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Transforming growth factor beta can be a parameter of aggressiveness of pT1 colorectal cancer

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Abstract

AIM: To evaluate the significance of transforming growth factor beta (TGF β) expression, in correlation with histopathological parameters, at the front of invasion in T1 colorectal cancer (CRC) and presence of metastases.

METHODS: TGF β immunohistochemical expression was studied in 34 specimens of colorectal adenocarcinomas (pT1). A three-step avidin-biotinylated immuno-peroxidase (ABCu-NCL) staining technique was performed on 4-µm paraffin-embedded tissue sections with a monoclonal antibody to TGF β (Novocastra, NCL-TGFB, clone TGFB 17, dilution 1:40).

RESULTS: Seventeen (50%) out of 34 lesions were positive for TGF β expression. The TGF β -positive rate in patients with vascular invasion was significantly higher than in those without vascular invasion (11/14 cases, *P*<0.01, *P* = 0.005). The TGF β -positive rate was observed in 91.7% of patients with presence of tumor budding at the front of invasion (11/12 cases, *P*<0.01, *P* = 0.0003). A statistically significant correlation was found between the presence of lymph node metastases and positive expression of TGF β (14/16 cases, *P*<0.01, *P* = 0.0001). We also observed that the TGF β -positive rates in groups with distant and non-distant metastases were 92.8% and 20% respectively, and a significant correlation between TGF β expression and distant metastasis was shown (*P*<0.01, *P* = 0.0003).

CONCLUSION: The evaluation of TGF β expression of protein in association with histological parameters can be used as a parameter of the aggressiveness of pT1 CRC.

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Key words: TGF_β; Tumor budding; Colorectal cancer

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INTRODUCTION

Colorectal cancer (CRC) is one of the most common types of cancer in Poland and other western countries. According to statistics, the mortality of CRC is ranking second in western countries and third in Poland amongst all cancers. That is why its biology is still being investigated by researchers^[1-3]. Transforming growth factor β (TGF β) is a multifunctional cytokine that can induce growth inhibition, apoptosis, and differentiation of intestinal epithelial cells^[4,5].

Generally, TGF β inhibits the growth of normal intestinal epithelial cells but can switch to a growth stimulator with tumor progression and enhance malignant transformation^[6,7].

Until now, the number of potential prognostic factors for CRC is continually increasing, which is associated with its biologically diverse malignancy, given the same stage of advancement. Some histological parameters such as vascular invasion and tumor budding at the front of invasion seem to give very important information about invasive potential and metastasis of CRC. A few published studies have associated tumor- budding with metastatic changes^[8-12]. These results seem to confirm that tumor-budding can play a crucial role as a prognostic factor in CRC. Also, the observed presence of neoplastic cells and nets of cells in lymphatic and venous vessels (vascular invasion) is an important feature of invasiveness in CRC. On the other hand, the presence of lymph node and distant metastasis is a recognized parameter of metastatic potential. This is why the aims of the study were, to evaluate the vascular invasion and tumor-budding in CRC growth zone, to analyze its relationship with the expression of TGF β protein in tumors, and to find any correlation between the expression of TGF β and the presence of metastases.

MATERIALS AND METHODS

Collection of samples

Thirty-four T1 colorectal carcinoma specimens from 34 patients, treated by radical surgery at the Department of Surgery, J. Śniadecki Hospital, Białystok, Poland, between 1999 and 2003, were examined retrospectively. Patients' clinical records and pathological reports were reviewed with special attention paid to the presence or absence of lymph node metastasis, local recurrence, and distant metastasis. Tissue specimens were collected immediately after tumor

removal, fixed in 100 g/L formaldehyde, embedded in paraffin, and then histopathologically examined using standard hematoxylin-eosin staining according to the TNM classification.

Vascular invasion, lymphatic and venous invasion were examined and grouped. Tumor budding was examined according to the criteria of Morodomi *et al*¹¹.

Immunohistochemistry

Slides of 4-µm-thick serial sections of the primary tumor were prepared. A standard avidin-biotin immunoperoxidase (Novostain Super ABC Kit (universal), No. NCL-ABCu, Biokom, Poland) method was used for the detection of TGF β expression. Briefly, the slides were dewaxed using xylene, transferred to alcohol, placed in citric acid buffer (10 mmol/L) and heated in a microwave oven (700 W) for 15 min to expose antigens. Endogenous peroxidase activity was inhibited by incubating the section with 3% hydrogen peroxide in methanol for 5 min. The slides were then washed three times with phosphate-buffered saline (PBS) and incubated in 10 g/L normal horse serum for 20 min to reduce nonspecific antibody binding. After washing with PBS, the slides were incubated overnight at 4 °C with monoclonal antibodies. Anti-human TGF β protein monoclonal antibody (Novocastra, NCL-TGFB, clone TGFB 17, dilution 1:40, Biokom, Poland) was used. Nonspecific mouse IgG was used as a negative control. The reaction products were visualized with diaminobenzidine DAB (DAKO \$3000, Dako, Poland). The sections were counterstained with hematoxylin, dehydrated, and mounted. The tissue sections known to be positive were used as positive controls.

Evaluation of samples

The immunostaining of cytoplasm was observed under a light microscope for TGF β . TGF β expression was semiquantitatively assessed in neoplastic cells of primary tumors. Cases were considered positive when >20% of cancerous cells was stained with TGF β and negative when <20% of cancerous cells was stained with TGF β . The percentage of TGF β -positive cells was calculated in at least 500 neoplastic cells per sample, under light microscope (×400). The overall results of immunohistochemical examinations are presented in Tables 1-4.

Table 1	TGF β expression	in main	mass	of CRC	with	and	without
vascular	invasion						

Parameter	n -	Expression of	Total (%)	
rarameter		Negative	Positive	Total (%)
With vascular invasion	14	3 (21.4)	11 (78.6)	41.2
Without vascular invasion	20	14 (70)	6 (30)	58.8

P = 0.005, P < 0.01

Table 2 TGF β expression in main mass of CRC with and without lymph node metastasis

Parameter	n	Expression of	- Total (%)	
i arameter	n ·	Negative	Positive	- 10tai (70)
With lymph node metastasis	16	2 (12.5)	14 (87.5)	47.1
Without lymph node metastasis	18	15 (83.3)	3 (16.7)	52.9

P = 0.0001, P < 0.01.

Table 3 TGF β expression in main mass of CRC with and without distant metastasis

Parameter	п	Expression of	Total (%)	
1 arameter		Negative	Positive	10tai (70)
With distant metastasis	14	1 (7.2)	13 (92.8)	41.2
Without distant metastasis	20	16 (80.0)	4 (20.0)	58.8

P = 0.00003, P < 0.01.

Table 4 TGF β expression in main mass of CRC with and without tumor budding

Dawanaataw	n	Expression of	Total (%)	
Parameter		Negative	Positive	10tal (<i>%</i>)
With tumor budding	12	1 (8.3)	11 (91.7)	35.3
Without tumor budding	22	16 (72.7)	6 (27.3)	64.7

P = 0.00003, P < 0.01.

Statistical analysis

The association between TGF β expression and clinicopathological parameters was examined using χ^2 test. Fisher's exact test was used for statistical analysis. *P*<0.05 was considered statistically significant.

RESULTS

In our study, 17 (50%) out of 34 colorectal carcinomas displayed positive cytoplasmatic TGF β protein reactivity (Figure 1). There was a significant difference between the expression of TGF β in patients with and without vascular invasion (78.6% and 30% respectively, *P*<0.01). A significant difference also existed between the expression of TGF β in primary tumor and lymph node metastases (87.5%, *P*<0.01) and distant metastases (92.8%, *P*<0.01). The presence of tumor- budding at the front of invasion showed a statistically significant correlation with the expression of TGF β in the primary tumor (*P*<0.01).

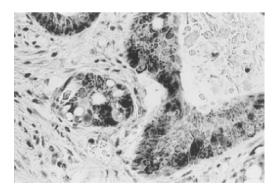


Figure 1 Strong cytoplasmic expression of TGF β in neoplastic cells ×200.

DISCUSSION

Recent studies reported that many tumor cells, including colon cancer cells, can secrete TGF $\beta^{[13,14]}$. It has been shown that elevated levels of human TGF β protein in colorectal

cancer correlate with an increased metastatic potential^[15]. There is a significant correlation between tumor expression of TGF β 1 and a shorter post-operative survival^[16]. It has been reported that plasma concentrations of active total TGF β 1 are significantly higher in patients with CRC than in healthy volunteers^[17]. Patients with CRC in stages C-D have significantly higher expressions of TGF β 1 in tumors ^[16]. Some authors observed that the expression of TGF β 1 is closely related to a higher rate of lymph node metastases in gastric cancer^[18]. Our results also showed that the expression of TGF β in CRC pT1 with lymph node and distant metastasis was higher. All these findings are consistent with the results from Robson et al^[19] who reported a positive expression in 58% of tumors. Similarly, Bellone et al^[20] reported that colon carcinoma progression is associated with gradual and significant increases in the expression of TGF- β 1, TGF- β 2 mRNA and proteins. Tsushima *et al*²¹ showed that preoperative TGF β 1 level is a predictive factor for liver metastasis after curative resection.

The novel histopathological parameter in CRC associated with invasion is known as 'tumor- budding'. Recent studies of the prognostic factors for CRC have paid attention to tumor-budding as a potential prognostic factor^[1,8-12]. We found a statistically significant correlation between the intensity of tumor- budding and lymph node involvement^[2]. Hase et al^[8] examined 663 patients with CRC for tumorbudding, and suggested that tumor- budding is an important prognostic factor in patients with CRC. Morodomi et al^[1] reported that tumor- budding represents the neoplastic cells that are directly involved in host tissue invasion. According to these authors, if the degree of differentiation in colorectal adenocarcinoma is moderate (G2), lymph node involvement is highly probable. However, even in these moderately differentiated tumors, lymphatic involvement is probable but less likely. In well-differentiated adenocarcinomas (G1), tumor-budding and lymphatic invasion are usually not observed. If tumor-budding occurs in such cases, then lymph node involvement is highly probable.

We found that the expression of TGF β in investigated tumors (T1) was strongly correlated with the presence of tumor -budding, vascular invasion at the front of invasion and presence of lymph node and distant metastases. These results suggest that TGF seems to be closely related to the aggressiveness of CRC.

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