

# Elevated levels of total cholesterol and high-density lipoprotein cholesterol are associated with better prognosis in patients with lung adenocarcinoma: a retrospective cohort study

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**Background:** As is well known, lipids play an important role in cellular metabolism and storage, and they have a significant impact on signal transduction during the growth and metastasis of cancer cells. Our study aimed to evaluate the role of the preoperative plasma lipid profile, including triglycerides (TG), total cholesterol (TC), low-density lipoprotein cholesterol (LDL-C), and high-density lipoprotein cholesterol (HDL-C) in prognosis, and to develop nomograms to predict overall survival (OS) and disease-free survival (DFS) based on the preoperative plasma lipid profile for patients with lung adenocarcinoma (LUAD) after complete resection.

**Methods:** Clinical data, including preoperative plasma profile levels, were retrospectively collected and reviewed in 304 patients with LUAD who underwent radical lung resection between 2016–2017. Approval of the study protocol was obtained from the Academic Ethics Committee of Shaoxing People's Hospital, and follow-up on all patients was completed in the clinic or by telephone. The OS and DFS were assessed by the Kaplan-Meier method and the Cox proportional hazards regression model. Clinicopathological factors and preoperative plasma lipid profile factors were integrated to construct nomograms. Calibration plots and concordance indexes (C-indexes) were used to evaluate the accuracy and discrimination of the models.

**Results:** TC level was significantly related to the sex of the patient ( $P=0.001$ ), history of smoking ( $P=0.04$ ), and death ( $P=0.007$ ), and the HDL-C level was significantly associated with sex ( $P=0.004$ ), history of smoking ( $P=0.02$ ), tumor recurrence ( $P=0.050$ ), and death ( $P=0.002$ ).  $TC \leq 3.58$  and  $HDL-C \leq 1.01$  were deemed as independent preoperative risk factors for OS, and  $HDL-C \leq 1.01$  was an independent preoperative risk factor for DFS. In the multivariate analyses involving OS and DFS, an increased TC level [hazard ratio (HR), 0.504; 95% confidence interval (CI): 0.324–0.782,  $P=0.002$ ] was significantly associated with better OS. Additionally, a decreased HDL-C level was significantly associated with worse OS (HR, 0.665; 95% CI: 0.443–0.999,  $P=0.049$ ) and DFS (HR, 0.619; 95% CI: 0.420–0.912,  $P=0.02$ ). Preoperative plasma lipid profile factors were involved in constructing the nomograms for predicting 1-, 3-, and 5-year OS and DFS. The C-index of the final nomograms was higher than that of the tumor node metastasis (TNM) staging system for predicting OS (0.735 *vs.* 0.689;  $P=0.009$ ). The performance of the nomograms for predicting OS (0.699 *vs.* 0.735;  $P=0.03$ ) and DFS (0.659 *vs.* 0.700;  $P=0.002$ ) was significantly lower when preoperative plasma lipid profile factors were excluded. These findings indicated that TC and HDL-C levels are associated with the prognosis in patients with LUAD.

**Conclusions:** In patients with LUAD, increased TC levels may predict better OS, while decreased levels of HDL-C may predict worse outcomes for both DFS and OS. These findings may aid in the identification of high-risk patients and allow them to take necessary measures in advance.

**Keywords:** Plasma lipid profile; lung adenocarcinoma (LUAD); prognosis

Submitted Jun 24, 2024. Accepted for publication Feb 11, 2025. Published online Mar 23, 2025.

doi: 10.21037/tcr-24-1062

View this article at: <https://dx.doi.org/10.21037/tcr-24-1062>

## Introduction

Lung cancer has a poor prognosis and accounts for the majority of new cases and cancer deaths worldwide (1). Lung adenocarcinoma (LUAD), a type of non-small cell lung cancer, is the most common pathological type of lung cancer in humans and the most common type of lung cancer among non-smokers and younger patients. An epidemiological study has reported that adenocarcinoma accounts for approximately 40% of lung cancer cases (2). Due to the high risk of metastasis and recurrence, the prognosis of patients with LUAD is poor. Thus far, a variety

of biomarkers have been studied to predict the prognosis of patients with LUAD (3,4). For example, Zhang *et al.* found that PSMD14 may serve as a potential prognostic marker and therapeutic target for LUAD patients (5), and Sun *et al.* suggested that downregulation of m6A reader YTHDC2 may promote tumor progression and predict poor prognosis in LUAD (6). Due to the high cost and complicated detection equipment required for existing methods, the development of accurate, fast, and convenient predictive biomarkers for the identification of patients with a high risk of metastasis and recurrence is emerging as a significant area of investigation, and it may present an opportunity for improving clinical prognosis and postoperative quality of life (7).

Findings from previous studies have demonstrated that patients with cancer often present with an altered serum lipid profile. A high level of serum triglycerides (TG) was reported to increase the risk of lung cancer (8), and Kitahara *et al.* showed a lower risk of lung cancer in males with higher serum total cholesterol (TC) than in the general population (9). Additionally, a study has indicated low high-density lipoprotein cholesterol (HDL-C) levels are associated with an increased risk of lung cancer (10), and a prospective cohort study of Chinese men showed that low levels of low-density lipoprotein cholesterol (LDL-C) are implicated in an increased risk of lung cancer (11). Several recent studies have confirmed that dyslipidemia is associated with an increased risk of tumor recurrence and reduced survival for multiple cancer types (7,12,13). The studies of lipid biomarkers provided clinical evidence for the treatment of tumors via lipid metabolism. It is widely known lipids are important for cellular metabolism and storage, they exert a critical influence on signal transduction in cancer cell processes, and regulate the growth, invasion, migration, metastasis, and survival of cancer cells through various signaling pathways (14,15). Shen *et al.* found anlotinib suppresses LUAD growth via inhibiting recombinant fatty acid synthase-mediated lipid metabolism (16). Our previous study found that preoperative lipid distribution has an important impact on the prognosis of patients with lung squamous cell carcinoma (7), but there is little evidence

### Highlight box

#### Key findings

- Our study finds that in patients with lung adenocarcinoma (LUAD), increased total cholesterol levels may predict better overall survival, while decreased levels of high-density lipoprotein cholesterol may predict worse outcomes for both disease-free survival and overall survival. These findings may aid in the identification of high-risk patients and allow them to take necessary measures in advance.

#### What is known and what is new?

- It is widely known lipids are important for cellular metabolism and storage, they exert a critical influence on signal transduction in cancer cell processes, and regulate the growth, invasion, migration, metastasis, and survival of cancer cells through various signaling pathways. Shen *et al.* found anlotinib suppresses LUAD growth via inhibiting recombinant fatty acid synthase-mediated lipid metabolism. Our previous study found that preoperative lipid distribution has an important impact on the prognosis of patients with lung squamous cell carcinoma.
- However, there is little evidence about the association between blood lipid levels and clinical characteristics of patients with LUAD and the effect of lipid profiles on prognosis of patients with LUAD. Our current study shows changes in lipid metabolism and its potential to become a survival biomarker in patients with LUAD.

#### What is the implication, and what should change now?

- It may be a therapeutic option for patients with LUAD via regulating lipid metabolism related signaling pathways or gene targets, and combining preoperative blood lipid factors into nomogram models may improve the prediction of the prognosis of LUAD patients compared to using a simple tumor node metastasis staging system.

about the effect of lipid profiles on prognosis of patients with LUAD, lipid metabolism changes and their potential to become biomarkers related to the survival of LUAD require further exploration.

Our current study aimed to develop nomograms to elucidate the effect of lipid distribution on predicting prognosis in patients with LUAD who have undergone complete lung resections. We present this article in accordance with the TRIPOD reporting checklist (available at <https://tcr.amegroups.com/article/view/10.21037/tcr-24-1062/rc>).

## Methods

### *Patient selection*

Records from a total of 304 patients with LUAD who underwent complete lung resections at Shaoxing People's Hospital from January 2016 through December 2017 were collected and reviewed in our retrospective cohort study. We performed a prior sample size estimation by referencing our past research. Blood samples were collected on an empty stomach before surgery, and plasma was prepared and analyzed using an automated biochemical detection instrument, and blood lipids were determined via enzymatic method. All operations were completed by professional personnel with relevant qualifications from our hospital. The tumor node metastasis (TNM) staging criteria for LUAD referred to the latest 2023 9<sup>th</sup> edition of the TNM staging criteria for lung cancer. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). Approval of the study protocol was obtained from the Academic Ethics Committee of Shaoxing People's Hospital (No. 078), and all of the methods were performed in accordance with the relevant guidelines and regulations. Written informed consent was not required due to the retrospective nature of this study. All of the patients met the following eligibility criteria: (I) pathological confirmation of LUAD (pathological diagnoses were confirmed by an independent experienced pathologist at Shaoxing People's Hospital); (II) no previous history of cancer; and (III) had serum samples obtained before treatment (including anti-cancer treatment or any other treatment that may affect blood lipids).

### *Patient follow-up*

Follow-up on all patients was completed in the clinic or by

telephone, and the study was discontinued after a total of 25 patients were unable to complete the designated follow-up. Physical examination and laboratory blood draws were conducted every 3–6 months in the first 2 years and every 6–12 months from years 3–5 (7). The main outcome of this study was overall survival (OS), which we defined as the time from curative resection to death. The secondary outcome was disease-free survival (DFS), which was calculated from the date of radical surgery to the date of disease recurrence or diagnosis of distant metastasis (13).

### *Nomogram construction*

We determined the univariate prognostic factors of OS and DFS using log-rank test. Variables with a  $P < 0.05$  were entered into the multivariate Cox proportional hazards model. The final model was selected using a backward stepdown selection process based on the Akaike information criterion (17). The independent prognostic factors determined through multivariate analysis were then used to construct nomograms for OS and DFS.

### *Nomogram validation and calibration*

The nomograms were subjected to 1,000 bootstrap resamples for validation. The concordance index (C-index) was used to assess the discrimination performance of the nomograms (17). The value of the C-index ranges from 0.5 to 1.0, with a higher C-index value indicating a better capacity to separate patients with different survival outcomes. We utilized previously introduced methods to compare the C-indexes of two different models (17). The TNM staging system in this study was the model, which included the tumor size, number of positive lymph nodes, and metastasis. Calibration represents the capacity of a model to accurately estimate outcomes (17). The observed rates were then compared to the nomogram-predicted probabilities of the models, and the differences between them were used to construct calibration curves. In a well-calibrated model, the predictions are expected to fall on a 45° diagonal line (17).

### *Statistical analysis*

Data analyses were conducted on SPSS 20 and R software, version 3.5.0 (<http://www.r-project.org>). Continuous variables were displayed as mean  $\pm$  standard deviation (SD). The optimal cut-off values for the lipid profile determined

Table 1 Characteristics of patients

Characteristics	Patients, n (%)
Age, years	
>60	136 (44.7)
≤60	168 (55.3)
Gender	
Female	133 (43.8)
Male	171 (56.3)
Smoking state	
Yes	138 (45.4)
No	166 (54.6)
Drinking state	
Yes	103 (33.9)
No	201 (66.1)
T stage	
I–II	263 (86.5)
III–IV	41 (13.5)
Lymph node metastasis	
Yes	139 (45.7)
No	165 (54.3)
TNM stage	
I–II	213 (70.1)
III–IV	91 (29.9)
Recurrence	
Yes	125 (41.1)
No	154 (50.7)
Unknown	25 (8.2)
Death	
Yes	117 (38.5)
No	174 (57.2)
Unknown	13 (4.3)
TG (mmol/L)	
>1.03	199 (65.5)
≤1.03	105 (34.5)
TC (mmol/L)	
>3.58	247 (81.3)
≤3.58	57 (18.8)

Table 1 (continued)

Table 1 (continued)

Characteristics	Patients, n (%)
HDL-C (mmol/L)	
>1.01	232 (76.3)
≤1.01	72 (23.7)
LDL-C (mmol/L)	
>2.20	224 (73.7)
≤2.20	80 (26.3)

TNM, tumor node metastasis; TG, triglycerides; TC, total cholesterol; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol.

by X-Tile software were used to dichotomize lipids for the Chi-squared test and Cox proportional analyses. The relationship between lipid profile and survival rates was obtained with the Kaplan-Meier survival approach and the Log-rank (Mantel-Cox) test. The risk factors for survival were identified using the Cox proportional hazards model. The association between serum lipid levels and clinicopathological parameters, including age, gender, history of smoking, history of drinking alcohol, T stage, lymph node metastasis, TNM stage, recurrence, and death, was tested with the Chi-squared test (7). The R packages cmprsk21 and rms22 were used for modeling and developing the nomograms (17). The rcorr.cens function in the R package Hmisc23 was used for comparing the C-index between two nomograms (17). In all statistical methods,  $P<0.05$  indicated a statistically significant difference.

Results

Patient characteristics

After the eligibility review, 304 patients with LUAD who had undergone radical surgical resection were enrolled in the analysis, and their characteristics are presented in Table 1. The median follow-up time was 42 months (range, 1–70 months). The median age at resection was 59 years (range, 40–81 years); and 171 (56.3%) of the patients were male and the remaining 133 (43.8%) were female. Cancer staging was conducted according to the TNM classification for LUAD. A total of 213 patients (70.1%) were diagnosed with stage I–II disease and 91 (29.9%) were diagnosed with stage III–IV disease. A total of 117 patients (38.5%) died

**Table 2** Clinical characteristics of 304 patients with lung adenocarcinoma based on TG/TC/HDL-C/LDL-C expression status

Characteristics	TG (mmol/L)		TC (mmol/L)		HDL-C (mmol/L)		LDL-C (mmol/L)	
	Mean ± SD	P value	Mean ± SD	P value	Mean ± SD	P value	Mean ± SD	P value
Age, years		0.55		0.65		0.52		0.70
>60	1.472±0.831		4.410±0.841		1.287±0.411		2.645±0.750	
≤60	1.541±1.101		4.363±0.947		1.259±0.351		2.610±0.790	
Gender		0.051		0.002		0.10		0.19
Female	1.636±1.087		4.563±0.902		1.313±0.366		2.692±0.795	
Male	1.412±0.896		4.244±0.876		1.240±0.386		2.574±0.751	
Smoking state		0.009		0.04		0.26		0.71
Yes	1.357±0.651		4.265±0.868		1.245±0.403		2.608±0.748	
No	1.638±1.185		4.483±0.917		1.294±0.356		2.641±0.792	
Drinking state		0.15		0.047		0.72		0.20
Yes	1.395±0.777		4.241±0.876		1.260±0.453		2.546±0.743	
No	1.569±1.078		4.457±0.906		1.278±0.335		2.667±0.784	
T stage		0.46		0.17		0.44		0.30
I–II	1.527±1.014		4.412±0.918		1.278±0.385		2.644±0.792	
III–IV	1.403±0.806		4.205±0.763		1.229±0.331		2.510±0.617	
Lymph node metastasis		0.67		0.65		0.03		0.08
Yes	1.483±1.042		4.409±0.923		1.222±0.317		2.710±0.758	
No	1.533±0.943		4.363±0.883		1.314±0.420		2.555±0.778	
TNM stage		0.12		0.46		0.07		0.43
I–II	1.567±1.055		4.409±0.893		1.297±0.399		2.603±0.773	
III–IV	1.376±0.800		4.325±0.919		1.212±0.317		2.679±0.768	
Recurrence		0.91		0.09		0.03		0.21
Yes	1.494±0.883		4.287±0.812		1.215±0.338		2.569±0.721	
No	1.507±0.988		4.468±0.936		1.317±0.403		2.686±0.800	
Death		0.14		0.09		0.004		0.53
Yes	1.393±0.689		4.289±0.923		1.204±0.326		2.594±0.788	
No	1.566±1.125		4.473±0.880		1.335±0.404		2.652±0.768	

TG, triglycerides; TC, total cholesterol; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; SD, standard deviation; TNM, tumor node metastasis.

during the follow-up period and 125 patients (41.1%) had disease recurrence or distant metastasis.

### *Preoperative plasma profile levels in patients with LUAD*

The relationship between preoperative plasma lipid profile levels and clinical parameters was analyzed (*Table 2*). TG

level ( $P=0.009$ ) and TC ( $P=0.04$ ) level were higher in patients who were non-smokers. A higher TC level was closely correlated with females ( $P=0.002$ ) and no current history of drinking alcohol ( $P=0.047$ ). In addition, HDL-C level was identified to be down-regulated in patients with lymph node metastasis ( $P=0.03$ ) or recurrence ( $P=0.03$ ) or death status ( $P=0.004$ ). Other clinical parameters were



**Table 3** Relationship between TC concentration and clinical characteristics in 304 patients with lung adenocarcinoma based on Chi-squared test

Characteristics	TC >3.58 mmol/L	TC ≤3.58 mmol/L	P
Age, years			0.88
>60	111	25	
≤60	136	32	
Gender			0.001
Female	119	14	
Male	128	43	
Smoking state			0.04
Yes	105	33	
No	142	24	
Drinking state			0.08
Yes	78	25	
No	169	32	
T stage			0.77
I–II	213	50	
III–IV	34	7	
Lymph node metastasis			0.54
Yes	115	24	
No	132	33	
TNM stage			0.73
I–II	172	41	
III–IV	75	16	
Recurrence			0.33
Yes	99	26	
No	129	25	
Death			0.007
Yes	87	30	
No	151	23	

A total of 25 patients were unable to complete the designated follow-up. TC, total cholesterol; TNM, tumor node metastasis.

not significantly associated with preoperative plasma lipid profile levels, such as age, T stage, and TNM stage ( $P>0.05$ ).

*Optimal cut-off determination of lipid metabolism*

The optimal cut-off value, derived with X-tile software

analysis, for TG was 1.03 mmol/L, 3.58 mmol/L for TC, 1.01 mmol/L for HDL-C, and 2.20 mmol/L for LDL-C.

*Association of TC and HDL-C levels with clinical characteristics*

The relationships between the clinicopathological features of LUAD and the levels of TC and HDL-C are summarized in *Tables 3,4*. The Chi-squared test showed that the TC level was significantly related to the sex of the patient ( $P=0.001$ ), history of smoking ( $P=0.04$ ), and death ( $P=0.007$ ) (*Table 3*). However, no significant associations were reported between TC levels and other clinicopathological parameters, including age, history of drinking alcohol, T stage, and tumor recurrence (all  $P>0.05$ ; *Table 3*). The HDL-C level was significantly associated with sex ( $P=0.004$ ), history of smoking ( $P=0.02$ ), tumor recurrence ( $P=0.050$ ), and death ( $P=0.002$ ) (*Table 4*). T stage, lymph node metastasis, and other clinical parameters were not significantly associated with preoperative plasma lipid profile levels (all  $P>0.05$ ; *Table 4*).

*Prognostic significance of clinical characteristics in LUAD*

In the univariate analyses, a significant correlation between history of smoking, T stage, lymph node metastasis, TNM stage, levels of TC, HDL-C, and LDL-C, and OS were detected. Additionally, we found a correlation between lymph node metastasis, TNM stage, HDL-C levels, and DFS. In the multivariate analysis, we observed significant associations between age, T stage, lymph node metastasis, and levels of TC and HDL-C with OS (*Table 5*). In addition, associations between lymph node metastasis and HDL-C levels with DFS were observed (*Table 6*). The multivariate analysis was conducted based on the age at resection, sex, history of smoking, history of drinking alcohol, T stage, lymph node metastasis, TNM stage, and levels of TG, TC, HDL-C, and LDL-C.

*Prognostic significance of the serum lipid profile in LUAD*

Among all 304 patients, 30 of 53 (56.6%) who had a TC level  $\leq 3.58$  mmol/L and 87 of 238 (36.6%) who had a TC level  $>3.58$  mmol/L died ( $P=0.007$ ). In addition, 37 of 65 (56.9%) patients who had an HDL-C level  $\leq 1.01$  mmol/L and 80 of 226 (35.4%) patients who had an HDL-C level  $>1.01$  mmol/L ( $P=0.002$ ) died. In the univariate Cox proportional analysis, a decreased TC [hazard ratio (HR), 0.545; 95% confidence

**Table 4** Relationship between HDL-C concentration and clinical characteristics in 304 patients with lung adenocarcinoma based on Chi-squared test

Characteristics	HDL-C >1.01 mmol/L	HDL-C ≤1.01 mmol/L	P
Age, years			0.83
>60	103	33	
≤60	129	39	
Gender			0.004
Female	112	21	
Male	120	51	
Smoking state			0.02
Yes	97	41	
No	135	31	
Drinking state			0.01
Yes	70	33	
No	162	39	
T stage			0.61
I–II	202	61	
III–IV	30	11	
Lymph node metastasis			0.98
Yes	106	33	
No	126	39	
TNM stage			0.31
I–II	166	47	
III–IV	66	25	
Recurrence			0.050
Yes	89	36	
No	125	29	
Death			0.002
Yes	80	37	
No	146	28	

A total of 25 patients were unable to complete the designated follow-up. HDL-C, high-density lipoprotein cholesterol; TNM, tumor node metastasis.

interval (CI): 0.359–0.826,  $P=0.004$ ] or HDL-C level (HR, 0.557; 95% CI: 0.377–0.823,  $P=0.003$ ) was significantly associated with decreased OS (*Table 5*; *Figure 1*), and this finding remained significant in the multivariate analysis that included T stage and lymph node metastasis. In addition, an

increased LDL-C level (HR, 0.658; 95% CI: 0.446–0.971,  $P=0.04$ ; *Table 5*) and III–IV TNM stage (HR, 0.393; 95% CI: 0.273–0.567,  $P<0.001$ ) were statistically linked with an increased OS in the univariate Cox proportional analysis. However, in the multivariate analysis, the LDL-C levels were not statistically significant, including TNM stage ( $P>0.05$ ; *Table 5*). Other clinical parameters showed no significant difference in the results from either the univariate or multivariate analysis, including sex and history of drinking alcohol ( $P>0.05$ ; *Table 5*).

Regarding DFS, local recurrence or distant metastasis after radical surgical resection was diagnosed in 26 of 51 (51.0%) patients with a TC level  $\leq 3.58$  mmol/L and in 99 of 228 (43.4%) patients with a TC level  $>3.58$  mmol/L ( $P>0.05$ ), as well as in 36 of 65 (55.4%) patients with an HDL-C level  $\leq 1.01$  mmol/L and in 89 of 214 (41.6%) patients with an HDL-C level  $>1.01$  mmol/L ( $P=0.050$ ). In the univariate Cox proportional analysis, a decreased HDL-C level (HR, 0.632; 95% CI: 0.429–0.931,  $P=0.02$ ; *Table 6*) was significantly correlated with decreased DFS, and the significance of this finding remained in the multivariate analysis (HR, 0.619; 95% CI: 0.420–0.912,  $P=0.02$ ), including the lymph node metastasis. In addition, III–IV TNM stage was statistically linked with increased DFS (HR, 0.572; 95% CI: 0.399–0.820,  $P=0.002$ ), although there was no statistical significance of TNM stage in the multivariate analysis ( $P>0.05$ ; *Table 6*). TG, TC, and LDL-C showed no significant difference in the results from either the univariate or multivariate analysis (*Table 6*, *Figure 2*), including age, gender, history of smoking, and history of drinking alcohol ( $P>0.05$ ; *Table 6*).

#### Construction and validation of nomograms for OS and DFS

Nomograms including significant prognostic variables for the OS and DFS of LUAD patients at 1, 3, and 5 years are presented in *Figure 3A, 3B*. Points in the nomograms were assigned based on the hierarchy of effects on survival. The highest points were assigned to lymph node metastasis status for both the OS and DFS nomograms. Although lymph node metastasis status contributed the most to the prognosis, preoperative blood lipid variables moderately impacted prognosis (*Figure 3A, 3B*). Calibration plots revealed a high consistency between the predicted and actual observed 1-, 3-, and 5-year OS and DFS for LUAD patients (*Figure 4A, 4B*). The C-index for the final nomograms for OS was higher than that for the TNM staging system (0.735 vs. 0.689;  $P=0.009$ ; *Table 7*). A lower C-index was generated

**Table 5** Univariate and multivariate Cox proportional analysis with overall survival

Parameters	Univariate analysis		Multivariate analysis	
	HR (95% CI)	P	HR (95% CI)	P
Age (>60 vs. ≤60 years)	1.421 (0.989–2.042)	0.06	1.596 (1.105–2.306)	0.01
Gender (female vs. male)	0.735 (0.506–1.066)	0.11		
Smoking state (yes vs. no)	1.442 (1.003–2.074)	0.048		
Drinking state (yes vs. no)	1.388 (0.954–2.018)	0.09		
T stage (I–II vs. III–IV)	0.499 (0.321–0.778)	0.002	0.575 (0.365–0.906)	0.02
Lymph node metastasis (yes vs. no)	2.773 (1.896–4.056)	<0.001	2.926 (1.975–4.335)	<0.001
TNM stage (I–II vs. III–IV)	0.393 (0.273–0.567)	<0.001		
TG (>1.03 vs. ≤1.03 mmol/L)	0.700 (0.484–1.010)	0.06		
TC (>3.58 vs. ≤3.58 mmol/L)	0.545 (0.359–0.826)	0.004	0.504 (0.324–0.782)	0.002
HDL-C (>1.01 vs. ≤1.01 mmol/L)	0.557 (0.377–0.823)	0.003	0.665 (0.443–0.999)	0.049
LDL-C (>2.20 vs. ≤2.20 mmol/L)	0.658 (0.446–0.971)	0.04		

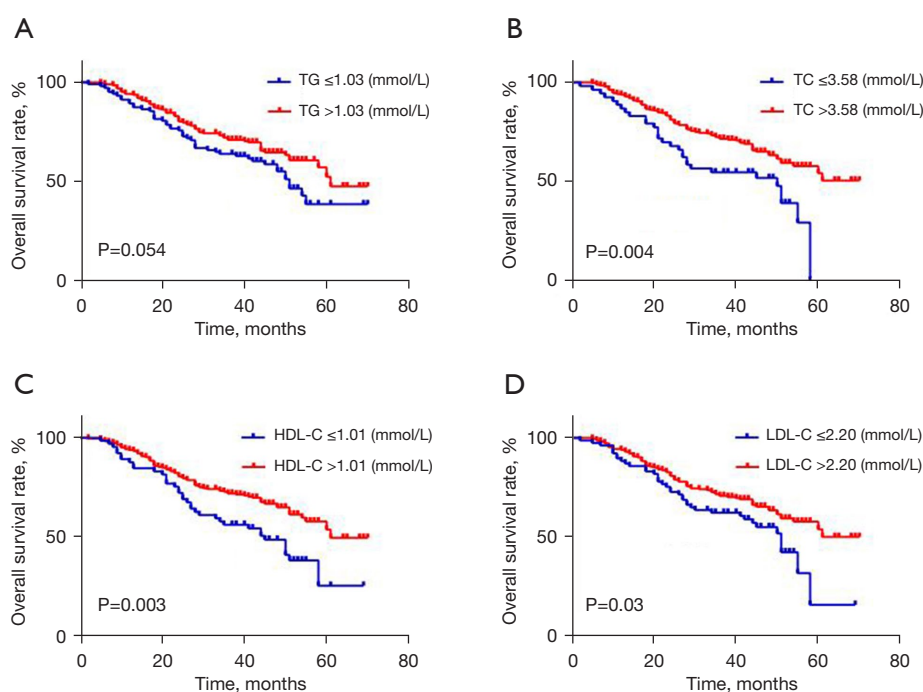
HR, hazard ratio; CI, confidence interval; TNM, tumor node metastasis; TG, triglycerides; TC, total cholesterol; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol.

**Table 6** Univariate and multivariate Cox proportional analysis with disease-free survival

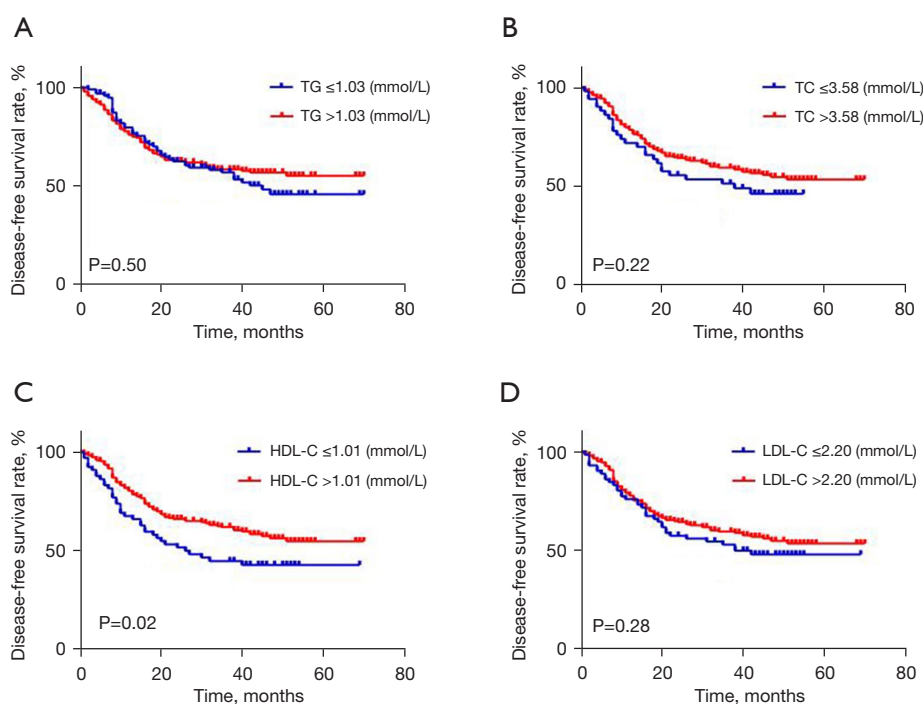
Parameters	Univariate analysis		Multivariate analysis	
	HR (95% CI)	P	HR (95% CI)	P
Age (>60 vs. ≤60 years)	1.030 (0.724–1.465)	0.87		
Gender (female vs. male)	0.803 (0.562–1.148)	0.23		
Smoking state (yes vs. no)	1.283 (0.904–1.823)	0.16		
Drinking state (yes vs. no)	1.174 (0.816–1.689)	0.39		
T stage (I–II vs. III–IV)	0.711 (0.445–1.137)	0.16		
Lymph node metastasis (yes vs. no)	2.005 (1.402–2.867)	<0.001	2.026 (1.417–2.899)	<0.001
TNM stage (I–II vs. III–IV)	0.572 (0.399–0.820)	0.002		
TG (>1.03 vs. ≤1.03 mmol/L)	0.884 (0.615–1.269)	0.50		
TC (>3.58 vs. ≤3.58 mmol/L)	0.767 (0.498–1.181)	0.23		
HDL-C (>1.01 vs. ≤1.01 mmol/L)	0.632 (0.429–0.931)	0.02	0.619 (0.420–0.912)	0.02
LDL-C (>2.20 vs. ≤2.20 mmol/L)	0.808 (0.549–1.190)	0.28		

HR, hazard ratio; CI, confidence interval; TNM, tumor node metastasis; TG, triglycerides; TC, total cholesterol; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol.

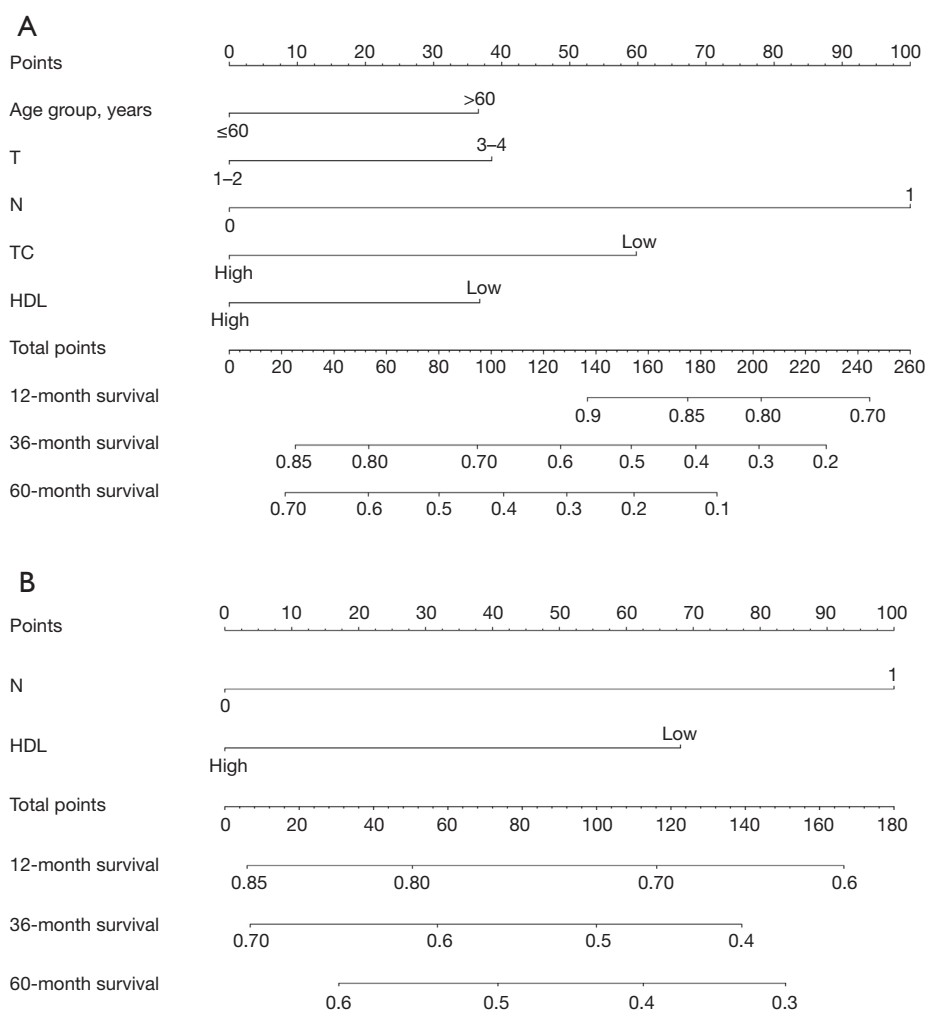




**Figure 1** Kaplan-Meier curves for overall survival regarding low *vs.* high TG levels ( $P=0.054$ ) (A); low *vs.* high TC levels ( $P=0.004$ ) (B); low *vs.* high HDL-C levels ( $P=0.003$ ) (C); and low *vs.* high LDL-C levels ( $P=0.03$ ) (D). TG, triglycerides; TC, total cholesterol; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol.



**Figure 2** Kaplan-Meier curves for disease-free survival regarding low *vs.* high TG levels ( $P=0.50$ ) (A); low *vs.* high TC levels ( $P=0.22$ ) (B); low *vs.* high HDL-C levels ( $P=0.02$ ) (C); and low *vs.* high LDL-C levels ( $P=0.28$ ) (D). TG, triglycerides; TC, total cholesterol; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol.



**Figure 3** Prognostic nomograms in patients with LUAD. Nomograms for OS (A); nomograms for DFS (B). Points of each variable can be estimated by drawing an upward vertical straight line from the variable value of the patient to the axis at the top flagged as “Points”. A vertical straight line is drawn downward from sum of all variable values on the axis of “Total points” to calculate 1-, 3-, and 5-year OS or DFS. LUAD, lung adenocarcinoma; OS, overall survival; DFS, disease-free survival; TC, total cholesterol; HDL, high-density lipoprotein.

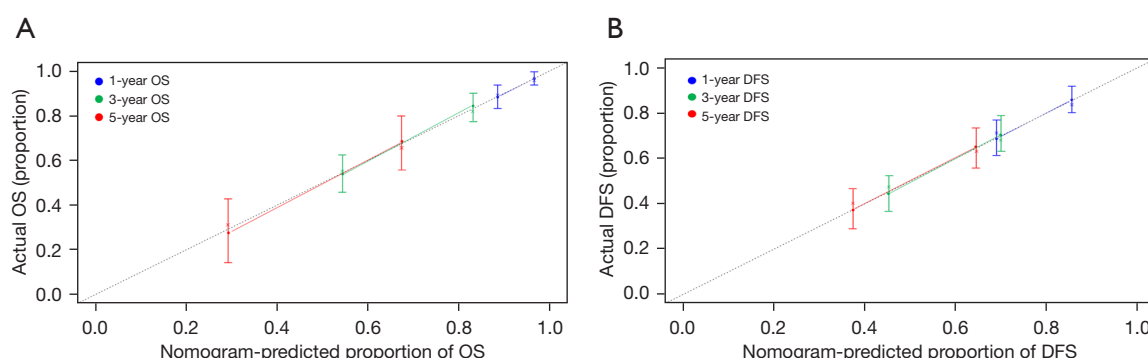
by the OS and DFS nomograms (0.699 *vs.* 0.735,  $P=0.03$ ; 0.659 *vs.* 0.700,  $P=0.002$ , respectively; *Table 7*) that excluded all preoperative blood lipid factors.

## Discussion

Exploring the risk factors of LUAD prognosis and recurrence can provide a clinical basis for postoperative monitoring and early intervention to reduce the tumor recurrence rate and prolong the survival time of patients (18,19). Research shows that blood lipid levels, which are easy and inexpensive to obtain, are closely related to tumor occurrence and recurrence. Therefore, we are interested in

understanding the relationship between blood lipid levels and the prognosis of LUAD patients in China.

There is increasing evidence that dyslipidemia is a diagnostic feature of tumor patients. In this study, lower TC levels indicated a poor OS and was an independent risk factor for the prognosis of patients with LUAD. In addition, lower HDL-C levels were related to sex and history of smoking but not stage of cancer. Cholesterol is essential to maintain cell integrity and various biological functions, and its level changes can affect various signal pathways and related proteins (20,21), such as cell survival kinase Akt (11). Low cholesterol levels are thought to be associated with immunosuppression, up-regulation of mevalonate pathway



**Figure 4** Calibration plots of survival probabilities at 1, 3, 5 years in patients with LUAD. Calibration plots for OS (A); calibration plots for DFS (B). In calibration plots, actual survival is plotted on the vertical axis and predicted survival is plotted on the horizontal. Dotted grey line represents the ideal calibration model in which the predict survival is identical to the actual survival. Vertical bars represent 95% confidence intervals. LUAD, lung adenocarcinoma; OS, overall survival; DFS, disease-free survival.

**Table 7** Comparison of C-indexes for the nomograms and TNM staging system in the patients with LUAD

Items	OS		DFS	
	C-index (95% CI)	P	C-index (95% CI)	P
Nomogram 1	0.735 (0.676–0.794)	Reference	0.700 (0.639–0.757)	Reference
TNM staging system	0.689 (0.627–0.750)	0.009	0.689 (0.627–0.750)	0.60
Nomogram 2 (excluding preoperative plasma lipid profile factors)	0.699 (0.638–0.760)	0.03	0.659 (0.602–0.717)	0.002
Nomogram 3 (excluding age factors)	0.718 (0.658–0.779)	0.08		

TNM, tumor node metastasis; LUAD, lung adenocarcinoma; OS, overall survival; DFS, disease-free survival; CI, confidence interval.

activity or reactivity, and increase in nuclear factor kappa B (NF- $\kappa$ B) activity, thereby promoting the occurrence and development of cancer (7,11). A previous prospective cohort study showed that serum TC level is negatively correlated with total cancer mortality (22). In recent years, research on the correlation between serum cholesterol levels and the risk of lung cancer has gradually increased. Our recent previous study found that preoperative low cholesterol levels are associated with poor prognosis in lung squamous cell carcinoma (7), and Zhang *et al.* also reported that high cholesterol levels reduce the risk of death in patients with advanced non-small cell lung cancer (NSCLC) and who have the epidermal growth factor receptor (*EGFR*) gene mutation (23), thus aligning with our results.

HDL-C is a major indicator of lipoprotein cholesterol (24,25). We revealed that increased HDL-C levels predicted increased DFS and OS after radical surgery. Lower HDL-C levels were related to sex, history of drinking

alcohol, and history of smoking but not stage of cancer. Based on data from previous studies, HDL-C may affect the occurrence of cancer by participating in the reverse transport of cholesterol (26), affecting cell cycle entry (27), and regulating apoptosis and the inflammatory response (7,28,29). Tumor cells increase their pathogenicity by using exogenous fatty acids to support their rapid division while generating large numbers of signal lipids, such as phosphatidylinositol (3,4,5)-triphosphate (PIP3), ceramide-1-phosphate (CIP), and prostaglandins (30). HDL-C can remove excess cholesterol and reduce the consumption and storage of cholesterol in tumor tissues (31). Additionally, HDL may reduce proinflammatory cytokines, such as interleukin-6 (IL-6), interleukin-1 (IL-1), and tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), and increase the level of anti-inflammatory cytokines, such as IL-10, thereby inhibiting cell proliferation and promoting cell apoptosis (13,32,33).

There are different research conclusions about the

influence of LDL-C on the prognosis of lung cancer patients. In small cell lung cancer, lower low-density lipoprotein (LDL) levels were significantly associated with higher OS (34). In lung squamous cell carcinoma, a higher preoperative LDL level indicates a better prognosis (7). This may be due to different tissue types of lung cancer and different treatment options. In our present study, we found that LUAD patients with low LDL had a worse prognosis, although this may not be an independent risk factor.

The abnormality of tumor lipid metabolism is a new field of concern that has gathered attention in recent years. On the one hand, the increase of lipid uptake, storage, and metabolism can promote the growth of tumor cells in various cancers (8-10); while on the other hand, low levels of cholesterol may impair the immune system's ability to weaken tumor inhibition (35). Our study showed that low levels of TC and HDL-C are independent risk factors for the poor prognosis of LUAD patients undergoing surgery.

Building on previous studies investigating the relationship between preoperative blood lipid levels and tumor patient prognosis, our study further attempted to establish a nomogram model to predict patient prognosis to facilitate more accurate decisions for subsequent treatment. Our research indicates that combining preoperative blood lipid factors into nomogram models improves the prediction of the prognosis of LUAD patients compared to using a simple TNM staging system.

As a retrospective study, our article has some shortcomings. For example, a lipid panel was not rechecked after surgery. In addition, some patients with dyslipidemia were administered statins following surgery. A previous study has shown that statins can reduce the risk of death and improve the prognosis of lung cancer patients (36). A systematic review and meta-analysis suggested that statins may be associated with reduced mortality risk and improved OS in observational studies, but such results have yet to be confirmed by randomized controlled clinical trials (37).

## Conclusions

In conclusion, our study showed that preoperative blood lipid levels are closely related to the occurrence and development of LUAD, revealing their potential as a predictor of poor prognosis. Therefore, preoperative TC and HDL-C levels can be considered prognostic factors of LUAD, and dyslipidemia may become a new topic of interest for lung cancer research.

## Acknowledgments

We thank LetPub ([www.letpub.com](http://www.letpub.com)) for its linguistic assistance during the preparation of this manuscript.

## Footnote

*Reporting Checklist:* The authors have completed the TRIPOD reporting checklist. Available at <https://tcr.amegroups.com/article/view/10.21037/tcr-24-1062/rc>

*Data Sharing Statement:* Available at <https://tcr.amegroups.com/article/view/10.21037/tcr-24-1062/dss>

*Peer Review File:* Available at <https://tcr.amegroups.com/article/view/10.21037/tcr-24-1062/prf>

*Funding:* This study was funded by the Zhejiang Medical and Health Science and Technology Projects (grant Nos. 2020KY331 and 2023KY346).

*Conflicts of Interest:* All authors have completed the ICMJE uniform disclosure form (available at <https://tcr.amegroups.com/article/view/10.21037/tcr-24-1062/coif>). 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 conducted in accordance with the Declaration of Helsinki (as revised in 2013). This study was approved by the Academic Ethics Committee of Shaoxing People's Hospital (No. 078), and all of the methods were performed in accordance with the relevant guidelines and regulations. Written informed consent was not required due to the retrospective nature of this study.

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**Cite this article as:** Li Z, Xu J, Wu J, Zhou J, Wang H, Xu H, Wang B, Ding J, Yu G, Xu P. Elevated levels of total cholesterol and high-density lipoprotein cholesterol are associated with better prognosis in patients with lung adenocarcinoma: a retrospective cohort study. *Transl Cancer Res* 2025;14(4):2381-2394. doi: 10.21037/tcr-24-1062