



Serum IL-6 as a vital predictor of severe lung cancer

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Background: Recent clinical studies have reported that some cytokines are associated with lung cancer prognosis and mortality. However, the relationship between cytokines and clinical outcomes in severe lung cancer patients was unclear. IL-6 as an important cytokine in inflammation, expression level in severe lung cancer patients was unknown.

Methods: A cohort of 55 severe lung cancer patients were enrolled retrospectively in this study. The clinical characteristics, including performance status (PS), therapeutic effect, and patients' adverse effects, were recorded. The association of cytokines and the concerned clinical outcomes were assessed by logistic regression analysis. The area under the curve (AUC) was assessed to evaluate the strength of prediction.

Results: The mean age of the patients was 59.8, and 42 patients were males. Increased IL-6 levels were associated with worse PS. Logistic regression analysis demonstrated that higher IL-6 was associated with an increased risk of progressive disease (PD) (OR = 1.03, 95% CI: 1.0–1.06). The area under the ROC curve (AUC) of the model used for predicting PD was 0.821.

Conclusions: Increased IL-6 levels are correlated with worse PS and are an essential predictor for PD in severe lung cancer patients. Monitoring the IL-6 level may represent an essential strategy in improving the prognosis of patients with severe lung cancer.

Keywords: Cytokines; severe lung cancer; IL-6, clinical outcomes

Submitted Sep 24, 2020. Accepted for publication Jan 08, 2021.

doi: 10.21037/apm-20-2229

View this article at: <http://dx.doi.org/10.21037/apm-20-2229>

Introduction

Lung cancer is one of the leading causes of cancer death (1) with a high overall death rate (1). The complicated tumor microenvironment in lung cancer, which consists of various immune cells and tumor cells, is widely considered as a vital factor in tumorigenesis and development (2,3). Large

amounts of inflammatory cells and pro-inflammatory factors cytokines (4), such as interleukin (IL), interferon (IFN), and tumor necrosis factor (TNF), play a crucial pro-inflammatory role in cancer progression. The tumor microenvironment provides a role in tumor progression which consist of vasculature, cancer-associated fibroblasts, immune cells, tumor-associated endothelial cells and extracellular matrix (5).

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In the system, IL-6, VEGF were secreted to activate the major pathways on survival, migration and invasion through MAPK, PI3K and PKC. IL-6 regulates all hallmark including DNA damage repair, cell death, tissue invasion, metastasis and angiogenesis. Stromal cells and tumor cells interaction was enhanced by such soluble factors like IL-6 which was mainly secreted by tumor-associated macrophage (TAM) (6).

IL-6 is also a kind of inflammatory cytokine that control of acute inflammation. In cells express IL-6R, classical IL-6 signalling was activated. Free IL-6 binds soluble IL-6 receptor- α which will bind glycoprotein 130 complex and activates the anti-inflammatory pathways. Study discovered IL-6 expression in TAMs of the tumor stroma which is an independent prognostic factor for survival (7).

Specific cytokines may also serve as therapeutic targets of solid cancer (8), such as IL-2 in metastatic renal carcinoma (9,10), and IFN- α in metastatic breast cancer (11) and renal cell carcinoma (12). Notably, a recent clinical trial study (13) demonstrated that IL-1 β inhibitors were associated with lower incidence and lung cancer mortality. Some previous studies have demonstrated that the levels of IL-2 (14) and IL-6 (15) are associated with the prognosis of non-small cell lung cancer (NSCLC).

Cytokines, including IL-1, IL-6, and TNF- α (16) derived from a dysfunctional immune system or tumor is believed to play an essential role in cancer progression. Severe lung cancer here we defined as a specific syndrome of lung cancer patients with critical factors, including worse performance status (PS) scores of Eastern Cooperative Oncology Group (ECOG) (17), or patients with rapid disease progression after treatments, and patients who underwent severe adverse events. Performance status (PS) scores between 3 and 4 mean severe clinical symptoms caused by cancer (17). Previously study mentioned that IL-6 was a prognostic factor in lung cancer (18). However, the relationship between IL-6 in severe lung cancer patients has not been focused. In the present study, we aim to determine the cytokines level in severe lung cancer patients. We present the following article in accordance with the MDAR reporting checklist (available at <http://dx.doi.org/10.21037/apm-20-2229>).

Methods

Patients and data collection

We reviewed patients with a defined lung cancer diagnosis between January 2019 and August 2020 in Xiangya Hospital. We retrospectively enrolled severe lung cancer patients with

critical factors, including lung cancer patients with worse ECOG PS scores, rapid disease progression after treatments, or patients who underwent severe adverse events. Patients with combining autoimmune disease or active infection were excluded. Finally, a cohort of 55 patients was included in the study. Demographics and clinical characteristics, including age, gender, pathology, and clinical stage, were collected. All procedures performed in this study involving human participants were in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by ethics committee of Xiangya Hospital. Individual consent for this retrospective analysis was waived.

Cytokines detection

The inflammation-associated cytokines were detected by enzyme-linked immunosorbent assay (ELISA) using fresh peripheral blood. The detections were performed following the chemiluminescence kit (SIEMENS, Siemens Healthcare Diagnostics Products Ltd.) specification, and the indexes were read on the IMMULITE 1000 Immunoassay System (SIEMENS, Immulite®1000). The cytokines and the reference ranges were as follows: IL-1 β : 0–5 pg/mL, TNF- α : 0–8.1 pg/mL, IL-6: 0–5.9 pg/mL, IL-10: 11.4 pg/mL. The Hospital laboratory detected ESR, CRP, PCT, and lymphocyte subsets.

Clinical outcomes

The patients' performance status (PS) was assessed by the Eastern Cooperative Oncology Group (ECOG) score, also named the WHO or Zubrod score. The scoring criteria divide patients' activity status into 6 levels ranging from 0 to 5: 0, asymptomatic; 1, symptomatic but completely ambulatory; 2, symptomatic, and <50% in bed during the day; 3, symptomatic and >50% in bed; 4, bedbound; and 5, death. The therapeutic effect was divided into four groups: complete release (CR), partial release (PR), stable disease (SD), and progression of the disease (PD), according to Response Evaluation Criteria in Solid Tumors (RECIST) 1.1. Three to four-degree therapy-related adverse effects were obtained, including myositis, hypotension, erosive gastritis, granulocytopenia, severe anemia, and gastrointestinal bleeding.

Statistical analysis

Descriptive analyses were performed with either means \pm standard deviation (continuous variables) or numbers/

Table 1 The demographics and clinical characteristics of patients with severe lung cancer in this study

Characteristic	Overall (n=55)
Age (y)	59.8±8.7
Gender, n (%)	
Male	42 (80.8)
Female	13 (19.2)
Pathology, n (%)	
Adenocarcinoma	25 (45.5)
Squamous cell carcinoma	23 (41.8)
Small cell lung cancer	5 (9.1)
Others	2 (3.6)
Staging, n (%)	
IIB	1 (1.8)
IIIA	4 (7.3)
IIIB	5 (9.1)
IIIC	1 (1.8)
IVA	24 (43.6)
IVB	19 (34.5)
IVC	1 (1.8)

percentages (categorical variables) to describe the patient's characteristics. The categorical variables were compared by Chi-square tests and continuous variables by ANOVA tests. The association of cytokines and the concerned clinical outcomes were assessed by logistic regression analysis. Odds ratios (OR) with 95% confidence intervals (CI) were calculated. The receiver operating characteristic (ROC) curve was calculated from the logistic regression model. The area under the curve (AUC) was assessed to evaluate the strength of prediction. A two-sided P value <0.05 was considered as statistically significant. All statistical analyses were performed using Stata software (version 16.0; StataCorp LLC, College Station, TX, USA).

Results

Baseline characteristics of lung cancer patients

The demographic and clinical characteristics of severe lung cancer patients enrolled in this study are present in *Table 1*. The

Table 2 Association between cytokines and performance status

ECOG score	PS		P value
	0–2	3–4	
IL-6, n (%)			0.012
Normal	35 (71.4)	10 (28.6)	
High	14 (40.0)	15 (60.0)	
TNF- α , n (%)			0.109
Normal	48 (68.5)	22 (31.5)	
High	1 (25.0)	3 (75.0)	
IL-1 β , n (%)			0.237
Normal	36 (63.1)	21 (36.9)	
High	13 (76.4)	4 (23.6)	
ESR/CRP/PCT, n (%)			0.5
Normal	23 (67.6)	11 (32.4)	
High	26 (65.0)	14 (35.0)	
Th/Ts, n (%)			0.51
Normal	49 (72.0)	19 (28.0)	
High	4 (66.7)	2 (33.3)	

ECOG, Eastern Cooperative Oncology Group; PS, performance status; IL, interleukin; TNF- α , tumor necrosis factor- α ; CD, leukocyte differentiation antigen; Ts, suppressor T-cell; Th, helper T-cell.

mean age was 59.8 years ranging from 38 to 74, and 80.8% of the patients were male. The pathologic type was distributed as adenocarcinoma 45.5%, squamous cell carcinoma 41.8%, small cell lung cancer 9.1%, and 3.6%. The clinical-stage was distributed as 18.2% for stage III and 79.9% for stage IV. Of 74 cytokines expression tests, 39.1% (29/74) showed a high expression level of IL-6 and 5.4% (4/74) with an elevated level of TNF- α . The average levels of CRP, ESR, and PCT were 12.3 mg/L, 53.5 mm/h, 0.1 ng/mL, respectively.

Association between cytokines and performance status

The association of cytokines and PS is shown in *Table 2*. Patients with ECOG score 0–2, 35 (71.4%) had normal expression of IL-6. In patients with ECOG score 3–4, 15 (60%) patients had a high expression of IL-6 (P=0.012). Other detected cytokines were not associated with PS. In patients with worse PS scores, only 3 in 22 patients had high TNF- α levels (P=0.1). Differences were not seen in IL-1 β , ESR/CRP/PCT, and Th/Ts (P=0.237/0.5/0.51).

Table 3 Comparisons of the demographic, clinical characteristics, and cytokines, by different treatment effects

Characteristic	CR (n=2)	PR (n=23)	SD (n=26)	PD (n=23)	P	P trend
Age (y)	54.5±23.3	58.0±8.4	63.4±8.2	59.4±7.5	0.294	0.682
Male	1 (50%)	14 (87.5%)	13 (100%)	14 (66.7%)	0.058	0.293
Staging					0.002*	0.001*
IIB	0	1 (5.6%)	0	0		
IIIA	2 (100%)	2 (11.1%)	0	0		
IIIB	0	3 (16.7%)	1 (7.1%)	1 (4.8%)		
IIIC	0	0	1 (7.1%)	0		
IVA	0	8 (44.4%)	7 (50.0%)	9 (42.9%)		
IVB	0	4 (22.2%)	4 (28.6%)	11 (52.4%)		
IVC	0	0	1 (7.1%)	0		
Metastasis	0	7 (38.9%)	7 (5%)	14 (66.7%)	0.157	0.030*
IL-6	3.1 (2.1–4.0)	2.7 (2.0–4.6)	9.0 (3.5–18.8)	9.5 (5.0–27.0)	0.001*	<0.001*
TNF- α	6.3 (5.0–7.6)	9.3 (6.0–11.3)	9.7 (6.9–12.3)	9.2 (7.0–13.0)	0.421	0.141
IL-1 β	5.0 (5.0–5.0)	5.0 (5.0–7.0)	5.0 (5.0–5.0)	5.0 (5.0–6.0)	0.719	0.754
IL-10	5.0 (5.0–5.0)	5.0 (5.0–5.0)	5.0 (5.0–5.0)	5.0 (5.0–5.0)	0.535	0.186
CD3	73.3±9.5	70.1±7.9	68.0±12.7	68.0±11.0	0.819	0.459
CD3CD4%	44.9±3.0	39.6±7.7	38.1±9.7	37.6±9.1	0.648	0.170
CD3CD8%	26.4±5.2	27.0±7.4	26.6±8.6	27.0±8.1	0.996	0.921
Th/Ts	1.7 (1.5–1.9)	1.4 (1.1–2.0)	1.2 (1.0–2.2)	1.3 (1.0–1.9)	0.897	0.562

*, IL-6 was higher in patients with SD and PD compared with CR and PR patients. $P<0.001$. IL, interleukin; TNF- α , tumor necrosis factor- α ; CD, leukocyte differentiation antigen; Ts, suppressor T-cell; Th, helper T-cell; CR, complete response; PR, partial response; PD, progressive disease; SD, stable disease.

Characteristics of study cohort by therapeutic effect classification

The comparison of demographic, clinical characteristics, and cytokines, by different treatment effects, are shown in *Table 3*. The stage IVA and IVB of PD patients were 42.9% and 52.4%, respectively. Patients with IIB and IIIA showed no progress. Most PR and SD patients were IV stage patients ($P=0.002$). The expression of IL-6 was higher in patients with SD and PD when compared to patients with CR and PR ($P=0.001$). The difference was not observed in age, gender, metastasis status, and other cytokines.

IL-6 as a predictor of progressive disease (PD) in severe lung cancer

The logistic regression analysis results for the association

of inflammatory cytokines and PD are presented in *Table 4*. Univariate analysis revealed that increasing IL-6 levels is associated with a higher risk of PD (OR =1.03, 95% CI: 1.00–1.06), while other inflammatory cytokines were not associated with PD in severe lung cancer. The predictive performance of the IL-6 was evaluated by ROC curve analysis, with a total of 74 evaluations. The area under the ROC curve (AUC) of the prediction model was 0.821 (*Figure 1*).

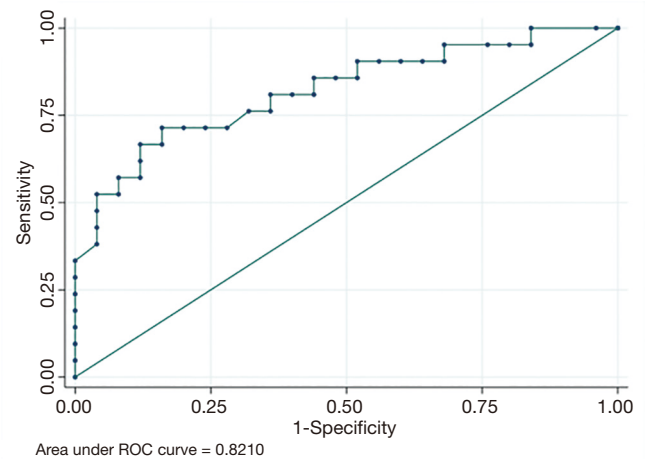
Association of cytokines and therapy-related adverse effect

The association of cytokines and the incidence of therapy-related adverse effects were assessed (*Table 5*). The total adverse event incident rate was 15%, and nine cases of SAE were observed, including two myositis, one hypotension, one erosive gastritis, three granulocytopenia, one severe

Table 4 The association of inflammatory cytokines and the progression of the disease

Variable	OR	95% CI	P value
CRP	1.04	0.97–1.11	0.276
ESR	1.02	0.99–1.04	0.100
IL-6	1.03	1.00–1.06	0.039
TNF- α	1.04	0.06–1.13	0.363
IL-1 β	1.00	0.99–1.02	0.530
IL-10	1.05	0.94–1.16	0.395
CD3	0.99	0.95–1.02	0.438
CD3CD4%	0.975	0.930–1.020	0.315
CD3CD8%	1.000	0.959–1.050	0.990
Th/Ts	0.940	0.567–1.557	0.810

IL, interleukin; TNF- α , tumor necrosis factor- α ; CD, leukocyte differentiation antigen; Ts, suppressor T-cell; Th, helper T-cell; CRP, C-reactive protein; ESR, erythrocyte sedimentation rate.

**Figure 1** ROC curve for IL-6 to predict the progression of disease in severe lung cancer patients.**Table 5** Association of cytokines and therapy related adverse effect

Adverse events	Total (N=74)	High TNF- α (n=30)	Normal TNF- α (n=44)	High IL-1 β (n=9)	High IL-1 β (n=65)	High IL-6 (n=40)	Normal IL-6 (n=34)
All events	10 (15%)	7 (23.3%)	2 (4.5%)	2 (22.2%)	8 (12.3%)	6 (15%)	4 (11.7%)
Myositis	2	0	1	0	2	1	1
Hypotension	1	1	0	1	0	1	0
Erosive gastritis	1	1	0	0	1	0	1
Granulocytopenia	3	2	1	0	3	2	1
Severe anemia	1	1	0	0	1	0	1
Gastrointestinal bleeding	1	1	0	0	1	1	0
Interstitial lung disease	1	1	0	1	0	1	0

anemia, and one gastrointestinal bleeding. The adverse event incident rates in high/normal TNF- α and IL-1 β patients were 23.3%/4.5% and 22.2%/12.3%, respectively; however, the rate of high/normal IL-6 patients was 15%/11.7%.

Discussion

There were a few markers which indicate the status of lung cancer patients. The balance between cancer cells and humoral/cellular immunity can be evaluated. Tocilizumab was reported to treat Cytokine Release Syndrome (CRS)

with fevers and multiorgan dysfunction caused by cancer immunotherapies (19,20). We noticed that some patients PS score deteriorate besides acute infection, most of them had high inflammatory status.

In the study, we evaluated whether cytokines are predictors of clinical outcomes in severe lung cancer patients. IL-1 β was secreted by T lymphocyte, ESR/CRP/PCT were usually correlated with infection of inflammation which might trigger signal activation. Th/Ts reflect the balance of cellular and humoral immunity. IL-6 is one of the most important cytokines. Elevated IL-6 was observed in patients with infective disease as SARS and COVID-19 were

believed related to T lymphocyte hyperactivated leading to vascular dysfunction and pulmonary damage (19). Several studies have reported the predictive value of IL-6 for the prognosis of different cancers. Lee *et al.* reported that IL-6 was a prognostic indicator for hepatocellular carcinoma in patients treated with transarterial chemoembolization among cytokines (21). An experimental study found that overexpression of IL-6 in colon tumor cells significantly increased tumor growth *in vivo* of colon tumors (22). Osuala *et al.* found that IL-6 signaling between preinvasive ductal carcinoma in situ cells (DCIS) and stromal CAFs represent an important factor in the initiation of DCIS progression to invasive breast carcinoma (23). IL-6 also reportedly promoted metastasis of non-small-cell lung cancer by up-regulating TIM-4 via NF- κ B (24). Our study discovered that IL-6 serves as an important biomarker to predict severe lung cancer's poor outcome.

We identified that patients with high IL-6 were at high risk of worse PS and PD compared to those with normal levels of IL-6. In a previous study, IL-6 levels were deemed as a biomarker to assess the activity of irAEs (19); however, in our study, the predictive value of IL-6 in AEs of severe lung cancer was not obvious. The correlation between IL-6 and severe lung cancer indicates it may be a potential therapeutic target. Tocilizumab is a recombinant humanized monoclonal anti-IL-6R antibody. It binds both soluble and membrane-bound IL-6R to inhibit IL-6-mediated signaling. Tocilizumab has been approved by The US Food and Drug Administration for the treatment of severe CAR T-cell-induced CRS (20). If IL-6 was elevated in severe lung cancer with worse PS or rapid PD, we should evaluate its role in treatment decisions, particularly in anti-IL-6 treatment like Tocilizumab.

In our data, there were no significant difference in IL-1 β , ESR/CRP/PCT, and Th/Ts in different performance status. The study revealed that TNF- α , IL-1 β , IL-10, lymphocyte subgroup had no relation with therapeutic effect.

Besides IL-6, other cytokines were reported to be related to the prognosis of different cancers. Lima *et al.* enrolled 74 patients (weight stable cancer n=31 and cachectic cancer n=43) diagnosed with colorectal cancer (CRC) and observed higher protein expression of inflammatory cytokines and growth factors in the tumor and serum of cachectic cancer patients when compared to weight-stable counterparts, such as epidermal growth factor, granulocyte-macrophage colony-stimulating factor, interferon- α , and interleukin (IL)-8 (25). Camargo *et al.* reported that NF- κ Bp65 and its target genes expression (TNF- α , IL-1 β ,

MCP-1, and I κ B- α) were significantly higher in cachectic cancer patients, which proved the function of white adipose tissue in inflammation (26). Martins *et al.* found that an increased level of IL-4 was associated with longer survival time in females with mammary tumors (27). IL-38 is a reliable and sensitive biomarker for distinguishing between CRC and non-cancer colonic tissue. There is a positive correlation between colonic IL-38 in CRC prognosis, particularly in advanced CRC. This supports IL-38 as a reliable and consistent independent factor in predicting CRC prognosis (28). However, we found no significant association between other cytokines and clinical outcomes, including TNF- α , IL-1 β , and IL-10. Additional cytokines should be studied in the future.

In conclusion, we determined that in severe lung cancer patients, a high level of IL-6 was associated with worse PS and PD. Also, IL-6 may act as an important predictor of PD in severe lung cancer patients. IL-6 is relatively simple to detect and interpret, and monitoring the IL-6 level may represent an important strategy to improve the prognosis of patients with severe lung cancer.

Acknowledgments

Funding: This investigation was supported by the National Multidisciplinary Cooperative Diagnosis and Treatment Capacity Building Project for Major Diseases (Lung Cancer), Natural Science Foundation of China (81903020), and China Postdoctoral Science Foundation (2019M652812).

Footnote

Reporting Checklist: The authors have completed the MDAR checklist. Available at <http://dx.doi.org/10.21037/apm-20-2229>

Data Sharing Statement: Available at <http://dx.doi.org/10.21037/apm-20-2229>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/apm-20-2229>). 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. All procedures performed in this study involving human participants were in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by ethics committee of Xiangya Hospital. Individual consent for this retrospective analysis was waived.

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- (English Language Editors: L. Gray and J. Chapnick)

Cite this article as: An J, Gu Q, Cao L, Yang H, Deng P, Hu C, Li M. Serum IL-6 as a vital predictor of severe lung cancer. *Ann Palliat Med* 2021;10(1):202-209. doi: 10.21037/apm-20-2229