Hypoxia-induced up-regulation of angiopoietin-2 in colorectal cancer

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Received September 2, 2005; Accepted November 7, 2005

Abstract. Angiogenesis is a compensatory mechanism that enables malignant tumors to survive in an oxygen-deficient environment. To test our hypothesis that hypoxia stimulates the production of angiopoietin-2 (Ang-2) in colorectal cancer (CRC), we investigated the expression of Ang-2 in three cultured CRC cell lines, and in specimens from 11 CRC metastatic liver tumors. Hypoxia-induced Ang-2 mRNA expression was clearly evident in HCT116 cells that did not express Ang-2 under normoxic conditions. Ang-2 mRNA was detected only after 48 h in hypoxic serum-deprived cultures in a LoVo cell line, and under both normoxic and hypoxic conditions without any noticeable difference in the HT29 cells. There was a stepwise increase in Ang-2 expression from the periphery to the central part of the liver metastatic foci, whereas an inverse result was noted in tumor blood vessels, with a gradual decrease in CD31-positive ECs from the edge to the central region of the metastatic lesion. An expression pattern similar to Ang-2 was found in glucose transporter 1 (Glut-1), a known hypoxia-induced factor. These findings suggest that hypoxia plays an important role in inducing the expression of Ang-2 in CRC.

Introduction

It is generally accepted that CRC growth and metastasis are dependent on neovascularization, a process called ‘angiogenesis’ (1). The angiogenic switch involves various factors and complex processes. There is evidence that angiopoietins (Ang) are implicated in tumor angiogenesis through a shift of balance towards Ang-2, which destabilizes the existing vasculature to a more plastic state (2).

Angiopoietins constitute a novel family of extracellular ligands that bind exclusively to the endothelium-specific tyrosine kinase receptor Tie-2 (3,4). Of the four known Ang molecules (Ang-1, Ang-2, Ang-3, and Ang-4), the best characterized are Ang-1 and Ang-2. Up-regulated expression of Ang-2 has been noted in many human malignancies, including CRC (5-10).

We reported previously that high expression of Ang-2 was significantly more common in metastatic CRC than in normal mucosa and primary CRC, and the expression of Ang-2 increased from the peripheral to intermediate region of the metastatic lesion. Thus, we hypothesized that worsening hypoxia in the center of the metastatic lesion would induce overexpression of Ang-2. To our knowledge, this has not been previously investigated in CRC, and we examined the role of hypoxia on Ang-2 mRNA induction in CRC cell lines and compared its distribution with that of hypoxia, as measured by the distribution of a hypoxia marker, Glut-1, in a CRC metastatic liver tumor. We also assessed the distribution of Ang-2 and Glut-1 in relation to the supply of blood vessels, as measured by antibodies to CD31.

Materials and methods

Clinical samples. We obtained surgical specimens from 11 patients with CRC and synchronous liver metastasis. Written informed consent was obtained from all patients before the study.

Cell lines and culture conditions. The human colon cancer cell lines, HCT116, LoVo, and HT29, were obtained from the American Type Culture Collection (Manassas, VA). These cells were grown in Dulbecco’s modified Eagle’s medium supplemented with 10% fetal bovine serum, 100 units/ml penicillin, and 100 μg/ml streptomycin at 37°C in a humidified incubator with 5% CO2. For cultures under hypoxic conditions, 16 h after the last medium change, the monolayer cultures were grown for 24 or 48 h at 37°C in a continuously monitored atmosphere of 1% O2, 5% CO2, and 94% N2 gas mixture using a multigas incubator, model 9200 (Wakenyaku...
Laser capture microdissection (LCM). LCM was performed in frozen tissue samples using the LM200 LCM system (Arcturus Engineering, Santa Clara, CA) as described previously (10).

RNA extraction and semi-quantitative duplex RT-PCR. Total RNA was extracted from colon cancer cells and clinical samples by a single-step method using Trizol reagent (Life Technologies, Gaithersburg, MD). Complementary DNA (cDNA) was generated using avian myeloblastosis virus reverse transcriptase (Promega, Madison, WI). Semiquantitative analyses for the expression of Ang-2 and Glut-1 mRNA were performed by the duplex RT-PCR techniques, as described previously (10). We used ß-actin as the internal standard. The PCR primer sequences and conditions are shown in Tables I and II.

Reagents and immunohistochemical assay. The anti-human antibodies used in this study were goat polyclonal Ang-2 (Santa Cruz Biotecnology), rabbit polyclonal Glut-1 (Dako, Carpinteria, CA), and mouse monoclonal CD31 (Dako). Immunohistochemical assay of Ang-2, Glut-1 and CD31 was performed using the Vectastain avidin-biotin complex peroxidase kit (Vector Laboratories, Burlingame, CA), as described previously (10).
Statistical analysis. Statistical analysis was performed using StatView J-5.0 software (Abacus Concepts, Berkeley, CA). Medians were compared using the Wilcoxon signed-rank test, and \( p < 0.05 \) was accepted as significant.

Results

Induction of Ang-2 and Glut-1 expression by hypoxia in the colorectal cancer cell lines. RT-PCR analyses did not detect Ang-2 mRNA in the HCT116 or LoVo cell lines under normoxic conditions. Ang-2 mRNA was detected in the HT29 cell line, under normoxic and hypoxic conditions without any noticeable difference. Hypoxia-induced Ang-2 mRNA expression was observed in the HCT116 cell line, whereas in the LoVo cell line, it was only induced after 48 h in hypoxic serum-deprived cultures supplemented with 1% or 0.1% FBS, but not 10% FBS (Fig. 1). The culture of HCT116 and HT29 cells enhanced the expression of Glut-1 mRNA after 24 and 48 h under hypoxic conditions, but not under normoxic conditions (Fig. 2). These experiments were repeated 3 times with reproducible results.

Expression of Ang-2 mRNA and Glut-1 mRNA in the liver metastatic tumors. We measured the levels of Ang-2 mRNA and Glut-1 mRNA by semi-quantitative RT-PCR after LCM. Ang-2 mRNA expression was detected in all 11 liver metastatic lesions, being stronger within the metastatic lesion than in bordering liver tissue (Fig. 3). The ranking intensity of Ang-2 mRNA expression was located in the intermediate area of the metastasis. We found a similar expression pattern in Glut-1 mRNA (Fig. 4A). A stepwise increase in Ang-2 and Glut-1 expression from the adjacent liver to the periphery, then to the intermediate area of the metastatic tumors with increased expression of Ang-2 and Glut-1, was normalized by \( \beta \)-actin (Fig. 4B).

Immunohistochemical findings in the liver metastatic tumors. We observed an increase in the expression of Ang-2 protein from the periphery to intermediate region of the metastatic tumor (Fig. 5). A similar pattern of staining was seen in Glut-1, whereas an inverse result was noted in CD31-positive ECs, with a gradual decrease from the edge to the central region of the metastatic lesion.

Discussion

Our data support the hypothesis that hypoxia induces Ang-2 expression in CRC. Immunohistochemical (IHC) assay showed low vascular density and high Ang-2 expression in the intermediate part of the metastatic lesion near the central area.
of necrosis, in accordance with our previous study (10). Considering the similar expression patterns of Glut-1 and the role of Ang-2 in tumor angiogenesis, it is reasonable to attribute this phenomenon to tissue hypoxia, an environmental factor relating to several tumor biological characteristics including angiogenesis. The presence of necrosis within tumors is thought to be a marker of hypoxia (11) and Glut-1 is well validated as a predictor of hypoxia in CRC (12). In tumor angiogenesis, Ang-2 acts as a destabilizing signal, expressed at the sites of vessel sprouting and regression (13). Thus, we assume that as the tumor enlarges, the hypoxia inside the tumor becomes more severe, which in turn stimulates tumor cells exposed to the most hypoxic conditions to produce and secrete Ang-2.

To ameliorate the hypoxic conditions, the hypoxia-driven Ang-2 induced the ECs of surrounding blood vessels to detach from the pericytes and basement membrane and migrate to the peripheral part of the tumor, resulting in neovascularization in undervascularized areas. Ang-2 was found to stimulate the migration and tubular formation of mouse brain capillary ECs (14). Although it is widely accepted that most tumors and metastases originate as an avascular mass, evidence suggests that tumors in more natural settings may initially grow by co-opting existing host vessels, particularly when they arise within or metastasize to vascularized tissue (15). These co-opted vessels regress, but the remaining tumor is rescued by angiogenesis at its margin. The over-expression of Ang-2 may be a critical regulator in vascular regression (13).

RT-PCR clearly showed that hypoxia is capable of inducing Ang-2 expression in HCT116 CRC cell lines. Hypoxic-induced Ang-2 expression has also been reported in epithelial cells and glioma cells (5,16,17). Unlike HCT116 cells and human glioma cells, in which hypoxic stimulation directly induces Ang-2 (18), a decrease in both the oxygen concentration and the nutrient supply induced Ang-2 expression in the LoVo CRC cell line, which suggests that poor nutrition may be another stimulus of Ang-2 induction. The action of Ang was also found to be cell-specific. Ahmad et al reported that 14 of 18 colon carcinoma cell lines expressed Ang-2, which may explain the expression pattern of Ang-2 in HT29 cells (9). Moreover, the effect of hypoxia on Ang-2 expression varies in different tumor cells. Hypoxic stimuli were not responsible for Ang-2 up-regulation in human hepatocellular carcinoma cells, and down-regulation was seen in renal cell carcinoma cells (19,20).
In summary, the findings of this study suggest that the expression of Ang-2 in CRC is a complex mechanism; however, hypoxia plays an important role in its induction. Moreover, the insufficient blood supply and resultant hypoxia in liver metastatic foci, especially in the intermediate region, might stimulate CRC cells to produce Ang-2.

Acknowledgements

This work was supported by a Grant-in-Aid for Cancer Research from the Ministry of Education, Science, Sports, and Culture Technology, Japan (to H.Y.). This study was also supported in part by a Japan-China Sasakawa Medical Fellowship (to Jinyu Gu).

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