Association of serum hemoglobin A1c, C-peptide and insulin-like growth factor-1 levels with the occurrence and development of lung cancer

MEIYUN ZHANG, XIAOFENG LI, XINYI ZHANG, YONGYAN YANG, ZHIQING FENG and XIAOYAN LIU

Baotou Cancer Hospital, Baotou, Inner Mongolia 014030, P.R. China

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Abstract. The present study aimed to investigate the correlation of the of hemoglobin A1c (HbA1c), C-peptide and insulin-like growth factor-1 (IGF-1) levels with the development and progression of lung cancer. The serum HbA1c, C-peptide and IGF-1 levels were measured and compared between 80 lung cancer patients and 80 healthy controls; furthermore, their correlation with histopathological type and tumor stage was analyzed in the 80 lung cancer patients. Our results suggested that the levels of HbA1c, C-peptide and IGF-1 were significantly increased in patients with lung cancer compared to those in the control group (P<0.05). In addition, the levels of C-peptide and IGF-1 were significantly higher in the small-cell lung cancer group (n=18), the stage III-IV (n=55) group and the lung cancer with diabetes mellitus group (n=43) compared to those in the non-small-cell lung cancer group (n=62), the stage I-II lung cancer group (n=25) and the lung cancer without diabetes group (n=37), respectively (P<0.05). Thus, the present study suggests that the increased serum HbA1c, C-peptide and IGF-1 levels are significantly correlated with the development and progression of lung cancer.

Introduction

Malignant tumors and diabetes share common characteristics with regards to changes in the cellular and humoral immunity (1). The concentration of hemoglobin A1c (HbA1c) reflects the mean blood glucose levels over the last 8-12 weeks and it is known that long-term high blood sugar levels may cause cell immune-regulatory dysfunction and disturb T-lymphocyte proportions. The concentration of C-peptide, a concomitant of insulin, is increased due to ectopic hormone secretion from the tumor tissues, with a high level detected in the blood of patients with malignant tumors, including liver, colorectal and lung cancer (2,3). Furthermore, the high expression levels of insulin-like growth factor-1 (IGF-1) in patients with diabetes mellitus may induce the overexpression of N-myc mRNA and protein and result in p53 dysfunction, abnormal cell senescence and cancer.

Among several risk factors associated with type II diabetes and lung cancer, lifestyle, diet and environmental factors bear significant similarities, suggesting that diabetes may increase the risk of lung carcinogenesis. In addition, the concentrations of HbA1c, C-peptide and IGF-1 are closely associated with the development and progression of diabetes. An increasing number of studies reported that diabetes mellitus may affect the risk of developing a variety of cancers (4-7); however, the number of studies investigating the association between diabetes and lung cancer is limited. Therefore, the present study aimed to investigate the correlation between the levels of serum HbA1c, C-peptide and IGF-1 and lung carcinogenesis.

Materials and methods

Subjects. The study population included 80 lung cancer patients (49 men and 31 women, with a mean age of 62.1 years) treated at the Baotou Cancer Hospital (Inner Mongolia, China) between 2009 and 2011. The control group comprised 80 healthy subjects (51 men and 29 women, with a mean age of 60.6 years). Subjects with cardiovascular and cerebrovascular diseases, diabetes, high blood pressure, liver, kidney and endocrine diseases on physical examination were excluded from this study. The lung cancer patients were staged according to the 7th edition of the International Staging of Lung Cancer, 2009 (8). The diagnosis was confirmed by histopathological examination and the cases were classified as 62 with non-small-cell lung cancer (NSCLC) (24 with squamous cell carcinoma, 35 with adenocarcinoma, 1 with large-cell carcinoma and 2 with adenosquamous carcinoma) and 18 with small-cell lung cancer (SCLC). Among the 80 lung cancer patients, 25 had stage I-II and 55 had stage III-IV disease. No patients presented with primary lipid metabolism disorders. The diagnostic criteria for diabetes followed the current general international criteria provided by the World Health Organization Diabetes Expert Committee 1999 (9). This study was approved by the Institution Review Board (Baotou, China).
Methods. Blood samples (2 ml) were collected from fasting subjects early in the morning. C-peptide levels were determined by immune chemiluminescence (10), HbA1c testing was conducted with high-pressure liquid chromatography analysis (11) and IGF-1 was measured using the double-antibody sandwich ELISA (12).

Statistical analysis. Statistical analysis was performed using SPSS statistical software, version 13.0 (SPSS, Inc., Chicago, IL, USA). The data are expressed as means ± standard deviation. Significance was assessed using the two-sample Student's t-test for comparison of the corresponding serum indicators between the control and lung cancer groups, the NSCLC and SCLC groups, the stage I-II and stage III-IV lung cancer groups, as well as the lung cancer alone and the lung cancer with diabetes groups. All the statistical tests were two-sided and P<0.05 was considered to indicate a statistically significant difference.

Results

Increased HbA1c, C-peptide and IGF-1 levels in the lung cancer group. The average fasting serum HbA1c was 6.11% in the control group (n=80) and 7.00% in the lung cancer group (n=80). Notably, the mean concentration of HbA1c in the lung cancer group was higher compared to the cut-off HbA1c level of ≥6.5% in the American Diabetes Association Clinical Practice Recommendations for the diagnosis of diabetes (13). The mean HbA1c level in the lung cancer group was significantly higher compared to that in the control group (P<0.05). Moreover, the levels of C-peptide (4.97 ng/ml) and IGF-1 (297.34 µg/ml) in the lung cancer group were significantly elevated compared to those in the control group (3.25 ng/ml and 121.81 µg/ml, respectively; P<0.01) (Table I).

Differences in C-peptide and IGF-1 levels between the SCLC and NSCLC groups. As summarized in Table II, the mean levels of fasting serum C-peptide and IGF-1 were 2.75 ng/ml and 198.73 µg/ml, respectively, in the NSCLC group (n=62). However, the levels of C-peptide and IGF-1 were 6.38 ng/ml and 338.97 µg/ml, respectively, in the SCLC group (n=18). Moreover, the mean C-peptide and IGF-1 levels in the SCLC group were significantly higher compared to those in the NSCLC group (P<0.01), suggesting that C-peptide and IGF-1 may be associated with different pathological types of lung cancer (Table II).

Increased HbA1c, C-peptide and IGF-1 levels in the advanced lung cancer group. The mean serum HbA1c was significantly increased in lung cancer patients with stage III-IV disease (7.58%, n=55) compared to that in patients with stage I-II disease (6.30%, P<0.05) (Table II). In the advanced lung cancer group, the mean fasting serum C-peptide level was 4.50 ng/ml, which was significantly higher compared to that in patients with stage I-II disease (3.03 ng/ml). Furthermore, the average IGF-1 level in the lung cancer group at stage III-IV was 303.41 µg/ml, which was approximately two-fold higher compared to the mean IGF-1 level in lung cancer patients with stage I-II disease (162.93 µg/ml; P<0.01) (Table III).

Table I. HbA1c, C-peptide and IGF-1 levels in patients with lung cancer.

<table>
<thead>
<tr>
<th>Serum indicators</th>
<th>Control group (n=80)</th>
<th>Lung cancer group (n=80)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>HbA1c, %</td>
<td>6.11±0.56</td>
<td>7.00±0.82</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>C-peptide, ng/ml</td>
<td>3.25±0.76</td>
<td>4.97±1.34</td>
<td>&lt;0.01</td>
</tr>
<tr>
<td>IGF-1, µg/ml</td>
<td>121.81±41.54</td>
<td>297.34±89.00</td>
<td>&lt;0.01</td>
</tr>
</tbody>
</table>

Table II. Differences in C-peptide and IGF-1 levels between small-cell and non-small-cell lung cancer.

<table>
<thead>
<tr>
<th>Serum markers</th>
<th>Non-small-cell lung cancer (n=62)</th>
<th>Small-cell lung cancer (n=18)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>C-peptide, ng/ml</td>
<td>2.75±1.87</td>
<td>6.38±1.54</td>
<td>&lt;0.01</td>
</tr>
<tr>
<td>IGF-1, µg/ml</td>
<td>198.73±89.72</td>
<td>338.97±168.34</td>
<td>&lt;0.01</td>
</tr>
</tbody>
</table>

Table III. Increased HbA1c, C-peptide and IGF-1 levels in the advanced lung cancer group.

<table>
<thead>
<tr>
<th>Serum indicators</th>
<th>Stage I-II (n=25)</th>
<th>Stage III-IV (n=55)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>HbA1c, %</td>
<td>6.30±0.46</td>
<td>7.58±0.78</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>C-peptide, ng/ml</td>
<td>3.03±1.75</td>
<td>4.50±1.49</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>IGF-1, µg/ml</td>
<td>162.93±84.84</td>
<td>303.41±112.45</td>
<td>&lt;0.01</td>
</tr>
</tbody>
</table>

Table IV. Differences in C-peptide and IGF-1 levels between lung cancer patients with and those without diabetes mellitus.

<table>
<thead>
<tr>
<th>Indicators</th>
<th>Lung cancer (n=37)</th>
<th>Lung cancer with diabetes mellitus (n=43)</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>C-peptide, ng/ml</td>
<td>2.07±0.83</td>
<td>4.66±1.69</td>
<td>&lt;0.05</td>
</tr>
<tr>
<td>IGF-1, µg/ml</td>
<td>182.28±91.76</td>
<td>349.72±198.88</td>
<td>&lt;0.01</td>
</tr>
</tbody>
</table>

Differences in C-peptide and IGF-1 levels between lung cancer patients with and those without diabetes. As shown in Table IV, the mean C-peptide level in lung cancer patients with diabetes mellitus (n=43) was 4.66 ng/ml, which was significantly higher compared to that in patients with lung cancer without diabetes mellitus (n=37; 2.07 ng/ml). C-peptide is secreted in...
the same amount as insulin and, therefore, accurately reflects the amount of insulin secretion. Therefore, the increased levels of C-peptide may be associated with the development of lung cancer in patients with type II diabetes. In addition, the mean level of IGF-1 was significantly increased in lung cancer patients with diabetes mellitus (349.72 µg/ml) compared to that in patients with lung cancer alone (182.28 µg/ml; P<0.01).

In future studies, we plan to increase the sample size; furthermore, by measuring islet function in patients with lung cancer, monitoring blood glucose, tracking the type and dose of hypoglycemic drugs and perform follow-ups on the survival of lung cancer patients with and without diabetes, we aim to investigate the correlation between diabetes mellitus and the risk of lung cancer.

**Discussion**

It was previously demonstrated that, with increasing levels of HbA1c, the levels of tumor markers CEA, CA199 and CA153 in patients are gradually increased, suggesting that high blood sugar levels may promote cell expression of tumor markers (14). In this study, the HbA1c level was significantly increased in patients with lung cancer, particularly those with advanced-stage disease (stage III-IV), suggesting that HbA1c may be associated and potentially contribute to the development of lung cancer.

Cust et al (15) reported that, in patients with lung cancer and liver metastases, the levels of fasting serum C-peptide were significantly increased. However, a correlation between the changes of C-peptide levels and lung cancer histology was not established. This study demonstrated that C-peptide levels were significantly higher in lung cancer patients compared to those in the control group. As lung cancer progresses from early- to late-stage, the C-peptide levels also increase significantly; the levels of C-peptide were significantly higher in the SCLC compared to that in the NSCLC group, suggesting a significant correlation between fasting serum C-peptide levels and the occurrence and development of lung cancer.

Casa et al (16) demonstrated that the serum IGF-1 levels in patients with NSCLC were significantly decreased following effective chemotherapy and were significantly higher in the ineffective chemotherapy group. The subjects with high IGF-1 levels exhibited a significantly higher risk of cancer development. Blocking the IGF-1 signaling pathway by antisense oligonucleotides or RNA interference technology may significantly inhibit the growth and proliferation of tumor cells. Previous evidence (17,18) demonstrated that the IGF-1 signaling pathway is involved in multiple mechanisms underlying tumorigenesis, mitosis, metastasis, angiogenesis and antiapoptosis, and is also involved in tumor resistance to chemotheray, radiotherapy, human epidermal growth factor receptor 2 and epidermal growth factor receptor-targeted therapy. In the present study, when comparing the lung cancer with the control group, the SCLC with the NSCLC group, the late- with the early-stage lung cancer group and the lung cancer alone with the lung cancer plus diabetes groups, the IGF-1 levels were found to be significantly increased in the study groups. Our results suggested that IGF-1 may be closely associated with the occurrence and development of lung cancer, a finding that may help improve chemotherapy and radiation therapy for lung cancer.

**References**