

Elevated level of some chemokines in plasma of gastric cancer patients

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Abstract

Introduction: Gastric cancer is one of the most common cancer-related causes of death. This is mainly due to the lack of good noninvasive method/biomarkers suitable for early-tumour diagnosis and planning of further therapy modalities. Chemokines play an important role in cancer progression and metastasis formation. In gastric cancer patients, clinical relevance of CXCL12 and CCL5 level has been postulated.

Aim of the study: Efforts were undertaken to examine whether expanded chemokine range may be relevant for evaluation of preoperative staging of gastric cancer patients.

Material and methods: Plasma from 66 gastric cancer patients and 11 healthy controls was obtained, and CCL2, CCL3, CCL4, CCL5, CXCL8, CXCL9, and CXCL10 levels were determined by flow cytometry FlexSet system.

Results: In gastric cancer patients' plasma an increased level of CCL2, CCL4, CCL5, CXCL8, CXCL9, and CXCL10 was observed. In the case of CCL2, CXCL9, and CXCL10, the chemokine levels correlated with advanced (III and IV in TNM classification) disease stage. In the case of CCL4, CCL5, and CXCL8, elevated levels were observed in all cancer patients in comparison to healthy donors.

Conclusions: The accuracy of preoperative diagnosis in gastric cancer may include the monitoring of a wide range of chemokines in patients' plasma. Increased levels of chemokines may warn that the disease is more advanced than conventional diagnostic procedures suggest.

Key words: gastric, cancer, chemokines.

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Introduction

Chemokines are a family of small (8-14kDa) cytokines that induce migration of various cell types [1, 2]. They mediate the biological effect on target cells through widely distributed G-protein-coupled receptors, which are structurally characterised by seven transmembrane spanning domains. Chemokines regulate infiltration of leukocytes to the sites of inflammation, injury, or tumour bed [1, 2]. The availability of chemokines in the site of inflammation is regulated at the level of their biosynthesis, by proteolytic processing, and by mobilisation of the chemokines by cell surface molecules. Chemokines are produced by many cell types in the tumour microenvironment including leukocytes, endothelial cells, fibroblasts, epithelial cells, and cancer cells themselves [3]. In the tumour bed, chemokines from CXCL (CXCL1, 2, 3, 6, 8, 12) and CCL (CCL2, 5, 7) families were previously detected [4, 6]. As a conse-

quence, chemokines are also present in cancer patients' blood.

The biological role of chemokines in tumour progression is important because they promote tumorigenesis and metastasis formation [2]. Chemokines are responsible for modulation of tumour growth by regulation of angiogenesis (e.g. secretion of metalloproteinases [MMP] [4-6]), tumour-specific immune response (e.g. polarisation of tumour-associated macrophages (TAMs) [2, 4]), and stimulation of tumour cell proliferation [7] in an autocrine or paracrine fashion [8].

Gastric cancer is the fourth most common cancer worldwide and one of the major health problem [9]. Surgical resection is still a primary treatment option, although overall survival outcomes remain unsatisfactory. Numerous studies were undertaken to find specific "markers" of gastric cancer progression to improve diagnostic procedure. The most promising data suggests that the production of SDF-1 α in the primary cancer may be such a prognostic

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factor [10]. High CXCR4/high SDF-1 α (CXCL12) expression in tumour cells is significantly associated with depth of cancer invasion, lymph node involvement, and more advanced stage of disease [10]. Also, elevated level of RANTES (CCL5) in serum of gastric cancer patients correlates with poor prognosis [11]. CCL5 is also a candidate biomarker because its measurement is a noninvasive method useful for diagnosis of gastric adenocarcinoma [12].

Aim of the study

The aim of the study was to determine if other chemokines, namely MCP-1 (CCL2), MIP-1 α (CCL3), MIP-1 β (CCL4), IL-8 (CXCL8), MIG (CXCL9), and IP-10 (CXCL10), are present in the plasma of gastric cancer patients (stage I-IV TNM), and if they might be useful as serum/plasma biomarkers to improve accuracy of preoperative staging.

Material and methods

Patients and healthy donors

Sixty-six consecutive patients (median age 66 years, range 40-76 years) with biopsy-proven gastric cancer at different clinical stages were studied (Table 1). All patients underwent a surgical treatment, and the pathological staging was completed according to UICC TNM 7th edition classification. In all patients with the fourth-stage palliative resection was performed. None of the patients received preoperative chemotherapy or chemoradiotherapy. No concomitant active inflammatory diseases were diagnosed. There were no statistically significant differences between patients in stages I+II vs. III+IV regarding age, sex, and tumour type according to the Lauren classification.

The control group consisted of 11 healthy donors (median age 41 years, range 33-61 years). All study participants signed an informed consent form, and the Bioethical Committee of the Jagiellonian University Medical College (KBET/491/B/2003) approved the study protocol.

Determination of chemokine levels

Plasma from gastric cancer patients was collected preoperatively, and the concentrations of chemokines CCL2, CCL3, CCL4, CCL5, CXCL8, CXCL9, and CXCL10 were measured simultaneously using a Flex Set bead array (BD Biosciences, Immunocytometry Systems, San Jose, CA) followed by flow cytometric analysis (FACSCanto, BD Biosciences), as described previously [13]. The Flex Set beads were discriminated in FL-4 and FL-5 fluorescence channels, while the concentration of specified chemokine was determined by the intensity of FL-2 fluorescence. The amount of chemokines was computed by using the respective standard reference curve and FCAP Array software

(BD Biosciences). For all chemokines the detection level was 10 pg/ml.

Statistical analysis

Statistical analysis was performed by nonparametric Mann-Whitney test. Differences were considered significant at p -values < 0.05 .

Results

By the use of bead array assay we detected significantly elevated levels of specified chemokines (CCL2, CCL4, CCL5, CXCL8, CXCL9, and CXCL10; Fig. 1A-F) in the plasma of gastric cancer patients, compared to healthy donors.

In our study, the levels of CCL2, CXCL9, and CCL10 correlated with clinical stage of disease, being highest in more advanced cancer patients (III and IV stage in TNM classification Fig. 1A, E, F). The levels of CCL4 and CCL5 were markedly elevated in the patients' plasma despite their clinical stage (Fig. 1B, C). The level of CXCL8 was significantly elevated in the plasma of patients with II-IV disease stage (Fig. 1D). The level of CCL2 was elevated only in plasma of patients at stage IV (Fig. 1A). In case of CCL3, its level was very low and we did not detect any differences between patients and healthy donors (data not shown). Chemokine levels did not correlate with histological type of cancer, but with the TNM stage.

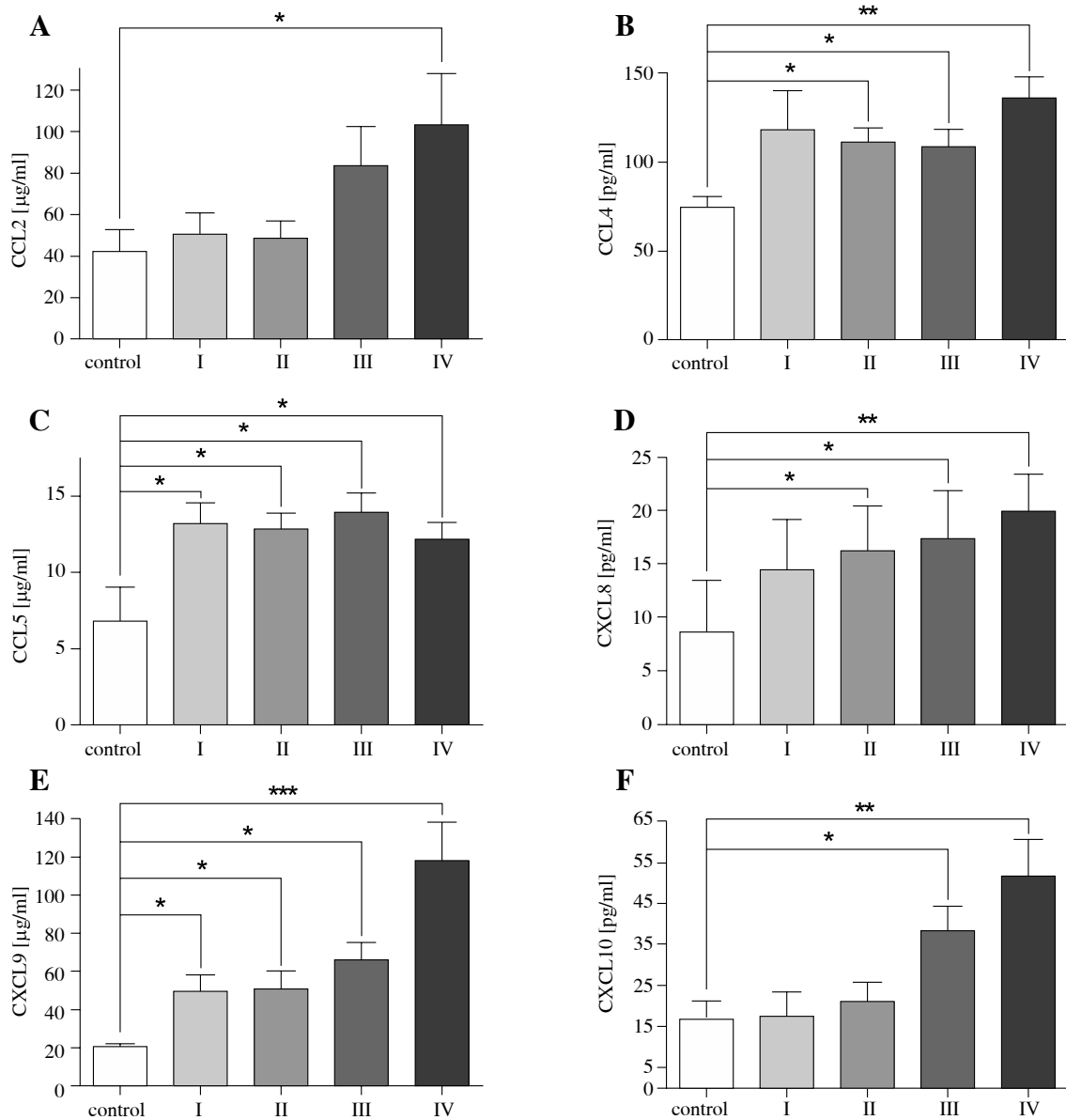
Discussion

The presented study shows that the level of several chemokines in the plasma of gastric cancer patients is elevated when compared to healthy donors. The differences in chemokine levels are related to cancer staging (TNM classification) rather than histological type of tumour (no differences between LI and LII in Lauren classification, data not shown). The elevated level was detected, not only in relation to well documented CCL5 and CXCL12 [11, 14, 15], but also to other chemokines from CC (CCL2, CCL4) and CXC (CXCL8, CXCL9, CXCL10) families.

In our study we observed an increase of CCL5 in plasma of gastric cancer patients, despite their clinical stage. Elevated levels of CCL5 were previously described by other authors [11, 12]. This was also the case in patients

Table 1. Patients' characteristics

Age	66 (40-76) \pm 9.8
Sex	K – 25 / M – 41
Stage I/II/III/IV	I – 11, II – 11, III – 22, IV – 22
Lauren I/II/III	I – 30 / II – 32 / III – 4
Total gastrectomy	56
Subtotal gastrectomy	10



* $p < 0.05$, ** $p < 0.001$ vs. control

Fig. 1. The concentration of chemokines: **A**) CCL2, **B**) CCL4, **C**) CCL5, **D**) CXCL8, **E**) CXCL9, **F**) CXCL10) in plasma from gastric cancer patients (stages are labelled below) and healthy donors (control) determined by FlexSet method. Data are presented as mean \pm SD

with breast cancer [16], where levels of CCL5 and CCL2 correlated with tumour progression [16]. In our study, CCL2 levels were increased, particularly in plasma of patients at stage IV. This corroborates with data published by Jianghong *et al.* [17] indicating that CCL2 may have a predictive value in such a group of patients. CCL2 has proangiogenic activity, and this activity results from macrophage recruitment and activation [18].

Increased levels of CCL3 in the plasma of gastric cancer patients was described for the first time by Rajkumar *et al.* as being involved in gastric cancer tumorigenesis [20]. However, in our data the CCL3 level was low and we did not observe differences between healthy donors and cancer patients. The discrepancies may result from the heterogeneity of the Rajkumar study group reflected in the wide range of CCL3 concentrations.

CCL4 was previously detected in gastric tumour cell cytoplasm (intracellular form). Its expression seems to be dependent on the tumour type (higher expression in non-solid type of poorly differentiated carcinoma) [19]. Herein we have shown that CCL4 is present in the plasma of gastric cancer patients, and its level is significantly elevated compared to healthy donors. This is in line with the data published by Rajkumar *et al.* [20].

Gastric cancer cells may produce CXCL8 both *in vitro* and *in vivo* [5, 21]. We observed elevated levels of CXCL8 in the plasma of all gastric cancer patients, which correlated with the progression of the disease. CXCL8 was previously identified as a strong angiogenic factor in lung, ovarian, prostate, and gastric cancer [22]. CXCL8 may also facilitate tumour growth by inducing chemotaxis of neutrophilic granulocytes, which may further promote angiogenesis, tumour growth, and metastasis formation by releasing metalloproteases (MMPs) and vascular endothelial growth factor (VEGF) [23].

The production of CXCL9, CXCL10, and CXCL11 is induced by other cytokines, including members of the interleukin family and interferons. These chemokines inhibit angiogenesis in colon carcinoma, melanoma, and uterine cervical cancers and are considered as angiostatic chemokines [22]. High levels of CXCL9 and CXCL10 were detected previously by immunohistochemistry in lymphocyte-rich gastric cancers, characterised by increased levels of lymphatic invasion and unfavourable prognosis [24]. CXCL10 and CXCL9 were expressed by peritumorous macrophages, which attract T cells [25]. Interestingly, plasma levels of CXCL9 and CXCL10 decreased significantly in gastric cancer patients after surgery [20].

In summary, the current state of knowledge leads to the conclusion that increased levels of chemokines in plasma may help tumour cells to migrate and invade. In this report we are trying to draw attention to a broad range of chemokines, the levels of which are simultaneously elevated in gastric cancer patients and, in consequence, may serve as adequate indicators of disease stage. Preoperative testing of chemokine levels may help to predict the advance of tumours, including the “biological” potential, which reflects the TNM classification.

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The authors declare no conflict of interest.

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