who underwent curative gastrectomy followed by adjuvant chemotherapy using S1. The FGFR2

				RFS		OS			
Variable	Category	No. of patients	5-year survival (%)	HR (95% CI)	P-value	5-year survival (%)	HR (95% CI)	P-value	
Age	<65	70	76.6	1.0	0.042	83.1	1.0	0.005	
	≥65	97	60.8	1.75 (1.02-3.10)		60.9	2.20 (1.28-3.94)		
Sex	Male	117	62.5	1.0	0.084	62.9	1.0	0.012	
	Female	50	79.5	0.59 (0.31-1.07)		85.8	0.47 (0.24-0.85)		
TNM stage	IIA/IIB	49	84.5	1.0	0.003	78.3	1.0	0.088	
-	IIIA/IIIB/IIIC	118	60.8	2.68 (1.38-5.87)		66.7	1.67 (0.93-3.22)		
Histologic type	Differentiated	58	70.4	1.0	0.410	61.9	1.0	0.096	
	Undifferentiated	109	62.4	0.80 (0.47-1.38)		74.6	0.64 (0.38-1.09)		
FGFR2 IHC status	1+, 2+	112	68.1	1.0	0.377	71.2	1.0	0.411	
	3+	55	66.6	1.27 (0.74-2.16)		67.8	1.25 (0.73-2.09)		

Table III. Cox proportional hazards analysis of clinicopathological factors for OS and RFS.

RFS, relapse-free survival; OS, overall survival; HR, hazard ratio; CI, confidence interval; TNM, tumor node metastasis; FGFR, fibroblast growth factor receptor; IHC, immunohistochemistry.



Figure 3. Kaplan-Meier curves for (A) relapse-free survival and (B) overall survival in patients with stage II/III gastric cancer stratified by the FGFR2 IHC staining levels. There is no relation of FGFR2 expression status to relapse-free survival or overall survival. Kaplan-Meier curves for (C and E) relapse-free survival and (D and F) overall survival in patients with diffuse type, intestinal type stage II/III gastric cancer stratified by the FGFR2 IHC staining levels. In patients with diffuse type cancer, the FGFR2 expression tended to be negatively correlated to (C) relapse-free survival (P=0.37). FGFR, fibroblast growth factor receptor; IHC, immunohistochemistry.



Figure 4. Kaplan-Meier curves for relapse-free survival in patients with stage II/III gastric cancer stratified by combination of (A) FGFR2 and HER3 expressions, and (B) FGFR2 and EGFR expressions. Prognostic impact of HER3 is augmented by FGFR2 expression with 5-year survival rate of 78 and 51% in patients with FGFR2/HER3 1+ or 2+/0 and FGFR2/HER 3+/1+ or 2+, respectively. When we combined FGFR2 and EGFR expression, no correlation was found in relapse-free survival. FGFR, fibroblast growth factor receptor; HER, human epidermal growth factor receptor; EGFR, EGFR, epidermal growth factor receptor.



Figure 5. Kaplan-Meier curves for relapse-free survival in patients with stage II/III gastric cancer stratified by combination of FGFR2, HER3, and EGFR expressions. Although patients with FGFR2 3+, HER3 1+/2+, and EGFR 3+ belongs to one of the worst survival groups, no significant difference is found between the FGFR2 3+, HER3 1+/2+, and EGFR 3+ group, FGFR2 1+/2+, HER3 1+/2+, and EGFR 3+ group, and FGFR2 3+, HER3 1+/2+, and EGFR 1+/2+, group (P=0.82). FGFR, fibroblast growth factor receptor; HER, human epidermal growth factor receptor; EGFR, EGFR, epidermal growth factor receptor.

expression was not significantly associated with RFS or OS. When we restricted patients to those with diffuse type cancer, FGFR2 expression levels tended to be negatively correlated with RFS. Moreover, proportion of patients who recurred more than 5 years after surgery was significantly larger in the high FGFR2 expression group than in the low FGFR2 expression group.

The FGFR2 expression level was not significantly associated with RFS or OS. Nagatsuma *et al* (24) reported that high expression of FGFR 2 correlated with tumor progression and survival in patients with gastric cancer. However, they did not refer to adjuvant chemotherapies and stage distribution. On the other hand, patients in this current study had pathological stage II or III and received adjuvant chemotherapy using S1. Adjuvant S1 chemotherapy might have some power to negate the difference of survival outcomes between different FGFR2 expression levels.

When we restricted patients to those with diffuse type gastric cancer, survival outcome was not significantly but slightly correlated with FGFR2 expression. High FGFR2 expression tended to be slightly worse survival outcomes. Inokuchi *et al* reported that high FGFR2 expression significantly correlated with tumor progression and survival only in diffuse type gastric cancer (25). If our study patients had

not been treated with S1 adjuvant chemotherapy, the negative effect of FGFR2 expression might have significance on survival outcomes.

We previously reported that HER3 expression was a significant independent prognostic factor in patients with stage II/III gastric cancer who receive curative resection and adjuvant chemotherapy with S-1. In our current study, the analyzed patients of which are identical to those of previous one, FGFR2 overexpression augmented prognostic impact of HER3 expression. EGFR family members are reported to be downstream targets of the amplified and highly activated FGFR2 kinase (26). That may explain the augmentation by FGFR2 expression. On the other hand, among the patients without expression of HER3, patients with high FGFR2 expression had higher survival than those with low FGFR2 expression. However, the difference between these patients is not statistically significant in the log-rank test (P=0.19). FGFR2 overexpression would not be as strongly associated with RFS as HER3 would be. That may be the reason why high FGFR2 expression did not have significantly lower RFS than low FGFR2 expression in patients without expression of HER3.

An interesting finding of our study was that patients who had recurrence more than 5 years after surgery had a significantly higher probability of having tumors with high FGFR2

Age	Sex	Histology	рТ	pN	pStage	FGFR2 IHC	HER3 IHC	EGFR IHC	Interval between surgery and recurrence (months)
38	F	Diffuse	4a	0	IIB	3+	2+	1+	67
55	F	Diffuse	4a	0	IIB	3+	1+	1+	124
66	F	Diffuse	4a	1	IIIA	3+	0	1+	64
63	Μ	Diffuse	4a	3	IIIC	1+	0	3+	68
66	М	Diffuse	4a	3	IIIC	3+	1+	1+	98

	Table IV. Patients	who had recurrence	e > 5 years after surgery.
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FGFR, fibroblast growth factor receptor; HER, human epidermal growth factor receptor; IHC, immunohistochemistry; EGFR, epidermal growth factor receptor; F, female; M, male.



Figure 6. Numbers of patients who recurred less than 5 years after surgery and more than 5 years after surgery per the FGFR2 IHC grade. Proportion of patients who recurred more than 5 years after surgery is significantly larger in the high FGFR2 expression group (FGFR2 3+) than in the low FGFR2 expression group (FGFR2 1+, 2+) [18% (4/22) vs. 2.9% (1/35), P=0.047]. FGFR, fibroblast growth factor receptor; IHC, immunohistochemistry.

expression. Recurrent sites of long-term failure mostly were peritoneum among diffuse type tumors. The patient with gastric cancer was assumed to be cured at 5 years after surgery unless any signs of recurrence were found. Then, the follow-up was usually terminated. However, we more recently have experienced late recurrence since the introduction of adjuvant chemotherapy using S1. Indeed, 18% of the recurrent patients with high FGFR2 expression recurred more than 5 years after surgery. This would affect the follow-up strategy for gastrectomized patients who underwent S1 adjuvant chemotherapy. A few studies have focused on the timing of recurrence and clinicopathological factors (27-29). Preoperative serum carcinoembryonic antigen level, tumor size, LN metastasis and venous invasion have been reported to be independent predictors of the timing of recurrence. However, no biomarker has been reported to be able to predict the timing of recurrence after surgery. To the best of our knowledge, this finding for the first time indicated that FGFR2 could predict long-term failure of postoperative adjuvant S1 chemotherapy in curative advanced gastric cancer. For patients with long-term failure, FGFR2 IHC level should be examined. If FGFR2 overexpression were confirmed, antibodies or small molecule inhibitors targeting FGFR2 might suppress progression of recurrent tumors.

One patient who recurred more than 5 years after surgery had tumors with FGFR2 IHC status of 1+. Instead, this patient had a tumor with EGFR IHC status of 3+. Moreover, EGFR expression level was negatively correlated with FGFR2 expression. This negative correlation might somewhat be associated with the reciprocal relation of the expressions of FGFR2 and EGFR in these patients. We cannot fully understand the underlying mechanism of this phenomenon, but one plausible reason is that both the EGFR and FGFR2-overexpressing cancer cells would have aggressive nature to cause early failure of S1 adjuvant chemotherapy, while either EGFR or FGFR2-overexpressing cancer cells would have relatively slow growing nature and could evade chemotherapy with the capacity of dormancy (30). That might be the reason why the reciprocal expression between the EGFR and the FGFR2 was found in patients who recurred more than 5 years after surgery.

It has been reported that Helicobacter pylori profoundly activates the epidermal growth factor receptor (EGFR) and its family members HER2 and HER3 (31). We only have checked Helicobacter pylori infection in 37 (22%) patients since we did not check it before 2009. In the analysis of these limited patients, no correlations were found between the Helicobacter infection and EGFR, HER3, or FGFR2 expression (data not shown).

Our study has some important limitations. First, the analysis was based on retrospective data collected at a single center, and the number of included patients was so small that we sometimes could not demonstrate statistically significant difference. Second, no rigid rule was applied in terms of dose reduction and termination of S1 administration. Administered dose of S1 differed between the studied patients. Third, not all the patients were followed up for 10 years, so exact recurrence rate may be different than the observed one.

In conclusion, there was no interaction between FGFR2 expression and patient survival outcomes in patients with stage II/III gastric cancer after the standard treatment in Japan. On the other hand, proportion of patients who recurred more than 5 years after surgery was significantly larger in the high FGFR2 expression group than in the low FGFR2 expression group. Patients with FGFR overexpression in stage II/III gastric cancer should be carefully followed-up for more than 5 years after surgery.

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