

TGF- β 1 Serum Concentration as a Complementary Diagnostic Biomarker of Lung Cancer: Establishment of a Cut-Point Value

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Lung cancer is a malignant disease with increasing mortality rates. Cytokines play a role in normal cell growth regulation and differentiation and are also implicated in malignant disease. Among these cytokines, Transforming Growth Factor β type 1 (TGF- β 1) acts as a tumor promoter in malignant cells. Several clinical studies have found high levels of TGF- β 1 in various cancer types. The aim of this study was to establish a TGF- β 1 cut-off point as a complementary diagnostic tool in lung cancer detection. Therefore, 72 clinically well-characterized individuals were studied, 41 lung cancer patients and 31 healthy subjects. Serum TGF- β 1 concentration was measured by an enzyme-linked immunosorbent assay (ELISA). We compared

statistically the serum TGF- β 1 concentration between both groups with analysis of variance, linear regression and receiver operating curve analysis. We observed that lung cancer patients produced higher TGF- β 1 levels than healthy individuals ($37,225 \pm 9,436$ vs. $28,416 \pm 9,324$ pg/ml, $P < 0.001$). The cut-point diagnostic value was 30,500 pg/ml with 80.5% sensitivity, 64.5% specificity and odds ratio: 7.5, 95% CI: 2.6–21.8. *Conclusions:* We found significantly higher TGF- β 1 levels in lung cancer patients than in healthy individuals. We propose the measurement of serum TGF- β 1 levels as a complementary diagnostic test in lung cancer detection. *J. Clin. Lab. Anal.* 25:238–243, 2011. © 2011 Wiley-Liss, Inc.

Key words: TGF- β 1 serum concentration; lung cancer; diagnosis; biomarker

INTRODUCTION

Worldwide lung cancer is a disease with high mortality rates (1). Risk factors like cigarette smoking, wood smoke exposure, and lung cancer family history, among others, have been reported for lung cancer (2). The detection of lung relies on both clinical history and other diagnostic test such as chest radiography and transthoracic needle aspiration (1). Lung cancer is histologically classified as nonsmall cell lung cancer (including adeno-, squamous cell-, and large cell-carcinoma) and Small Cell Lung Cancer. Each histological type requires specific treatment and has different prognosis (1). The biological changes that occur in carcinogenesis, depends on alterations in oncogene activation, tumor gene suppressor inactivation, as well

as alterations in signaling pathways that affect growth factors. Included in these, cytokines play a key role in normal cell growth regulation and differentiation and are involved in many types of malignant disease (3). Transforming Growth Factor β type 1 (TGF- β 1) is a cytokine member of a super-family that includes TGF- β 1 through 5, bone morphogenic proteins, activins,

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and inhibins. Human TGF- β 1 is a 25 kDa, disulfide-linked, nonglycosylated homodimer (4). This cytokine acts as tumor suppressor and arrests cell growth in epithelial normal cells and in cells with an early malignant state (5). In contrast, in an advanced tumoral phase, TGF- β 1 produces a positive effect for survival, progression, and tumor metastasis, promoting epithelium–mesenchymal transition and angiogenesis as well as avoiding immuno-surveillance (5).

Several clinical studies have found high levels of this cytokine in patients with colorectal carcinoma, esophagus carcinoma, gastric carcinoma, glioblastoma, and lung cancer (6–11). Given the increasing incidence of lung cancer and the low effectiveness in the treatment for advanced lung cancer cases, it is important to investigate biomarkers that could complement and improve the diagnostic stage of lung cancer. Therefore, the aim of this study was to quantify TGF- β 1 serum concentration in lung cancer patients and clinically healthy individuals. A positive correlation between high TGF- β 1 serum levels and lung cancer disease was demonstrated. In addition, we found a TGF- β 1 concentration cut-point value with acceptable sensibility and specificity to use as a complementary biomarker in the lung cancer diagnosis.

MATERIAL AND METHODS

Subjects

Eligible cases were identified at the Department of Neumology and Pulmonary Physiology, Centro Médico Nacional de Occidente, Instituto Mexicano del Seguro Social (IMSS), from August 2006 to December 2007. We analyzed clinically and histologically 102 patients suspicious for lung cancer; however, only patients with histopathological confirmation of lung cancer (gold standard diagnosis) were considered in the study ($n = 41$). Patients diagnosed with lung cancer did not receive any treatment, drug, and/or chemotherapy before a blood sample was obtained. Pathological diagnosis was classified according to WHO classification of lung tumors (1), and a staging (TNM classification groups) was performed by an expert pathologist in lung diseases. Also, we studied 157 volunteers, from these we excluded individuals with the following conditions (as these conditions could alter the TGF- β 1 concentration (12–22)): body mass index up to 30 kg/m², alcoholism history, cigarette smoking exposure, wood smoke exposure, chronic hepatic disease, chronic renal disease, acute or chronic respiratory diseases, diabetes mellitus, hypertension, immunodeficiency, patients previously transplanted, patients treated with anti-inflammatory drugs or having any malignant disease. Only individuals

without these criteria were considered as the control group ($n = 31$). All participants were age matched.

Ethic Considerations

The internal Ethics Committee of Centro Médico Nacional de Occidente approved this study. A standardized, structured questionnaire was used to collect clinical data. All participants (patients and controls) gave informed consent.

Sample Collection and TGF- β 1 Measurement

Blood samples were obtained by vein puncture (5 ml), and serum was separated by centrifugation at 3,500 rpm for 15 min. Serum samples were stored at -20°C until TGF- β 1 analysis. TGF- β 1 serum concentration was quantified by enzyme-linked immunosorbent assay (ELISA TGF- β 1 kit, R&D Systems, Minneapolis, MN) according to manufacturer's instructions. Briefly, to activate latent TGF- β 1 to the immunoreactive form, serum samples were acidified and lately neutralized with 1N HCl and 1.2N NaOH/0.5 M HEPES, respectively. 50 μ l of assay diluent was added to each well and subsequently, standard, control, or activated serum samples were added and incubated for 2 hr at room temperature. Each well was aspirated and washed four times with wash buffer. Later, 100 μ l conjugate was added to each well and incubated for 2 hr, and the last washing step was repeated. Subsequently, substrate solution was added to each well and incubated 30 min at room temperature. The reaction was stopped and the plate was read at 450 nm with a λ correction of 540 nm. To calculate TGF- β 1 concentration, the duplicate readings for each standard, control, or activated serum sample were averaged and subtracted to the average zero standard optical density. A standard curve was constructed and the best fit line was determined by regression analysis.

Statistical Analysis

An analysis of variance was used to compare TGF- β 1 serum concentration by several clinical data (age groups, gender, cigarette smoking, wood smoke exposure, signs and symptoms related to poor prognosis). Also, we performed a receiver operating curve (ROC) analysis to estimate a TGF- β 1 level cut-point value and assess its sensibility and specificity as a diagnostic test. These analyses were processed by the statistical software SPSS version 15.0 for WindowsTM (SPSS Inc., Chicago, IL). In addition, we analyzed positive predictive value (PPV) and negative predictive value (NPV) to establish the proportion of patients correctly or incorrectly diagnosed. We also calculated an accuracy determination to

analyze the probability of the test to classify correctly the patients.

Finally, we calculated odds ratio (OR) and a likelihood ratio in order to estimate the risk and to provide a direct estimation of how much this test result would change the probability of having lung cancer. These analyses were processed with the following software: Diagnostic test calculator (version 2006032401) 2002–2007 by Alan Schwartz (alansz@uic.edu), and Graph Pad PRISM version 5.0 for Windows™ (GraphPad Software Inc., La Jolla, CA).

RESULTS

The association between serum TGF- β 1 levels and age, gender, cigarette smoking, wood smoke exposure, and lung cancer family history signs and symptoms related to poor prognosis status was evaluated in healthy individuals and lung cancer patients. The clinical features of both groups are shown in Table 1.

The mean age \pm SD was 64.3 ± 10.6 and 61.5 ± 19.9 years old for the lung cancer and healthy individuals groups, respectively. Other clinical symptoms present at the moment of the diagnostic included cough (16.9%), dyspnea (14.6%), chest pain (11.8%), weight loss (10.6%), haemoptysis (9.6%), weakness (6.5%), anorexia

TABLE 1. Relevant Clinical Features of Lung Cancer Patients and Healthy Individuals

	Lung cancer <i>n</i> = 41 (%)	Control group <i>n</i> = 31 (%)
Age (years, mean age \pm SD)	64.3 ± 10.6	61.5 ± 19.9
Gender		
Male	29 (71)	17 (55)
Female	12 (29)	14 (45)
Wood smoke exposure	8 (20)	0 (0)
Tobacco smoking	30 (73)	0 (0)
Cancer familial history	15 (37)	0 (0)
Cancer Stage ^a		
IIIb	11 (30)	–
IV	26 (70)	–
Signs and symptoms related to poor prognosis		
Weight loss	4 (9.8)	0 (0)
Anorexia	2 (4.8)	0 (0)
Fatigue	3 (7.3)	0 (0)
Histological type		
Adenocarcinoma	16 (39.0)	–
Squamous cell carcinoma	11 (26.9)	–
Small cell carcinoma	4 (9.7)	–
Large cell carcinoma	2 (4.9)	–
Other types ^b	8 (19.5)	–

^aFour patients could not be classified in any stage (TNM staging of lung cancer, 1).

^bTumors which cytological or in a small biopsy specimen do not allow specific differentiation according to the WHO lung tumors classification (1).

(5.2%), and others (24.8%) (data not shown). Similarly, 73% of lung cancer patients were cigarette smokers, 20% were exposed to wood smoke (indoor contamination), and 37% had cancer family history (first-degree relatives). Furthermore, at the time of the diagnosis, 70% of patients with lung cancer presented metastatic disease (stage IV), and 30% had locally advanced disease (stage IIIb). We also studied the anatomical localization of the tumors, because this information is critical to establish a prognosis and treatment. We found that left hemithorax (46.3%) was the most frequent tumor location, followed by right hemithorax (39.0%), mediastinum (12.2%), and diffuse micronodular pattern (2.5%). In regards to TGF- β 1 serum concentration, we observed that lung cancer patients produced a higher concentration than healthy individuals ($37,225 \pm 9,436$ vs. $28,416 \pm 9,324$ pg/ml, $P < 0.001$, Fig. 1). We also analyzed the association of TGF- β 1 serum concentration with age, gender, cigarette smoking, wood smoke exposure, lung cancer family history, histological lung cancer types, and signs and symptoms related to poor prognosis of lung cancer status, but no significant differences were found ($P > 0.05$, data not shown).

In addition, to determine if the TGF- β 1 serum concentration could be useful in the lung cancer diagnosis, we performed a ROC analysis. We found that the data distribution generate a cut-point value of TGF- β 1 concentration: 30,500 pg/ml, with 80.5% sensitivity, 64.5% specificity, and OR 7.5 (CI 95%: 2.6–21.8) (Table 2 and Fig. 2).

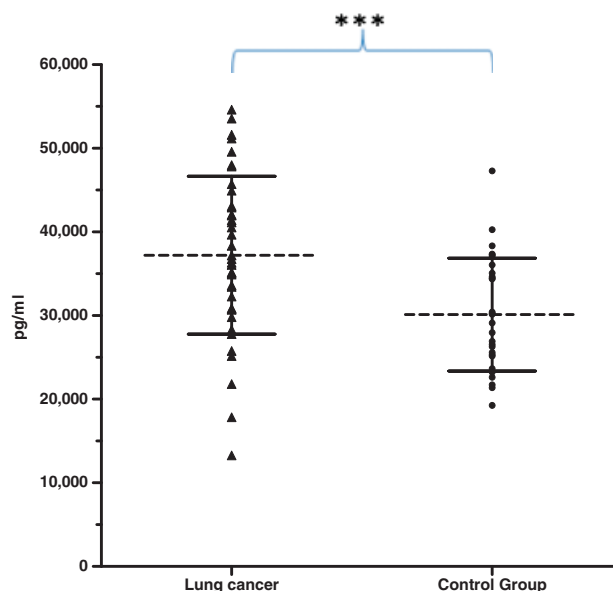


Fig. 1. Distribution of TGF- β 1 serum concentration values (mean \pm SD) in patients with lung cancer ($37,225 \pm 9,436$ pg/ml) and control group ($28,416 \pm 9,324$ pg/ml). *** $P < 0.001$.

TABLE 2. Statistical Parameters for the Use of TGF- β 1 Serum Levels as Diagnostic Test

	Value	95% CI
Sensitivity	80.5%	66.0–89.8%
Specificity	64.5%	46.9–78.9%
Positive predictive value	75.0%	60.6–85.4%
Negative predictive value	71.4%	52.9–84.7%
Accuracy	73.6%	62.4–82.4%
Odds ratio	7.5	2.6–21.8
Likelihood ratio	2.27	1.38–3.73

CI = Confidence interval.

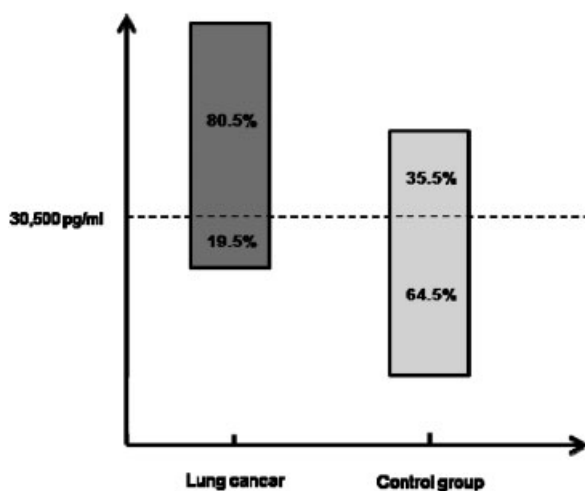


Fig. 2. Sensitivity and specificity of the determination of TGF- β 1 levels in lung cancer patients and control subjects, according to ROC analysis.

After ROC curve evaluation, we also evaluated data with PTP that indicates the proportion of patients with positive test results who are correctly diagnosed. In addition, we tested NPV that is the proportion of patients with negative test results who are correctly diagnosed. Serum TGF- β 1 concentration had PTP values of 75.0% and a NPV of 71.4%. The accuracy of the test was calculated (probability to correctly classify patients) with a value of 73.6%. We calculated a positive likelihood ratio of 2.27 (95% CI: 1.38–3.73) (Table 2).

DISCUSSION

TGF- β 1 has been related to cancer promotion in later cancer stages. In our study, the majority of lung cancer patients (70%) exhibited advanced disease (presence of metastasis) and the rest (30%) locally advanced disease. Accordingly, we found that these patients had a higher TGF- β 1 serum concentration than healthy individuals (control group). Our data is in agreement with data previously reported (10,11). The increase in TGF- β 1 serum concentration could be explained by alterations in

several signaling pathways, including the TGF- β 1 signaling pathway (23) and/or gene expression of cell cycle proteins. Moreover, the microenvironmental changes of tumor cells can promote a high rate of TGF- β 1 production and its activation (5). On the other hand, it has been reported that this cytokine has a dual role in carcinogenesis exerting a tumor suppressor activity (normal cells) as well as promoting cancer progression (malignant cells) (5,24). Its metastatic effect has been demonstrated in in vitro experiments where TGF- β 1 treatment of metastatic breast cancer cell lines enhanced metastases (25). Furthermore, Domagała-Kulawik et al. (26) reported an association between high levels of TGF- β 1 in bronchioalveolar fluid and advanced lung cancer stage that could be related to metastasis. In summary, significantly higher TGF- β 1 production was found in several lung cancer cell lines as well as in lung cancer patients (10,11,27). Kong et al. reported a TGF- β 1 levels reduction after radiotherapy in lung cancer patients proposing this measurement as a marker to monitor disease persistence and recurrence after treatment (10). However, these authors did not find a correlation between TGF- β 1 plasma levels before therapy and histological lung cancer types (11). Similar results were found in this study (Data not shown, P value = 0.846).

These results could be explained by the sample size and the heterogeneous distribution of the histological lung cancer types.

Otherwise, TGF- β 1 expression by tumor cells may inhibit immune response and enhance its tumorigenicity promoting angiogenesis and metastasis. To evaluate the role of TGF- β 1 in the disease progression, we compared the TGF- β 1 concentration with the symptoms related to poor prognosis (anorexia, weight loss, and fatigue) (28) but could not establish a correlation (P value = 0.134).

The current diagnostic strategies have not impacted lung cancer mortality rates (29). As 70% of the lung cancer patients present locally advanced or metastatic disease at the time of diagnosis, new markers and diagnostic methods are needed to detect lung cancer at early stages (1). Currently, sputum cytology, transthoracic needle aspiration, bronchial biopsy, bronchial washing, chest radiography, computed tomography, 18F-FDG Positron Emission Tomography, and/or Magnetic Resonance are the most common diagnostic tests (Table 3). All of these methods have different sensitivities and specificities, its selection depends on tumor localization (central, peripheral, and/or spread disease, Table 3), and its combinatorial use could increase the diagnostic accuracy (30–35). In this study, the measurement of TGF- β 1 concentration yielded enough sensitivity to detect lung cancer (80.5%). The detection of high TGF- β 1 levels provided a risk of 7.5 fold (OR) associated with lung cancer. Besides, we found that if these levels are up to

TABLE 3. Lung Cancer Diagnostic Methods

Diagnostic method	Sensitivity (%)	Specificity (%)	Indication	Comments
<i>Invasive diagnostic test</i>				
Thoracocentesis	80	>90	Pleural spillage	Pleural fluid cytology
Thoracotomy	–	–	Only clearly resectable tumors	Recommended for diagnosis and treatment of early nonsmall cell carcinoma
Excisional biopsy of an accessible node	–	–	Palpable lymphadenopathy	–
Flexible bronchoscopy with or without transbronchial needle aspiration	Central tumors: 88 Peripheral tumors: 60–70	90	Central or peripheral tumors and mediastinal lymphadenopathy	Fluoroscopic or CT guidance; transbronchial needle aspiration improves sensitivity in peripheral tumors
Transthoracic needle aspiration	Peripheral tumors: 90	97	Peripheral tumor in nonsurgical candidates or when transbronchial needle aspiration is inconclusive	Fluoroscopic or CT guidance; the assistance of a cytopathologist improves diagnostic yield
Video-assisted thoracoscopy	–	–	Small peripheral tumors (<2 cm in diameter), pleural tumors, or pleural effusions	May prevent the need for thoracotomy
<i>Noninvasive diagnostic test</i>				
Sputum cytology (at least three specimens)	Central tumors: 71 Peripheral tumors <50	99	Central tumor and haemoptysis	Noninvasive; further testing needed after negative result
Computed tomography	80–90%	–	–	Positive predictive value less than 20% in lung cancer screening
Magnetic resonance imaging	94%	95%	To differentiate malignant SPNs from benign SPNs	To evaluate microvessel density, staging lung cancer and for a treatment followup
Chest radiography	54–84%	90–99%	To detect presymptomatic disease and initial diagnosis	3–5% of lung lesions are identifiable only with lateral X-rays, and 5–17% can be observed better laterally than frontally
¹⁸ F-FDG positron emission tomography	96%	–	Evaluating on indeterminate SPNs	Emerging as a pre-operative assessment in NSCLC. Combined with CT Scan improves accuracy

SPNs, solitary pulmonary nodules. –: Nonspecified.

^aAdapted with permission from “Lung Cancer: Diagnosis and Management,” January 1, 2007, American Family Physician. Copyright © 2007 American Academy of Family Physicians. All rights reserved” Information added from 30, 32–35.

30,500 pg/ml (cut-point value) augment the lung cancer risk. Nevertheless, it is important also consider the clinical condition and lung cancer target symptoms. Actually, to improve the lung cancer algorithm diagnosis is necessary the combination of several clinical tests and markers.

In conclusion, our results indicate that the quantification of TGF- β 1 serum concentration could be useful as complementary diagnostic test, because of the following reasons: easy perform, high sensitivity and specificity, and high cost-effectiveness. In addition, the combination of an expanded number of biomarkers, clinical evaluation, and current diagnostic tests will allow more accurate detection of lung cancer.

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Conflict of interest statement: None of the authors has a financial relationship with a commercial entity that has an interest in the subject of this manuscript.

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