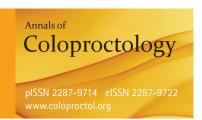
Original Article

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ERCC1 as a Predictive Marker for FOLFOX Chemotherapy in an Adjuvant Setting

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Purpose: The purpose of this study was to identify the excision repair cross-complementation group 1 (ERCC1) as a predictive marker for FOLFOX adjuvant chemotherapy in stages II and III colon cancer patients.

Methods: A total of 166 high risk stages II and III colon cancer patients were retrospectively enrolled in this study, and data were collected prospectively. They underwent a curative resection followed by FOLFOX4 adjuvant chemotherapy. We analyzed ERCC1 expression in the primary colon tumor by using immunohistochemical staining. The oncological outcomes included the 5-year disease-free survival (DFS) rate. The DFS was analyzed by using the Kaplan-Meier method with the log-rank test. A Cox proportional hazard model was used for the prognostic analysis.

Results: ERCC1-positive expression was statistically significant in the older patients (P = 0.032). In the multivariate analysis, the prognostic factors for DFS were female sex (P = 0.016), N stage (P = 0.009), and postoperative carcinoembryonic antigen level (P = 0.001), but ERCC1 expression was not a statistically significant prognostic factor for DFS in the univariate analysis (P = 0.397). The 5-year DFS rate was not significantly associated with the ERCC1 expression in all patients (P = 0.396) or with stage III disease (P = 0.582).

Conclusion: We found that ERCC1 expression was not significantly correlated with the 5-year DFS as reflected by the oncologic outcomes in patients with high-risk stages II and III colon cancer treated with FOLFOX adjuvant chemotherapy.

Keywords: Colon neoplasms; FOLFOX; ERCC1

INTRODUCTION

Colorectal cancer is the second and third most common cancer among Korean men and women, respectively [1]. The 2013 National Comprehensive Cancer Network guidelines recommended adjuvant FOLFOX or XELOX chemotherapy for patients with high-risk stages II and III colon cancer after surgery [2]. The MOSAIC (Multicenter International Study of Oxaliplatin/5-Fluoro-

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uracil/Leucovorin in the Adjuvant Treatment of Colon Cancer) trial reported that the FOLFOX chemotherapeutic regimen, in which oxaliplatin (a third-generation platinum-derivative alkylating agent) was added to 5-fluorouracil (5-FU) and leucovorin (LV), demonstrated a superior disease-free survival (DFS) rate than the LV5FU2 (5-FU and LV regimen) [3]. However, 40% to 50% of postsurgical patients with colorectal cancer eventually experienced recurrence and died of metastatic lesions [4, 5]. Even though the FOLFOX regimen may reduce the risk of recurrence in some patients and increase the survival time, there should be a resistance to oxaliplatin-added chemotherapy in major advanced cases.

Oxaliplatin is a third-generation 1, 2-diaminocyclohexane platinum analogue that causes DNA intrastrand crosslinks that trigger a series of intracellular events that ultimately result in cell death [6, 7]. However, there are several DNA repair systems in cancer cells, such as the base-excision repair, nucleotide-excision repair (NER), mismatch repair, and double-strand-break repair [8]. Many recent

studies reported that the NER capacity may have a major impact on the emergence of resistance to the cytotoxic effect of oxaliplatin [9-13]. Especially, a key player and a rate-limiting enzyme of the NER pathway, excision repair cross-complementation group 1 (ERCC1), is closely related to the risk factors of many patients with cancer, including colorectal cancer [14].

Numerous studies have reported that ERCC1 plays a prognostic role in treatment with oxaliplatin-based chemotherapy in patients with metastatic colorectal cancer [15-19]. However, scarce data were reported in an adjuvant setting. The purpose of this study was to identify the risk factors by using ERCC1 for FOLFOX adjuvant chemotherapy in stages II and III colon cancer patients

METHODS

Subjects

This is a retrospective, single-armed, observational study with analysis of data prospectively collected from the colorectal cancer registry database. A total of 166 patients were enrolled. They were all colon cancer patients who had been treated with FOLFOX4 adjuvant chemotherapy after a curative resection from April 2006 to December 2010. The colon cancer stage was classified in accordance with the 6th edition of the American Joint Committee on Cancer TNM staging system, and patients with high-risk stage II and III cancer were included in the study. The stage II high-risk group was defined as having at least one of the following factors: stage T4a/4b cancer, tumor perforation, bowel obstruction, a poorly differentiated tumor, or venous, perineural, or lymphatic invasion.

Chemotherapy method and follow-up observations

All the patients were treated with chemotherapy after curative surgery. LV at 200 mg/m²/day was administered intravenously for 2 hours. An intravenous bolus of 5-FU at 400 mg/m² was then administered, followed by continuous intravenous administration of 5-FU at 600 mg/m² for the remaining 22 hours. This regimen was continued for 2 days. Oxaliplatin at 85 mg/m² was infused for 2 hours only on day 1. A prophylactic antiemetic and sufficient fluid were infused on days 1 and 2 of chemotherapy. The FOLFOX regimen was administered every 2 weeks. The adjuvant chemotherapeutic regimen was carried out for a total of 12 cycles. Patients were followed up every 3 months for the first 2 years after surgery and every 6 months thereafter for 3 years, for a total follow-up period of 5 years.

History taking, physical examination, and measurement of the serum carcinoembryonic antigen (CEA) level were performed at each follow-up visit. Chest x-ray and abdominopelvic computed tomography examinations were performed to assess the efficacy of chemotherapy every four cycles, and every 6 months after completion of chemotherapy. Colonoscopy was performed annually. Recurrence was identified by imaging studies and colonoscopy and was confirmed by colonoscopic or percutaneous biopsy.

Radiologically-identified tumor growth within the previous surgical field was considered to indicate a recurrence when histological confirmation was not possible.

Tissue microarray and immunohistochemistry

All hematoxylin and eosin-stained slides were reviewed, and the diagnosis of an adenocarcinoma was confirmed. Representative paraffin blocks were selected, and two tissue cores (each 2.0 mm in diameter) obtained from different parts of the tumor were placed into a recipient paraffin block. Immunohistochemical staining was performed by using antihuman ERCC1 antibody (dilution 1:100: E18550, Spring Bioscience, Pleasanton, CA, USA). Four-micrometer-thick tissue sections from the tissue microarray block were incubated in tris-ethylenediaminetetraacetic acid buffer (pH 8.0) and heated at 99°C for 30 minutes. After endogenous peroxidase activity had been quenched with 3% hydrogen peroxide, the sections were treated with a UV inhibitor (Ventana Medical Systems Inc., Tucson, AZ, USA). The sections were incubated with primary antibody at 37°C for 30 minutes and then with HRP-multimer (Ventana Medical Systems Inc.) for 8 minutes. Finally, counterstaining was performed with Mayer's hematoxylin. Staining intensity was graded as 1 for weak, 2 for moderate, or 3 for strong, and the percentage of cells stained was graded as 1 for 0% to <10%, 2 for 10% to <50%, or 3 for 50% to 100%. Staining for ERCC1 was considered to be positive when tumor cells showed nuclear reactivity and both scores were greater than or equal to 2 (Fig. 1) [20].

Statistical analysis

All data were presented as count and percentage for discrete variables. The clinical parameters were compared between ERCC1 groups by using the chi-square test or the Fisher exact test, as appropriate. The DFS was analyzed by using the Kaplan-Meier method with the log-rank test. A Cox proportional hazard model with stepwise selection was conducted to identify risk factors for DFS. P-values of <0.05 were considered to indicate statistical significance. The statistical analysis was performed by using SAS ver. 9.3 (SAS Institute Inc., Cary, NC, USA).

RESULTS

Clinical parameters correlated with ERCC1 expression

The demographic details according to the ERCC1 expression investigated in this study are shown in Table 1. ERCC1-positive expression was detected in 104 patients (ERCC1-positive group), but 62 patients had ERCC1-negative tumors (ERCC1-negative group). The median follow-up time at the time of this analysis was 38.87 months. No significant differences in sex, T stage, N stage, histological differentiation, lymphovascular invasion, neural invasion, or postoperative CEA levels were observed between the two groups. However, ERCC1-positive expression was statistically significant in older patients (P = 0.031). Although not statis-

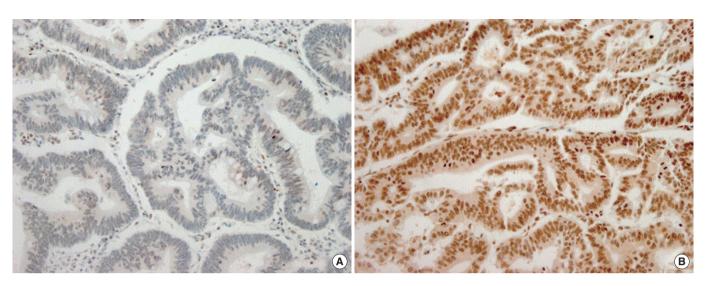


Fig. 1. Immunohistochemical stain for excision repair cross-complementation group 1 (×200): (A) negative (intensity score 1, percentage score 1) and (B) positive (intensity score 3, percentage score 3).

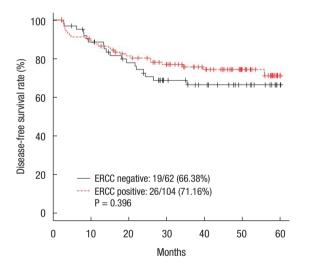


Fig. 2. Five-year disease-free survival rate of all the patients. ERCC, excision repair cross-complementation.

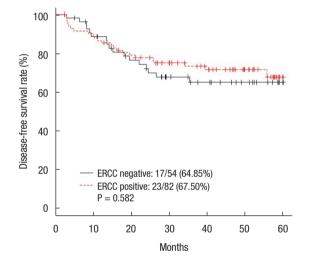


Fig. 3. Five-year disease-free survival rate of stage III patients. ERCC, excision repair cross-complementation.

tically significant, ERCC1 expression exhibited a tendency to be elevated in patients with preoperative high CEA levels (P = 0.095) (Table 1).

In both groups, stage T3 tumors (88.55%) were the most common. Among all patients, stage III tumors (81.93%) were more common than stage II tumors (18.07%). In both groups, moderately differentiated tumors (87.95%) were the most common. On histopathological examination, lymphovascular invasion was seen in 55 (52.88%) and 34 patients (54.84%) in the ERCCI-negative and the ERCC1-positive groups, respectively (Table 1).

Analysis of DFS and prognostic factors

The 5-year DFS rate was 66.38% in the ERCC1-negative group

and 71.16% in the ERCC1-positive group (P = 0.396) (Fig. 2). The 5-year DFS rate for patients with stage III disease was not significantly different between the two groups (64.85% in the negative group and 67.50% in the positive group, P = 0.582) (Fig. 3). The 5-year DFS rate was not significantly associated with the ERCC1 expression in all patients or in those with stage III disease. According to the univariate analysis, the prognostic factors for DFS were female sex (P = 0.047), P = 0.002, elevated preoperative CEA level (P = 0.008), and elevated postoperative CEA level (P = 0.001). In the multivariate analysis, the prognostic factors for DFS were female sex (P = 0.016), P = 0.009, and postoperative CEA level (P = 0.001) (Table 2). However, ERCC1 expression was not a statistically significant prognostic factor for

Table 1. Clinical parameters correlated with ERCC1 expression

Parameter	Negative (n = 62)	Positive (n =	104) P-value
Age (yr)			0.032
≥65	38	80	
<65	24	24	
Sex			0.660
Male	33	59	
Female	29	45	
T stage			0.582
1	1	0	
2	1	3	
3	55	92	
4	5	9	
N stage			0.212
0	8	22	
1	33	42	
2	21	40	
Stage			0.181
II	8	22	
III	54	82	
Differentiation			0.905
Well	3	3	
Moderately	54	92	
Poorly	3	5	
Mucinous cell	2	3	
Signet-ring cell	0	1	
Lymphovascular invasion			0.807
Absent	28	49	
Present	34	55	
Neural invasion			0.394
Absent	37	55	
Present	25	49	
Preoperative CEA level			0.095
<5	41	55	
≥5	21	49	
Postoperative CEA level			0.615
<5	52	84	
≥5	10	20	

ERCC1, excision repair cross-complementation group 1; CEA, carcinoembryonic antinen

DFS in the univariate analysis (P = 0.397) (Table 2).

DISCUSSION

The FOLFOX chemotherapeutic regimen is commonly used to treat colon cancer and has been shown to be an effective adjuvant

Table 2. Univariate and multivariate analyses of prognostic factors for disease-free survival

Classification	Univariate	Multivariate	
	P-value	HR (95% CI)	P-value
Sex			0.016
Female			
Male	0.047	0.48 (0.27-0.87)	
Age (yr)			0.313
<65			
≥65	0.337	1.238 (0.817-1.876)	
Differentiation	0.334		
Well			
Moderately			
Poorly			
Signet-ring cell			
Mucinous cell			
T stage	0.103		
1			
2			
3			
4a			
4b			
N stage	0.002		0.009
0		Reference	
1		1.11 (0.40–3.09)	
2		2.83 (1.08-7.42)	
Lymphovascular invasion			
Absent			
Present	0.478		
Neural invasion			
Absent			
Present	0.475		
Preoperative CEA level			
<5			
≥5	0.008		
Postoperative CEA level			0.001
<5		Reference	
≥5	< 0.001	2.96 (1.59-5.54)	
ERCC1 expression			
Negative			
Positive	0.397		

HR, hazard ratio; Cl, confidence interval; ERCC1, excision repair cross-complementation group 1; CEA, carcinoembryonic antigen.

or palliative chemotherapy [3]. However, treatment failure still exists due to a resistance to oxaliplatin, and many patients treated with this regimen develop long-term side effects such as peripheral neuropathy, as well. Thus, surgical oncologists are investigating predictive markers that can influence the oncological outcomes for patients with colon cancer who receive FOL8FOX chemotherapy. ERCC1 is a gene encoding a protein of the NER complex, which is a group of proteins that are able to repair the DNA damage induced by substances forming adducts, such as platinum [13]. Viguier et al. [17] determined the effect of the ERCC1 codon 118 polymorphism on the tumor response of advanced metastatic colorectal cancer treated by using the FOLFOX regimen. Several studies have also shown that ERCC1 expression can be a prognostic factor for cisplatin- or oxaliplatin-based chemotherapy in patients with non-small-cell lung cancer [21], bladder cancer [22], and gastric cancer [23]. Shirota et al. [15] reported a significant correlation between ERCC1 mRNA levels and the survival of patients with unresectable colorectal cancer who received FOLFOX chemotherapy after failure of FOLFIRI chemotherapy (P < 0.01). Kim et al. [20] analyzed ERCC1 by using immunohistochemistry in 50 patients with colorectal cancer with unresectable metastases that had been treated with the FOLFOX regimen and reported that the median overall survival was significantly longer in patients without ERCC1 expression (P = 0.047). However, our data showed no significant association between the 5-year DFS and ERCC1 expression in patients with high-risk stages II and III colon cancer treated with FOLFOX adjuvant chemotherapy, not even in subgroup analysis with stage III colon cancer.

This study has some notable limitations. First, the sample size was too small to access reliably the association between ERCC1 expression and DFS. Thus, more studies with larger sample sizes are required. Second, the ERCC1 expression level was investigated only though immunohistochemistry without polymerase chain reaction for mRNA expression. Third, no further evaluations were done with thymidylate synthase expression or k-ras status, so this potential prognostic factor must be investigated in future studies.

Interestingly, Li et al. [24] reported a similar outcome to that of the current study. They determined that the mRNA expression level of ERCC1 was not significantly associated with the DFS in Chinese patients with stage II to III colorectal cancer who received 5-FU and oxaliplatin-based adjuvant chemotherapy; neither did they find a significant association between DFS and thymidylate synthase. On the other hand, another study of patients with stage III colon cancer who received oxaliplatin-based adjuvant chemotherapy reported that the 5-year DFS was significantly higher in patients with ERCC1-negative tumors (72%) than in patients with ERCC1-positive tumors (54%) (hazard ratio, 1.98; 95% confidence interval, 1.19–3.31; P = 0.009) [25].

Although ERCC1 expression was not a statistically significant prognostic factor for DFS in this study, we found that lymphnode metastasis and a high postoperative CEA level were significant prognostic factors of 5-year DFS in patients with colon cancer with high-risk stages II and III disease treated with FOLFOX adjuvant chemotherapy. Several studies have demonstrated that lymph-node metastasis could be a prognostic factor for stage III colorectal cancer treated with FOLFOX adjuvant chemotherapy [26, 27]. Additionally, many studies have shown that abnormal postoperative CEA levels are significantly associated with postoperative relapse [28, 29]. Huang et al. [29] reported that a higher postoperative CEA level was a prognostic factor in patients with stage III colorectal cancer who received FOLFOX4 chemotherapy. Therefore, the node stage and the postoperative CEA level should be considered in future studies that perform prognostic analyses of FOLFOX adjuvant chemotherapy for treating patients with colorectal cancer.

In conclusion, we found that ERCC1 expression was not significantly correlated with the 5-year DFS as reflected by the oncologic outcomes in patients with high-risk stages II and III colon cancer treated with FOLFOX adjuvant chemotherapy. Nevertheless, several studies have reported various outcomes regarding the prognostic correlation between ERCC1 expression and FOLFOX chemotherapy in an adjuvant setting. Therefore, a prospective, randomized, controlled trial that includes both ERCC1 and variable biomarkers is needed.

CONFLICT OF INTEREST

No potential conflict of interest relevant to this article was reported.

ACKNOWLEDGMENTS

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