



The clinical significance of serum adipocytokines level in patients with lung cancer

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Background: Adipocytokines were known to play a relevant role in metabolism, inflammation responses and carcinogenesis of several malignancies. Our aims were to detect the expression of serum adipocytokines, explore their potential diagnostic ability and relationship with clinicopathological characteristics of lung cancer.

Methods: Adipocytokines, insulin-like growth factor binding protein 1 (IGFBP-1), resistin, tumor necrosis factors (TNF α), TNF RI and TNF RII, vascular endothelial growth factor (VEGF), leptin, interleukin (IL)-6 and IL-10, chemerin, brain-derived neurotrophic factor (BDNF), plasminogen activator inhibitor-1 (PAI-1) were assessed in 49 untreated lung cancer patients and 20 healthy controls. The protein chip was used to detect the serum levels of adipocytokines.

Results: Lung cancer patients exhibited significantly elevated serum IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6 levels compared to controls ($P < 0.05$) and most of these adipocytokines revealed a modest discriminative ability for the diagnosis of lung cancer, while BDNF were lower in patients ($P < 0.05$). TNF RI was associated with distant metastasis of lung cancer, while there was no relation between other adipocytokines and the patient clinicopathological features.

Conclusions: These results suggest that cytokines IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6 may be involved in the development and progression of lung cancer, and TNF RI may be involved in distant metastasis of lung cancer. Additionally, IGFBP-1, TNF RI, VEGF and TNF RII probably represent potentially useful biomarkers for the diagnosis of lung cancer.

Keywords: Lung cancer; adiponectin; metastasis; tumor necrosis factor RI (TNF RI); biomarker

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Introduction

Lung cancer is the highest morbidity and mortality malignant tumor in the world (1). The etiopathogenesis of lung cancer includes external environmental and inherent decisive factors including Family history, smoking, lifestyle, professional and dietary risk factors (2). Despite substantial advances in comprehensive diagnostic methods and

therapeutic protocols in lung cancer, the 5-year survival rate was less than 20% because of the deficiency of early diagnosis and easy to recurrence and distant metastasis (3). Therefore, it is crucial to study early diagnosis biomarkers and explore new therapeutic modalities of lung cancer.

Obesity is related to an increased risk of developing multiple cancers, such as breast, prostate, endometrial

cancer, and colon cancer (4-6). In obesity, dysfunctional adipose tissue causes metabolic disorders and a diminished ability of fat cells to store free fatty acids, as well as an increased secretion of different adipokines and cytokines which are known associated with tumor development (7). Adipocytes are the main components of adipose tissue, which was regarded as a site of maintain energy homeostasis by storing and mobilizing lipids during times of increased energy demand (8). Lipids are major components of the cell membrane and are essential for malignant tumors. Adipocyte-ovarian cancer cell co-culture identified up-regulation of fatty acid-binding protein 4 (a lipid chaperone) in ovarian cancer cells and promoted *in vitro* and *in vivo* tumor growth, suggesting adipocytes are an energy source of cancer cells (9). In adipocytes and lung cancer cells co-culture system, the results demonstrated a reciprocal interaction between adipocytes and lung cancer cells, which significantly enhanced lung cancer cell proliferation and metastasis (10). Adipocytes were also shown to promote tumor cells proliferation and the direct migration through interactions with certain cancer cells, including prostate, colon, and breast cancer (11-13). In addition, numerous studies have revealed that adipocytes are not only an effective source of energy but also secrete a variety of bioactive substances called adipocytokine, such as leptin, adiponectin, IL-6, VEGF, TNF- α , these secreted factors are able to influence tumor cell survival and behavior (7,14).

Adipocytokines regulate a variety of physiological and pathological processes by acting in autocrine, paracrine or endocrine manner. The accumulating evidence indicates that adipocytokines are associated with the development, progression and immune responses of several malignancies, and are attracting a lot of attention as potential tumor markers (15). In the tumor microenvironment, the binding of PD-1 and PD-L1 inhibits cytokine release and promotes tumor cell escaping from host immune attack (16). PD-L1 overexpression is associated with poor prognosis in non-small cell lung cancer (17). Although lung cancer is not considered to be an obesity-related disease (18), there is a complex interaction between adipose tissue, adipocytokines, insulin resistance and lung cancer pathogenesis (19-21). To date, the interaction between specific adipocytokines and the progression of lung cancer remains unclear and inconclusive. The aim of this study is to detect the expression of serum adipocytokines, explore their potential diagnostic ability and connection with clinicopathological features of lung cancer. In the present study, the serum levels of 12 obesity-associated adipocytokines were studied

and we demonstrate that the serum levels of several of these adipocytokines correlated with the clinicopathological characteristics of lung cancer patients.

Methods

Study population

Forty-nine patients and twenty consecutive healthy controls treated in Department of The Second Affiliated Hospital of Anhui Medical University between July 2016 and May 2017 were enrolled into this prospective study. All cases were newly diagnosed and confirmed by pathological examination as lung cancer. The clinicopathological features (tumor size, pathological type, lymph node and distal metastasis) were determined by physical examination, pathological examination and imaging examinations. Patients who had received chemotherapy or radiotherapy and accompanying other malignancies were excluded from the study. The study was approved by the hospital ethical committee and all participants provided informed consent (approval number: PJ-YX2018-045).

We collected fasting whole blood from forty-nine patients and twenty healthy volunteers into a common centrifuge tubes or vacuum tubes (the blood collection tubes contain no anticoagulant, preservative or separating agent), stored at 4 °C for 30–45 min, centrifuge at 3,000–5,000 rpm/min for 10min, and take the supernatant for detection or stored at –80 °C for further analysis.

Among 49 patients with lung cancer there were 33 males, 16 females; median age: 64, range, 42–77 years. The sex and BMI in healthy controls were resemble to those in lung cancer patient groups, while there are significant differences in ages ($P < 0.01$). Among all the patients were 25 (51.0%) with smoking history, 20 (40.8%) with drinking history and 23 (46.9%) with weight loss. Pathological examination revealed: 30 (61.2%) adenocarcinoma; 14 (28.6%) squamous cell; 5 (10.2%) small cell carcinoma.

Laboratory evaluation

The expression levels of adipocytokines in the fasting whole blood supernatants were measured by Human Obesity Array 3 Glass Slide box (RayBiotech, GA, USA). The experiment was performed according to the manufacturer's instructions. Adipocytokines of different types were detected: the insulin-like growth factor binding protein-1 (IGFBP-1), tumor necrosis factor (TNF α), proangiogenic vascular endothelial growth factor (VEGF), TNF receptor I (TNF RI), resistin,

TNF receptor II (TNF RII), brain-derived neurotrophic factor (BDNF), interleukin-6 (IL-6), plasminogen activator inhibitor-1 (PAI-1), leptin, IL-10, chemerin.

Cytokine assays

Take out the glass slide from the box, and let it air dry for another 1–2 h. Add 100 μ L Sample diluent into each well and incubate at room temperature for 30 min to block slides. Decant buffer from each well. Add 100 μ L of sample to each well. Incubate arrays at room temperature for 1–2 h, wash, decant the samples from each well, and wash 5 times (5 min each) with 150 μ L of 1 \times Wash Buffer I at room temperature with gentle shaking, completely remove wash buffer. Reconstitute the detection antibody by adding 1.4 mL of sample diluent to the tube. Add 80 μ L of the detection antibody cocktail to each well. Incubate at room temperature for 1–2 h, wash. After briefly spinning down, add 1.4 mL of Sample diluent to Cy3 equivalent dye-conjugated streptavidin tube, mix gently. Add 80 μ L Cy3 equivalent dye-conjugated streptavidin to each well. Cover the device with aluminum foil to avoid exposure to light or incubate in dark room. Incubate at room temperature for 1 h, wash, remove water droplets completely, imaging. The raw data were analyzed with GSH-ADI-3 software (Agilent Technologies).

Statistical analyses

All the statistical analyses were performed using SPSS (Statistic Package for Social Sciences) 23.0 and GraphPad Prism 7.0 (GraphPad Software, Inc., La Jolla, CA, USA). Values are presented as the mean \pm standard error of the mean from at least three independent experiments, Student's *t*-test and Chi-square test were used for comparisons between two groups. The cytokine assay raw numerical data was analyzed by RayBiotech software, volcano plot was from the ggplot function. Using receiver operating characteristic (ROC) curves to estimate the diagnostic sensitivity and specificity of cytokine, differences between areas under the curves (AUCs) were assessed by the Wilcoxon test. All tests were two-sided; $P < 0.05$ was considered as being statistically significant.

Results

Patients and healthy volunteers characteristics

The research summarizes the sex, age, body mass index

(BMI) and history of past (history of smoking, history of drinking) of all study participants as well as clinical and pathological features of lung cancer patients (*Table 1*). Compared to patients, healthy cases presented a lower age ($P < 0.01$), while there was no significant difference in BMI and previous history (including the smoking and drinking). Of 49 patients with lung cancer, 5 patients were diagnosed with small cell lung cancer and 44 with non-small cell lung cancer (30 had adenocarcinoma and 14 squamous cell carcinoma). Thirty-two patients (65.3%) were diagnosed with stages I and II, 17 patients (34.7%) stages III and IV, most patients had good performance status (PS) (29 with ECOG PS = 0 and 13 with ECOG PS = 1), while 23 out of 49 patients were classified as weight loss.

Comparison of adipocytokines levels in lung cancer patients and controls

As shown in *Table 2* and *Figure 1*, serum levels in the following 7 of the 12 adipocytokines assayed were found to be statistically different between patients and healthy volunteers. Significantly elevated serum levels of 6 adipocytokines (IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6) ($P < 0.05$) were found in lung cancer patients versus control subjects, whereas the serum BDNF levels were significantly lower in patients than in controls ($P < 0.01$). In addition, no statistical difference was detected between the two groups concerning the serum levels of resistin, leptin, TNF α , IL-10 and chemerin.

Correlation between serum adipocytokine levels and clinicopathological features of lung cancer

We further analyzed the possible association between serum adipocytokines levels and clinicopathological features of patients, and the results are summarized in *Table 3*. The statistical difference in adipocytokine serum levels were detected between groups of patients divided according to histology, tumor size, lymphatic and distant metastasis and TNM classification. Significantly higher levels of circulating TNF RI were found in patients with distant metastasis ($P < 0.05$) (*Figure 2*). We did not find any correlation of the adipocytokines levels with tumor size and lymphatic metastasis of lung cancer patients. There was positive correlated tendency between several adipocytokines levels and patients in TNM classification, but no significant association was

Table 1 The baseline characteristics of lung cancer patients and healthy volunteers included in the study

Baseline characteristics	Patients (N=49)	Healthy (N=20)	P value
Sex (males), n (%)	33 (67.3)	12 (60.0)	0.56
Age, years	64.43±9.73	37.63±11.32	0.00
BMI, kg/m ²	22.72±3.11	23.45±3.37	0.54
Smoker, n (%)	25 (51.0)	6 (30.0)	0.11
Drinker, n (%)	20 (40.8)	9 (45.0)	0.74
Histology, n (%)			-
Adenocarcinoma	30 (61.2)	-	
Squamous cell	14 (28.6)	-	
Small cell carcinoma	5 (10.2)	-	
Stage, n (%)			-
I + II	32 (65.3)	-	
III + IV	17 (34.7)	-	
Weight loss, n (%)	23 (46.9)	-	-
Performance status, n (%)			-
0	29 (59.2)	-	
1	13 (26.5)	-	
2	7 (14.3)	-	

BMI, body mass index.

Table 2 Serum levels of adipocytokines (pg/mL) in lung cancer patients and controls

Adipocytokines	Patients (mean)	Controls (mean)	P value
IGFBP-1	148,214.9	32,188.9	0.00
Resistin	254,917.2	192,906.2	0.06
TNF RI	45,126.3	21,141.4	0.00
VEGF	12,093.9	4,977.0	0.00
TNF RII	50,876.0	33,383.3	0.00
Leptin	11,636.0	19,326.2	0.38
BDNF	49,441.4	83,516.8	0.00
PAI-1	13,266.6	9,322.7	0.01
IL-6	12,404.7	10,374.1	0.01
TNF α	3,547.9	3,597.3	0.97
IL-10	2,725.7	2,360.6	0.43
Chemerin	2,160.6	2,386.8	0.73

P<0.05: the difference was statistically significant. IGFBP-1, insulin-like growth factor binding protein 1; TNF, tumor necrosis factor; VEGF, vascular endothelial growth factor; IL, interleukin; BDNF, brain-derived neurotrophic factor; PAI-1, plasminogen activator inhibitor-1.

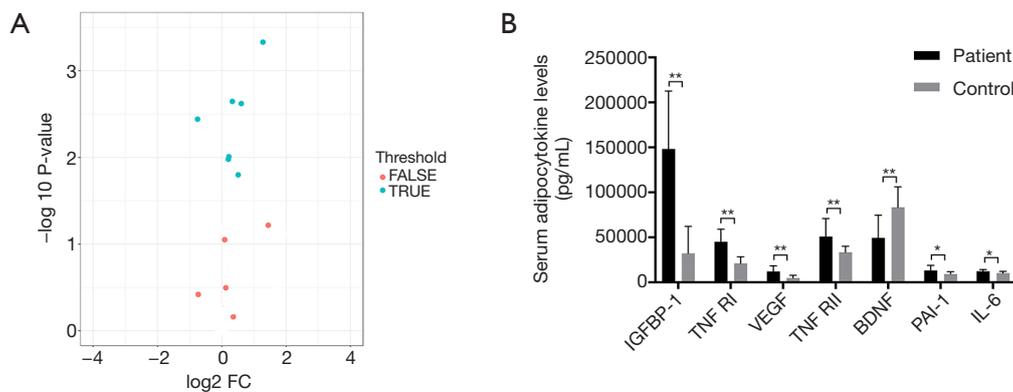


Figure 1 The expression of difference cytokines in patients and controls. (A) Volcano plot. TRUE presented differential expression cytokines. FALSE presented non-differentially expressed cytokines (P value < 0.05 and absolute Log_2 FC > 0.263). To detect differentially expressed factors, we performed the student's t -test (P value) and foldchange on each factor individually between two groups. Differentially expressed factors were defined as those with P value less than 0.05, foldchange over 1.2 and less than 0.83, and fluorescent value over 150. There were 7 differential expression cytokines. Differential expression factors were defined according to P value and foldchange between groups, and presented as volcano plot. (B) The expression levels of 7 differential expression adipocytokines between patients and controls. *, $P < 0.05$; **, $P < 0.01$.

found ($P > 0.05$).

Evaluation of diagnostic sensitivity and specificity of adipocytokines in lung cancer

The ROC curves analysis of different activities cytokines is given in *Figure 3*. The area under the ROC curve of serum IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6 for the diagnosis of lung cancer are 0.961, 0.961, 0.829, 0.803, 0.789 and 0.783 (95% CI: 0.880–1.041, 0.893–1.028, 0.672–0.986, 0.636–0.969, 0.590–0.989 and 0.592–0.974. $P < 0.05$), the best cut-off point (according to Youden index) for circulating adipocytokines levels were 0.32, 0.83, 0.65, 0.72, 0.73 and 0.68 pg/mL, and the sensitivity and specificity of diagnosis of lung cancer were more than 70%.

Discussion

Lung cancer is the leading cause of cancer-related death in worldwide, usually with high metastasis rate and poor prognosis (1). Therefore, it is significant to research the effective biomarkers in diagnosis and prognosis of lung cancer. In the present study, we tested the expression levels of wide adipocytokines serum profile, explore its diagnostic potential and assess its clinical significance of newly diagnosed lung cancer patients. We demonstrated here that several circulating adipocytokines are elevated in the

sera of lung cancer patients compared to controls, namely IGFBP-1, TNF RI, VEGF, TNF RII, BDNF, PAI-1 and IL-6, while BDNF were lower in patients. But only a few of these adipocytokines were of clinical relevance.

Adipocytokines are mainly synthesized and released by adipocytes and have various biological functions that regulate energy balance, lipid metabolism, and participate in inflammatory reactions (22). Numerous studies have demonstrated that adipocytokines are involved in the pathogenesis of several cancer types, including breast cancer (23), prostate cancer (24) and ovarian cancer (9). Moreover, there is accumulating evidence manifest the associativities between adipocytokines and lung cancer, such as leptin and adiponectin which are the most abundant adipocytokines secreted by adipose tissue. Many experts believed that high expression levels of leptin in serum and tissue of tumor patients is considered to promote tumor growth and progression (25–27), especially in lung cancer (20), but these research findings are inconsistent. In my study the leptin levels increase in the patients are not found. The same results were obtained in serum resistin, TNF α , IL-10 and chemerin levels, this may be due to the non-negligible heterogeneity of the studied patients.

By comparing the serum concentrations of adipocytokines between lung cancer patients and healthy controls, the results showed that circulating IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6 presents a significantly elevated

Table 3 Relationship between serum adipocytokines levels (pg/mL) and clinicopathological features of lung cancer

Clinicopathological parameters	IGFBP-1	TNF RI	VEGF	TNF RII	BDNF	PAI-1	IL-6
Histology							
Adenocarcinoma	158,243	47,944	12,449	55,344	55,768	14,434	12,452
Squamous cell	175,741	40,599	12,483	46,102	42,168	12,497	12,135
Small cell carcinoma	99,137	46,050	11,058	49,454	48,045	12,320	12,652
P value	0.24	0.63	0.96	0.82	0.74	0.89	0.73
Tumor size							
≤3 cm	142,571	43,508	12,378	42,374	65,235	16,820	13,377
3–5 cm	150,059	35,629	9,223	48,704	44,592	11,894	11,463
>5 cm	153,325	51,301	13,530	56,805	43,361	12,054	12,238
P value	0.95	0.10	0.48	0.64	0.42	0.65	0.19
Lymphatic metastasis							
Positive	150,259	46,861	11,898	51,247	47,688	13,936	12,194
Negative	140,549	38,617	12,826	49,482	56,015	10,753	13,193
P value	0.83	0.36	0.88	0.94	0.44	0.05	0.61
Distant metastasis							
Yes	151,356	50,511	13,633	53,130	56,360	12,620	12,597
No	143,895	36,783	9,977	47,775	46,193	14,539	12,140
P value	0.05	0.03	0.33	0.10	0.57	0.30	0.27
TNM stage							
I	147,430	30,024	12,086	43,820	56,692	11,614	12,300
II	172,561	49,723	5,652	58,446	32,619	23,094	12,955
III	108,157	37,363	10,084	45,015	38,771	11,836	11,005
IV	151,356	50,511	13,633	53,130	56,360	12,620	12,597
P value	0.31	0.08	0.37	0.16	0.32	0.71	0.07

Median values are given. Using the Kruskal-Wallis test calculate the P values). BMI, body mass index; NS Non-significant. IGFBP-1, insulin-like growth factor binding protein 1; TNF, tumor necrosis factor; VEGF, vascular endothelial growth factor; IL, interleukin; BDNF, brain-derived neurotrophic factor; PAI-1, plasminogen activator inhibitor-1.

in patients with lung cancer, while BDNF were lower in patients. Kimura *et al.* (28). have shown that BDNF and its receptor TrkB are highly expressed in small cell lung cancer tissues, promote cell proliferation and migration through the BDNF/TrkB signaling pathway and that high expression is related to poor prognosis. Although it is not consistent with our experimental results, we also pointed out that the expression of BDNF in small cell lung cancer was higher than non-small cell lung cancer. The difference may be due to only a spot of patients with small cell lung cancer which

were enrolled in this trial and more samples are needed in the future.

Besides, we further analyzed the relationship between these differential cytokines and the clinicopathological features of lung cancer patients, including pathological type, tumor size, lymph node and distant metastasis, and TNM stage. But only TNF RI was found clinically relevant, the expression of TNF RI was positively correlated with distant metastasis, the higher values of TNF RI in patients with advanced lung cancer. This means that TNF RI may

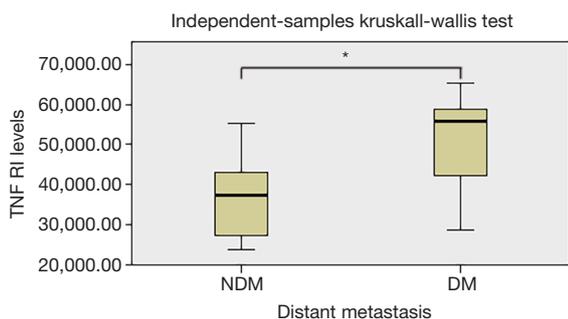


Figure 2 Relationships between distant metastasis of serum concentrations of TNF RI in patients with lung cancer. P values were calculated with the Kruskal-Wallis test. The DM in the abscissa represents the distant metastasis; the NDM in the abscissa represents the non-distant metastasis. *, $P < 0.05$. TNF, tumor necrosis factor.

involve in the development, progression and metastasis of lung cancer.

Metastasis is a complex process and the main cause of cancer-related deaths, especially in lung cancer. It is important to determine the metastasis related markers and potential molecular mechanisms for improving the prognosis of patients with lung cancer. It is well known that TNF- α is one of the factors regulating tumor microenvironment, which has both pro- and anti-tumoral effects through two major receptors (TNF RI and TNF RII) and play important roles in angiogenesis, apoptosis, inflammation and immune response (29). In addition, cytokines that associated with inflammatory responses were usually those most frequently correlated to clinicopathological features of cancers. Our study of the relationship between TNFR1 and the

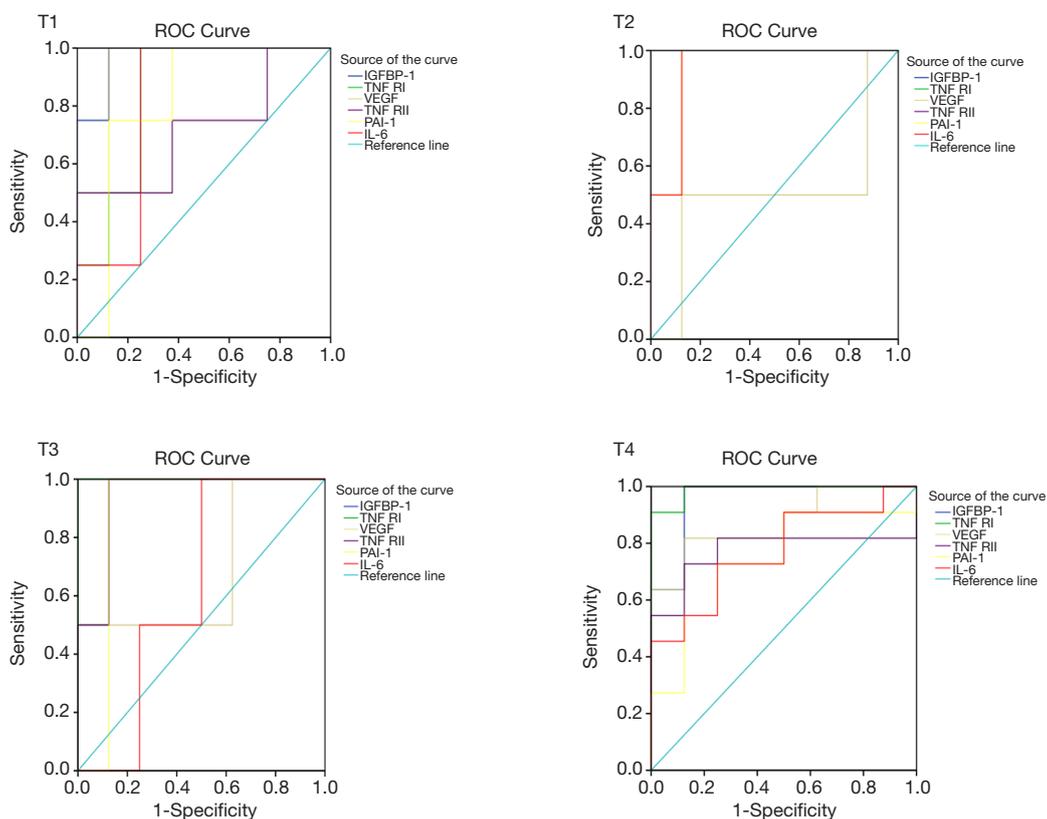


Figure 3 ROC curves for serum difference adipocytokines in untreated patients with lung cancer. Evaluation of sensitivity and specificity of adipocytokines in the diagnosis of lung cancer. The area under the ROC curve of serum IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6 for the diagnosis of lung cancer are 0.961, 0.961, 0.829, 0.803, 0.789 and 0.783 (95% CI: 0.880–1.041, 0.893–1.028, 0.672–0.986, 0.636–0.969, 0.590–0.989 and 0.592–0.974. $P < 0.05$), the best cut-off point (according to Youden index) for circulating adipocytokines levels were 0.32, 0.83, 0.65, 0.72, 0.73 and 0.68 pg/mL, and the sensitivity and specificity of diagnosis of lung cancer were more than 70%. ROC, receiver operating characteristic; IGFBP-1, insulin-like growth factor binding protein 1; TNE, tumor necrosis factor; VEGF, vascular endothelial growth factor; IL, interleukin; BDNF, brain-derived neurotrophic factor; PAI-1, plasminogen activator inhibitor-1.

clinicopathological features of lung cancer is in accordance with this view. Studies on the cell level points out the expression of TNF RI can promote carcinogenesis. TNF RI directly promotes breast cancer cells proliferation by activating p42/p44 MAPK, JNK, PI3-K/Akt pathway (30). On the other hand, TNF RI enhanced cell death induction by CD95L and attenuated the apoptotic effect of TRAIL (31). Moreover, TNF RI is also involved in the metastasis of many malignancies. Charles's experimental results showed that TNF RI promotes peritoneal metastasis of mice ovarian cancer (32). In a colon tumor model, TNF RI-deficient mice cause a reduction in liver metastasis (33). In a word, TNF RI plays an important role in tumor growth and metastasis. Similarly, our experimental results also show that TNF RI is significantly elevated in the serum of lung cancer patients with distant metastasis compared to patients without distant metastases, which means TNF RI may involve in the metastasis of lung cancer and represents a poor prognosis of patients.

The key finding of the research is that the ROC curves point out serum IGFBP-1, TNF RI, VEGF and TNF RII, which present the high sensitivity and specificity, with AUCs of 0.962, 0.961, 0.829 and 0.803, respectively. Diagnostic accuracy of PAI-1 and IL-6 was lower with AUCs of 0.798 and 0.783 ($P < 0.05$, *Figure 2*). In the present study population, there are newly diagnosed and not accompany other diseases and malignancies in all cases. Therefore, it is a good diagnostic method to detect serum levels of these cytokines for newly diagnosed lung cancer patients.

In summary, our study provides evidence that serum levels of several obesity-related cytokines are increased in lung cancer patients (namely IGFBP-1, TNF RI, VEGF, TNF RII, PAI-1 and IL-6), suggest that these cytokines play an important role in the occurrence and development of lung cancer. Moreover, serum levels of TNF RI were correlated to distant metastases of lung cancer and represents a poor prognosis of patients, may be a new target for lung carcinoma therapy. In addition, current results suggest that IGFBP-1, TNF RI, VEGF and TNF RII probably provide certain reference value for early diagnosis of lung cancer.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was approved by the hospital ethical committee (approval number: PJ-YX2018-045) and all participants provided informed consent.

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