Elevated serum levels of vascular endothelial growth factor is effective as a marker for malnutrition and inflammation in patients with ovarian cancer

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Abstract. Vascular endothelial growth factor (VEGF) reportedly plays an important role in the progression of malignant neoplasms. In the present study, measured serum concentrations of VEGF were measured in patients with ovarian cancer and correlations with nutritional damage and chronic inflammation were analyzed. A significant increase in serum levels in patients compared to healthy volunteers was observed. Levels of VEGF were inversely correlated with serum concentrations of prealbumin, transferrin and retinol-binding protein. VEGF levels were also correlated with serum levels of c-reactive protein (CRP), an effective marker of inflammation. CRP levels were significantly elevated in patients with stage III and IV disease and inversely correlated with serum concentrations of total protein, prealbumin, transferrin and retinol-binding protein. These results demonstrated that an increased production of VEGF correlated with nutritional impairment and inflammation.

Introduction

Ovarian cancer is the most frequent cause of death from gynecological cancer and the fourth most frequent cause of cancer-related death in women worldwide (1). Vascular endothelial growth factor (VEGF) plays a crucial role through its involvement in ovarian biology. Since it is closely associated with the normal function of ovaries, VEGF is also involved in ovarian pathologies, including malignant neoplasms (2). The observation that angiogenesis occurs around a neoplastic tumor was made over a century ago (3). Tumor growth and metastasis were subsequently suggested to depend on angiogenesis and blockage of angiogenesis may thus provide one strategy for inhibiting tumor growth. The participation of angiogenesis and VEGF in the pathogenesis of neoplastic diseases has been previously described (4).

VEGF, previously known as vascular permeability factor, has a mass of 45 kDa and belongs to a family of platelet-derived growth factors. Thus far, several forms of VEGF have been distinguished, including isoforms A, B, C, D and E (5,6). The biological significance of the different forms of VEGF has yet to be definitively determined. Elevated VEGF levels are reportedly associated with advanced-stage melanoma, together with negative immune reactions, including type 2 helper T-cell (Th2) dominance and impaired dendritic cell function (7). Recent studies have reported that immune-suppressing myeloid cells that in increase in number in cancer are expandable by VEGF (8,9).

Cachexia due to cancer is a complex metabolic disorder that includes loss of adipose tissue due to lipolysis, loss of skeletal muscle, elevation of resting energy consumption, anorexia and reduction of oral food intake (10). Acute-phase response proteins including VEGF are reportedly associated with the development of cachexia in patients with cancer (11). The present study investigated the status of VEGF and examined relationships in patients with ovarian cancer between serum levels of VEGF and markers of nutrition and inflammation.

Materials and methods

Sample collection. Blood samples were collected from 27 patients with ovarian cancer. The patient group included 4 patients with stage I disease, 2 with stage II, 13 with stage III and 8 with stage IV disease. The enrolled patients had received surgery or chemotherapy in the Department of Obstetrics and Gynecology at Fukushima Medical University Hospital (Fukushima, Japan) between May, 2011 and August, 2012 and were 38-83 years old (median, 58.5 years) with histological confirmation of the diagnosis. The patients were newly diagnosed and blood samples were collected prior to initiation of any treatment.

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This study was approved by the ethics committee at Fukushima Medical University (2010-2014) and written informed consent was obtained from the subjects enrolled in this study.

Serum levels of VEGF. Peripheral venous blood sera from the subjects were stored at -80°C until use. Serum concentrations of VEGF were measured by enzyme-linked immunosorbent assay (ELISA) (R&D Systems, Minneapolis, MN, USA) according to the manufacturer's instructions.

Markers for nutritional status. Nutritional status was determined by measuring serum concentrations of albumin (nephelometry), prealbumin (turbidimetric immunoassay), retinol-binding protein (latex agglutination immunoassay) and transferrin (turbidimetric immunoassay).

Statistical analysis. Differences between groups were determined using the Student’s t-test. Relationships between two variables were quantified by the Spearman’s rank correlation coefficient. P<0.05 was considered statistically significant. Notably, not all the blood samples were of sufficient volume for the measurements.

Results

We tested sera from 27 patients with ovarian cancer and from 18 healthy volunteers. The serum levels of VEGF in whole

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Results

We tested sera from 27 patients with ovarian cancer and from 18 healthy volunteers. The serum levels of VEGF in whole
Serum levels were significantly increased for whole patients (P<0.01) and stage IV patients (P<0.0001) compared to healthy volunteers (238.5±27.5 pg/ml) (Fig. 1). Serum levels of stage IV patients were significantly higher compared to those of stage II (P<0.00005) and stage III (P<0.05) patients. The serum levels of CRP in stage I, II, III and IV patients were 0.64±0.04, 0.34±0.27, 6.76±1.76 and 4.15±1.92 mg/dl, with significant increases being detected in the serum levels for stage III compared to stage I (P<0.01) and stage II (P<0.01) patients, and for stage IV patients compared to stage I (P<0.05) and stage II (P<0.05) patients (Fig. 2).

These data were analyzed for correlations with parameters of nutritional status and inflammation. VEGF levels showed significant inverse correlations with serum concentrations of prealbumin (r=‑0.467, P<0.005; Fig. 3A), transferrin (r=‑0.640, P<0.00005; Fig. 3B) and retinol-binding protein (r=‑0.457, P<0.005; Fig. 3C). A strong positive correlation was identified with CRP (r=0.422, P<0.01; Fig. 4) and CRP levels showed significant inverse correlations with serum concentrations of total protein (r=‑0.402, P<0.05; Fig. 5A), albumin (r=‑0.668, P<0.0005; Fig. 5B), prealbumin (r=‑0.511, P=0.005; Fig. 5C), transferrin (r=‑0.623, P<0.0005; Fig. 5D) and retinol-binding protein (r=‑0.410, P<0.05; Fig. 5E).

**Discussion**

VEGF is important in the progression of malignant neoplasms (12). In the present study, we investigated the association of VEGF with nutritional damage and systemic inflammation. VEGF levels were significantly higher in patients with ovarian cancer than in healthy volunteers as well as in patients with stage IV patients. VEGF levels showed significant inverse correlations with nutritional status as reflected by prealbumin, transferrin and retinol-binding protein and the marker of inflammation, CRP, which has a strong association with nutritional damage. CRP levels were also significantly elevated in advanced diseases. These results strongly support the hypothesis that VEGF plays a significant role in malnutrition and inflammation, which are essential factors for the progression of ovarian cancer, supporting the possible involvement of VEGF in the pathogenesis of cachexia.

Decreased albumin concentrations are involved with cachexia and are common laboratory features in malignant
diseases. Hypoalbuminemia has been demonstrated to be a predictive factor for poor responsiveness (13,14). The ongoing systemic inflammatory response in terms of CRP has recently gained some interest, as an easily measured and standardized predictor of outcomes in patients after treatment (15,16).

The immunosuppressive properties of malignant tumors have been previously reported (17). Central to this hypothesis is the polarization of the immune system towards a state of inflammation driven by immunological mediators produced by tumor and immune cells (17,18). CRP has been reported as a marker of systemic inflammatory response and an independent predictor of clinical benefit, good prognosis and survival in patients receiving cancer chemotherapy. We recently reported that myeloid-derived suppressor cells (MDSCs) are increased in various types of cancer (19,20) and strong correlations of MDSC with malnutrition were identified in patients with digestive system cancer. Although the exact mechanisms involved in the increased production of immature myeloid cells in cancer patients remain unclear, it has been reported that VEGF is one of the key molecules involved in the induction and expansion of MDSC (21-23). In conclusion, results of this study have demonstrated that an increased production of VEGF correlated with nutritional impairment and systemic inflammation. Future studies should be conducted to investigate possibilities for clinical control of chronic inflammation through the modulation of VEGF.

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References