

Correlation identification between radiotracer uptake and pathological features in non-small cell lung cancer (NSCLC)

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ABSTRACT

Background: SUV values from PET/CT can reflect disease progression; however, there are few detailed and comprehensive studies. In this study, we hope to provide data to complement the analysis of diagnostic indicators for NSCLC, LUAD, and LUSC by analyzing the correlation between SUVmax and various pathological parameters.

Patients and Methods: Patients (n=298) with lung lesions were retrospectively studied and clinicopathological index of these patients were collected. Radiomics texture features were extracted by PET/CT scanning and histological features of each patient was collected. The values of SUVmax were obtained and the inner correlation was analyzed. Evaluation and scoring were performed by calculating area under the curve (AUC), sensitivity, specificity and accuracy. **Results:** PET data were counted and analyzed for positive/negative relationship between SUV value and T stage and histology (P<0.05). Mean SUVmax was 13.32±6.41 mm; the SUVmax was 14.01±5.72 for male and 11.19±7.95 for female. The results showed that LUSC tumors were smaller and more homogeneous, but with more uptake and greater PET variability. In contrast, LUAD have lower and weaker uptake, variability and homogeneity.

Conclusion: By meticulously grouping nearly 300 NSCLC samples, AUC values were calculated to indicate the diagnostic value in NSCLC, LUAD and LUSC. It provides ideas and basis for pathological staging analysis of NSCLC.

INTRODUCTION

Lung cancer ranks among the TOP5 global tumor mortality rates, with NSCLC in particular being the most prevalent, accounting for roughly four-fifths of lung cancers ⁽¹⁻⁵⁾. By histological features, lung squamous cell carcinoma (LUSC) and lung adenocarcinoma (LUAD) were two subtype, but the prognosis and recurrence rates of these two are very different ^(6, 7). Therefore, accurate diagnosis is not only meaningful for improving the treatment effect, but also for avoiding unnecessary side effects.

Nowadays, conventional morphological imaging or laboratory indicators have made us increasingly clear about the evolution of diseases, but it is equally important to analyze the changes *in vivo* through non-invasive, specific molecular imaging methods to learn biochemical information ⁽⁸⁻¹⁰⁾. CT or PET/CT is often the first-line imaging method for many diseases such as tumors and is the primary source of baseline data for most NSCLC ⁽¹¹⁾. In NSCLC, FDG (18F-fluoro-2-deoxy-D-glucose) PET/CT (positron emission tomography/computed tomography) has been proved powerful means for staging judgement and efficacy assessment in last decade ^(12, 13). Metrics like metabolic tumor volume (MTV) and total disease glycolysis (TLG) derived from PET/CT imaging [18F] FDG uptake have been reported and confirmed to be

important in the study of metabolic activity of NSCLC ^(14, 15). Other studies have demonstrated that the use of quantitative CT images to describe gene expression data enables prediction of NSCLC survival ^(16, 17). More and more aspects of radiomics, including benign and malignant analysis, prognosis prediction, and metastasis, are showing potential, suggesting the prospect of personalized treatment of tumors ⁽¹⁸⁻²²⁾.

The present study aims to investigate the correlation between pathological indicators and SUV values in NSCLC patients through 18F-FDG PET/CT results. How clinicopathology of lung adenocarcinoma and squamous lung cancer can be organically combined with SUV and be effectively utilized in diagnosis and efficacy is what we have been trying to reveal. In this study, the mean and median values of SUVmax, 40% SUVmax, 50% SUVmax, 60% SUVmax, and 70% SUVmax were firstly finely calculated in nearly 300 NSCLC samples apparently ingested. Secondly, meticulous grouping was performed to calculate whether the clinicopathological indicators had diagnostic value in NSCLC, LUAD and LUSC one by one, i.e., AUC values were counted. This was not available in previous studies. In addition, the correlation between SUV and prognosis was analyzed, which provided ideas and basis for pathological staging analysis of NSCLC.

MATERIALS AND METHODS

Patient cohort and data collection

All enrolled patients were screened who underwent FDG PET/CT at Affiliated Jinhua Hospital, Zhejiang University School of Medicine for characterization or staging a suspected lung tumor (from 2021.05 to 2022.12). A PET/CT scans (before any treatment) baseline has been evaluated and all scans were performed. Inclusion criteria: a) >18 years; b) Presence of a single lung lesion recognisable by both CT and PET; c) a histologically confirmed diagnosis of NSCLC. Exclusion criteria: a) Unknown diagnosis due to tumor tissue too small, or inaccurate diagnosis due to sampling errors; b) Non-malignant tumors; c) FDG uptake below or approximating background activity in healthy lung parenchyma. Available clinical parameters of enrolled samples, including gender, age, smoking history, T stage, lymphatic metastasis, histological type and PET/CT characterizations, were recorded.

Histological types of NSCLC

Tumor histopathology was obtained from each patient by endobronchial ultrasound biopsy or thoracotomy. Histopathological examination was performed according to institutional standards: tumours were classified as LUAD and LUSC based on hematoxylin and eosin (HE) staining. The current WHO/IARC diagnostic criteria were adopted. Staging was conducted according to the VIIth edition TNM classification of malignant tumors. 137 patients were LUAD and 131 were LUSC; small cell carcinoma: 22, large cell carcinoma: 1 and not otherwise specified: 7. This was a retrospective study and all subjects included signed a written informed consent. Ethics approval number: [2021 Ethical Review No. (312)] was obtained from the Ethics Committee of Affiliated Jinhua Hospital.

PET/CT image acquisition

As reported by Boellaard *et al.*, the patients should be positioned supine and raise his or her arms above the head as far as tolerable (23). This means that beam-hardening artefacts in the abdominal and pelvic regions can be avoided, as well as artefacts caused by truncation of the measurement area. The CT scan is performed from the top of the head to the mid-thighs, and the PET acquisition is performed afterwards. The PET scan extends from the mid-thigh to the head. PET covers the same area as a CT scan. PET data and CT data were acquired during shallow breathing and during the baseline respiratory period, respectively. Before receiving ^{18}F -FDG (29702-43, Bayer Healthcare, Germany), patients were well-hydrated. Patients were asked to fast (blood glucose level: 150 mg/dL) 6-8 hours prior to ^{18}F -FDG PET/CT imaging and were imaged with 4 MBq/kg ^{18}F -FDG administered intravenously over approximately 60±5

minutes. PET/CT scanning were obtained by commercial PET/CT scanner (Biograph mCT-S (64); SIE-MENS, Germany) after tracer injection for sixty minutes. The detailed parameters of PET/CT scanner used are listed in table 1.

Table 1. Detailed specifications of the Biograph Vision PET/CT system.

	Biograph Vision
Detector material	LSO
Detector element dimension (mm ³)	3.2×3.2 x 20
Detector elements per block	16×16
Total number of detector elements	60,800
Signal readout	SiPM (2×2 per block)
Axial field-of-view (mm)	263
Transaxial field-of-view (mm)	780
Plane spacing (mm)	1.65
Image planes	119
Coincidence time window (ns)	4.7
Energy window (keV)	435-585
Energy resolution (%)	9
System time resolution (ps)	210
NEMA sensitivity (kcps/MBq)	16.4

Image reconstruction

PET scanning: 4 mins/bed, 6-8 beds/patient (rely on height). The raw CT data were reconstructed into 3.75 mm transverse slices, on the basis of which sagittal and coronal CT images were obtained. The PET/CT analyses were interpreted by two nuclear medicine physicians (random order and independent review), and they all have more than a decade of experience in PET/CT interpretation. All data for patients other than those with suspected NSCLC were not relevant for interpretation. The review committee first analyzed the PET-sole dataset without looking at CT images and then analyzed the PET/CT dataset in following order. After each PET and PET/CT analysis, TNM staging was divided on the basis of consensus within the committee. If no clear distinction could be made between the two stages (T, N, or M), both stages were recorded and considered equivocal.

SUVmax analysis

^{18}F -FDG PET/CT positive lesions were obtained by outlining region of interest (ROI) along the lesion periphery (avoiding the area of radioactive defects such as hemorrhage and necrosis within the lesion) and then automatically calculating the lesion by the computer and obtaining the standardized uptake value (SUV). The calculation formula of SUV is radioactivity per unit volume of lesion tissue / (amount of developer injection / weight of patient). ^{18}F -FDG uptake were evaluated using SUVmax according to the scanning criteria, which determined and evaluated systematically the metabolic activity of Lymph nodes. SUVmax means the maximum standardized uptake value when scanning was performed; a quantitative indicator of radioactive uptake in a lesion. SUVmax was equivalent to developer activity rate per unit volume of lesion tissue to the amount of developer injected. Static

emission scans were acquired with maximum standardized uptake values and weight corrections (for dead time, scattering and attenuation, and attenuation corrections by unenhanced low dose) calculated by the same software. A threshold of 2.5 for SUVmax is considered to allow efficient and specific detection of lesions.

Statistical analysis

All analyses in this study were done with the help of SPSS software (version 26.0, SPSS Inc, Chicago, IL, USA): the unpaired Student's *t*-test (two-tailed probability) and the Wilcoxon rank sum test were carried out using for means comparisons. ROC analysis, which were used to differentiate between groups at high risk of developing end-points, were employed to perform the best critical value estimation for SUVmax. Multivariate binary logistic regression analysis was employed to analyses the relationship between multiple independent variables and the dependent variable, with selected independent variables. *P* value < 0.05 was considered statistically significant.

RESULTS

Patient characteristics

Table 2 summarizes the basic information and NSCLC pathological features included in this retrospective study. Specifically, it includes: (1) The ratio of male to female in the samples collected is about 3: 1 (212: 86). (2) The sample size of patients with LUSC and LUAD was basically the same (LUAD: LUSC=137:131); The other 30 samples were small cell carcinoma: 22 (7.38%), large cell carcinoma: 1 (0.34%) and not otherwise specified: 7 (2.35%). (3) Among these 298 samples, the samples of T1/T2/T3/T4 stages was 95 (31.88%), 75 (25.17%), 45 (15.10%) and 83 (27.85%), the samples in N0/N1/N2/N3 stage was 110 (36.91%), 34 (11.42%), 76 (25.50%) and 78 (26.17%). (4) In addition, the percentage profile of smoking history was: current: 132 (44.30%); former: 30 (10.06%); never: 136 (45.64%).

PET/CT image acquisition

Two hundred and ninety-eight conventional (non-contrast-enhanced) baseline CT scans of NSCLC were included in this study. Two representative scanning sample was shown in figure 1 and figure 2. The imaging diagnosis of the patient in figure 1 was: there was an irregular soft tissue mass of about 4.3×3.9 cm in the left hilum with an SUVmax of 14.8. It was accompanied by bronchial obstruction in upper lobe of left lung; lamellar soft tissue shadows and multiple patchy increased density shadows in the upper left lung, with SUVmax of 13.0 and 2.3, respectively (figure 1A). Increased transparency of both lungs, irregular soft tissue mass in the left hilum, patchy soft

tissue shadow in upper lobe of left lung, accompanied by bronchial obstruction in upper lobe of left lung. Multiple patchy increased density shadows were observed in upper lobe of left lung, and multiple small nodular shadows were observed in the right lung (figure 1B-1C). The imaging diagnosis of the patient in figure 2 was: there was an irregular soft tissue mass in inferior lobe of left lung with a size of 7.6×6.3 cm, SUVmax: 18.8. With distal strip and patchy high-density shadows, the demarcation between the lesion and the adjacent tissue is unclear. In addition, multiple enlarged lymph nodes were seen in the left hilum and mediastinum (group 5 and 7), with an SUVmax of 13.5, and multiple nodular thickenings in the left pleura with an SUVmax of 8.5 (figure 2A). Mild breast imaging on both sides, roughly symmetrical on both sides, increased uptake of strip-like imaging agent in the lower esophagus, and no abnormal imaging agent concentration in other parts of the lungs (figure 2B-2C).

Table 2. The tumor and patient characteristics.

Characteristic	Value
Age (range) years	64.81±10.60 (27-86)
Gender, n(%)	
Male	212 (71.14%)
Female	86 (28.86%)
T Stage, n(%)	
T1	95 (31.88%)
T2	75 (25.17%)
T3	45 (15.10%)
T4	83 (27.85%)
N Stage, n(%)	
N0/Nx	110 (36.91%)
N1	34 (11.42%)
N2	76 (25.50%)
N3	78 (26.17%)
Smoking History, n(%)	
Current	132 (44.30%)
Former	30 (10.06%)
Never	136 (45.64%)
Histology, n(%)	
Adenocarcinoma	137 (45.97%)
Squamous cell carcinoma	131 (43.96%)
Small cell carcinoma	22 (7.38%)
Large cell carcinoma	1 (0.34%)
Not otherwise specified	7 (2.35%)

Note: T1: T1 stage; T2: T2 stage; T3: T3 stage; T4: T4 stage; N0/Nx: N0/Nx stage; N1: N1 stage; N2: N2 stage; N3: N3 stage

Prediction and discriminating values of SUVmax

In this study, the SUVmax was reported higher than 2.5 (i.e. significant uptake) in 296 patients by PET/CT, and only two patients reported a SUVmax of less than 2.5 as measured by PET/CT. For PET features, the mean value and median value of SUVmax, 40% SUVmax, 50% SUVmax, 60% SUVmax and 70% SUVmax were calculated. The specific mean value were 12.31 ± 7.11, 4.92 ± 2.84, 6.16 ± 3.55, 7.39 ± 4.27 and 8.62 ± 4.98, respectively; the specific median value were 11.20 (range: 0-41), 4.48 (range: 0-16.4), 5.60 (range: 0-20.5), 6.72 (range: 0-24.6) and 7.84 (range: 0-28.7), respectively (table 3).

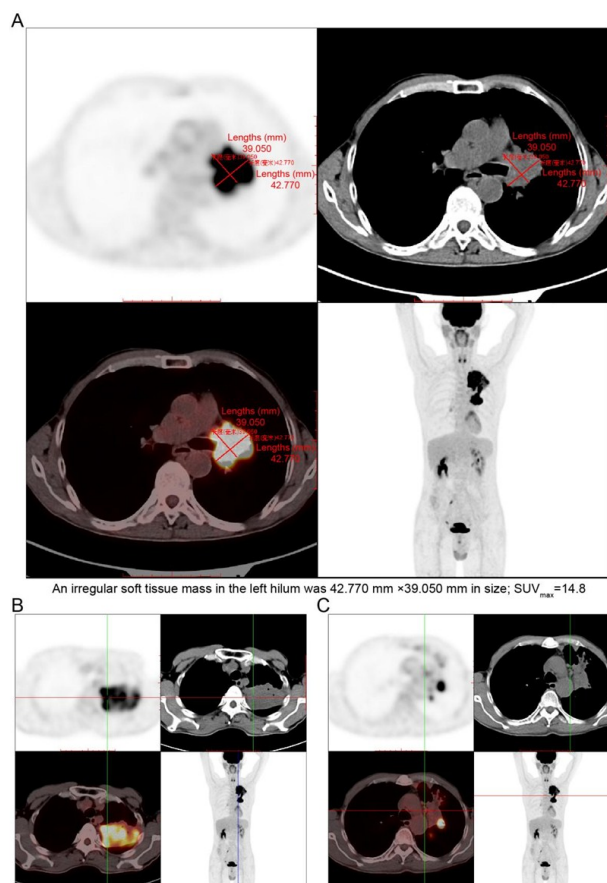


Figure 1. Computed tomographic slices from a patient with non-small-cell lung cancer. **A.** An irregular soft tissue mass in the hilum of the left lung, 4.3×3.9 cm in size; SUV_{max}: 14.8; **B-C.** Increased transparency of both lungs, irregular soft tissue mass in the left hilum, patchy soft tissue shadow in the upper lobe of the left lung, accompanied by bronchial obstruction in the upper lobe of the left lung. Multiple patchy increased density shadows were observed in the upper lobe of the left lung, and multiple small nodular shadows were observed in the right lung.

Table 3. Intra-modality correlation between PET features.

	Mean±SD	Median, range
SUV _{max}	12.31±7.11	11.20, 0-41
40%SUV _{max}	4.92±2.84	4.48, 0-16.4
50%SUV _{max}	6.16±3.55	5.60, 0-20.5
60%SUV _{max}	7.39±4.27	6.72, 0-24.6
70%SUV _{max}	8.62±4.98	7.84, 0-28.7

Correlation between clinical characteristics and SUV_{max}

The low SUV_{max} and high SUV_{max} groups were grouped according to the median SUV_{max}; and univariate analysis and multivariate analysis were performed, subsequently. The univariate analysis revealed that age (95% CI: 0.043-0.139; OR=0.078; $P<0.01$) and gender (95% CI: 0.080-0.555; OR=0.211; $P<0.01$) were different between low SUV_{max} group and high SUV_{max} group (table 4). The results of the multivariate analysis showed that differences in smoking history (95% CI: 0.053-0.486; OR=0.161; $P<0.01$), T stage (95% CI: 1.980-114.220; OR=15.039; $P<0.01$) N stage (95% CI: 3.846-75.143; OR=17.000;

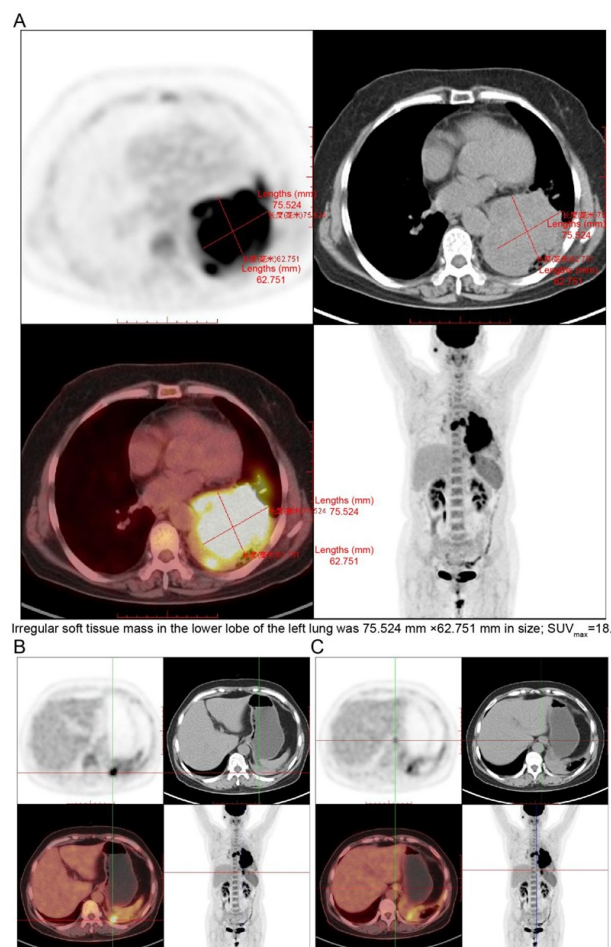


Figure 2. Computed tomographic slices from a patient with non-small-cell lung cancer. **A.** An irregular soft tissue mass in the lower lobe of the left lung, size 7.6×6.3 cm, SUV_{max}: 18.8; **B-C.** Mild breast imaging on both sides, roughly symmetrical on both sides, increased uptake of strip like imaging agent in the lower esophagus, and no abnormal imaging agent concentration in other parts of the lungs.

$P<0.01$), and histology features (95% CI: 1.604-19.849; OR=5.642; $P<0.01$) between low SUV_{max} group and high SUV_{max} group (table 4). In addition, to further confirm whether T stage, N stage and smoking history was associated with histological characteristics (LUAD or LUSC) according to SUV_{max} value, we performed the following statistical analysis (table 5). The analysis revealed that in LUAD, high SUV_{max} group and low SUV_{max} group presented differences on age ($P<0.01$), T stage ($P<0.01$) and N stage ($P<0.01$), respectively. In contrast, in LUSC, only age ($P<0.01$) differed between high SUV_{max} group and low SUV_{max} group (table 5).

ROC curve analysis of SUV_{max} by pathological features

According to the above analysis, the median value of SUV_{max} was used as the cut off for gender, age, smoking history, TNM stage and histology grouping based on the results analysed above. The ROC curve analysis revealed that significant difference was calculated in four indexes: gender, smoking history, T

stage and histology feature ($p < 0.01$, figure 3, D, F, J and L); AUC were 0.673, 0.666, 0.632 and 0.718, respectively. Also, the ROC curve analysis of age showed that the AUC were 0.573 and the difference is significant ($p < 0.05$, figure 3E). The ROC curve analysis in N stage have no significant difference, and the AUC were 0.603, respectively (figure 3K). The distribution of SUVmax in each subgroup was shown in the figure 3A-C, 3G-I. The median value of SUVmax was used as the cut off for gender, age, smoking history, TNM stage grouping based on the results analysed above in LUSC. The ROC curve analysis revealed that the AUC were 0.627, 0.558, 0.527, 0.563 and 0.598, respectively (figure 4D-F, 4I and J); no

significant difference was found in these indexes. The distribution of SUVmax in each subgroup was shown in the figure 4A-C, 4G-H. The median value of SUVmax was used as the cut off for gender, age, smoking history, TNM stage grouping based on the results analyzed above in LUAD. The ROC curve analysis revealed that there was a significant difference in T stage and its AUC value was 0.641 ($p < 0.01$, figure 5I). The ROC curve analysis in other four indexes showed no significant difference and AUC value were 0.592, 0.559, 0.606 and 0.595, respectively (figure 5D-F, 5J). Besides, the distribution of SUVmax in each subgroup was shown in the figure 5A-C, 5G-H.

Table 4. Logistic regression analysis.

	Age (mean)	Gender (M/F)	Smoking History (C/F/N)	T Stage (1/2/3/4)	N Stage (0/1/2/3)	Histology (A/S/Not)
Low SUVmax Group	57.42±9.19	7/12	4/0/15	17/1/0/1	17/2/0/0	16/3/0
High SUVmax Group	65.32±14.02	205/74	128/30/121	78/74/45/82	93/34/74/78	121/128/7
Uni-analysis	OR	0.078	0.211	-	-	-
	95% CI	0.043-0.139	0.080-0.555	-	-	-
	P	0.000**	0.001**	-	-	-
Multi-analysis	OR	-	-	0.161	15.039	5.642
	95% CI	-	-	0.053-0.486	1.980-114.220	1.604-19.849
	P	-	-	0.000**	0.001**	0.003**

Note: M: Male; F: Female; C: Current; F: Former; N: Never; 0: NO stage; 1: T1 stage/N1 stage; 2: T2 stage/N2 stage; 3: T3 stage/N3 stage; 4: T4 stage; A: Adenocarcinoma; S: Squamous cell carcinoma; Not: Not otherwise specified; Uni-analysis: Univariate analysis; Multi-analysis: Multivariate analysis.

Table 5. The results of correlation and regression analysis.

	LUAD		LUSC	
Items	Low Group vs. High Group n (%)	χ^2/P	Low Group vs. High Group n (%)	χ^2/P
Age (mean)	54.81±7.16; 64.65±11.32	37.322/0.000**	71.33±14.19; 65.18±7.73	82.987/0.000**
Gender		2.796/0.094		0.122/0.727
Male	4 (25.00%); 57 (47.11%)		3 (100%); 123 (96.09%)	
Female	12 (75.00%); 64 (52.89%)		0 (0%); 5 (3.91%)	
Smoking		1.194/0.550		0.992/0.609
Current	2 (12.50%); 23 (19.01%)		2 (66.67%); 86 (67.18%)	
Former	0 (0%); 5 (4.13%)		0 (0%); 21 (16.41%)	
Never	14 (87.50%); 93 (76.86%)		1 (33.33%); 21 (16.41%)	
T Stage		21.279/0.000**		0.860/0.835
T1	16 (100%); 47 (38.84%)		1 (33.33%); 28 (21.88%)	
T2	0 (0%); 19 (15.70%)		1 (33.33%); 44 (34.38%)	
T3	0 (0%); 14 (11.57%)		0 (0%); 25 (19.53%)	
T4	0 (0%); 41 (33.88%)		1 (33.33%); 31 (24.22%)	
N Stage		20.663/0.000**		2.032/0.566
N0	16 (100%); 48 (39.67%)		1 (33.33%); 42 (32.81%)	
N1	0 (0%); 12 (9.92%)		0 (0%); 17 (13.28%)	
N2	0 (0%); 18 (14.88%)		2 (66.67%); 42 (32.81%)	
N3	0 (0%); 43 (35.54%)		0 (0%); 27 (21.09%)	

Note: LUAD: Adenocarcinoma; LUSC: Squamous cell carcinoma; SCLC: small cell lung cancer; LCLC: Large cell lung cancer; Not: Not otherwise specified.

Table 4. Logistic regression analysis.

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Uni-analysis	OR	0.078	0.211	-	-	-
	95% CI	0.043-0.139	0.080-0.555	-	-	-
	P	0.000**	0.001**	-	-	-
Multi-analysis	OR	-	-	0.161	15.039	5.642
	95% CI	-	-	0.053-0.486	1.980-114.220	1.604-19.849
	P	-	-	0.000**	0.001**	0.003**

Note: LUAD: Adenocarcinoma; LUSC: Squamous cell carcinoma; SCLC: small cell lung cancer; LCLC: Large cell lung cancer; Not: Not otherwise specified.

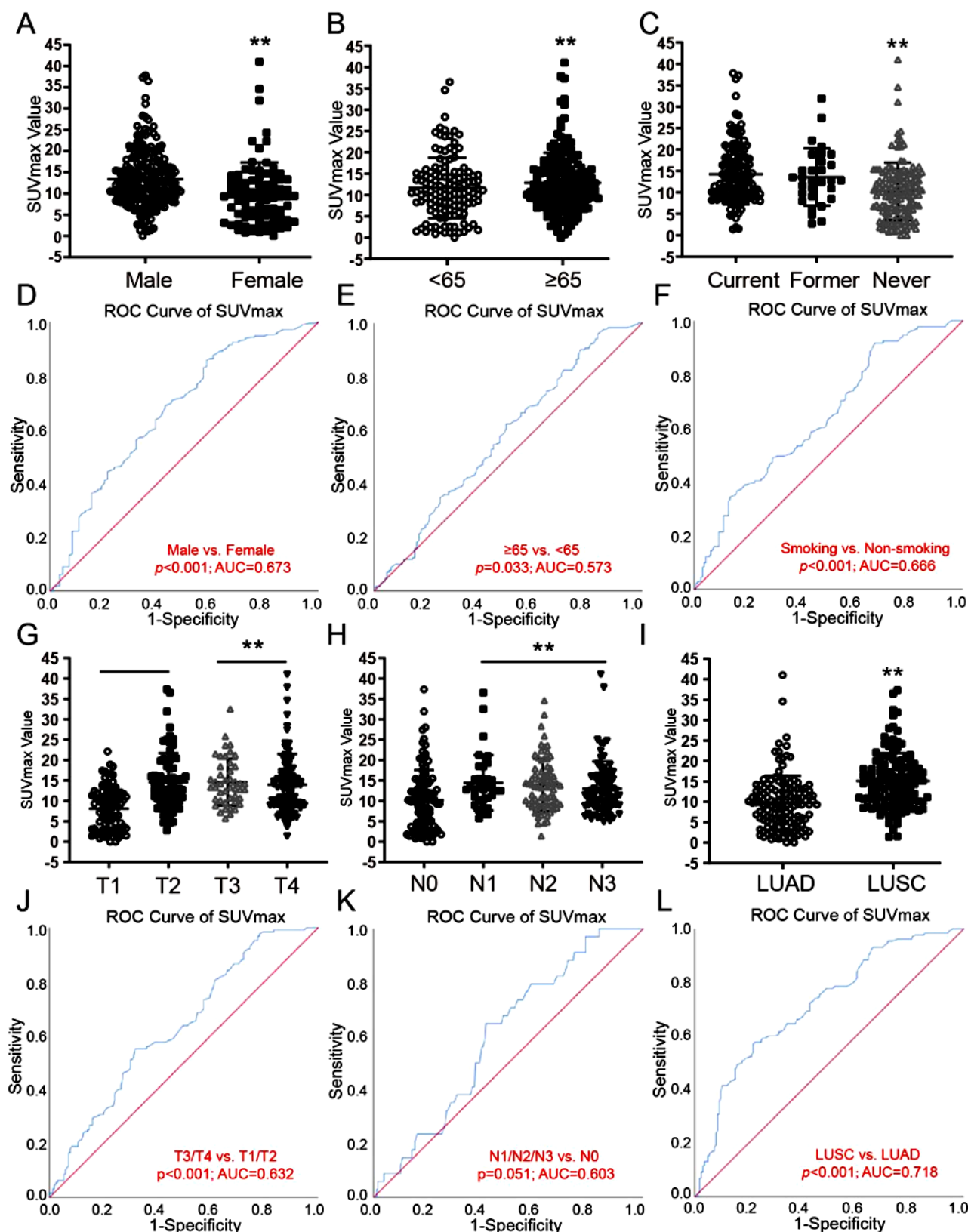


Figure 3. ROC curve analysis establishing the cut off value of SUVmax. (A&D) The distribution and ROC curve of SUVmax in male and female; (B&E) The distribution and ROC curve of SUVmax in <65 years and ≥ 65 years; (C&F) The distribution and ROC curve of SUVmax in current smoker, former smoker and never smoker; (G&J) The distribution and ROC curve of SUVmax in T1/T2/T3/T4; (H&K) The distribution and ROC curve of SUVmax in N0/N1/N2/N3; (I&L) The distribution and ROC curve of SUVmax in LUAD and LUSC.

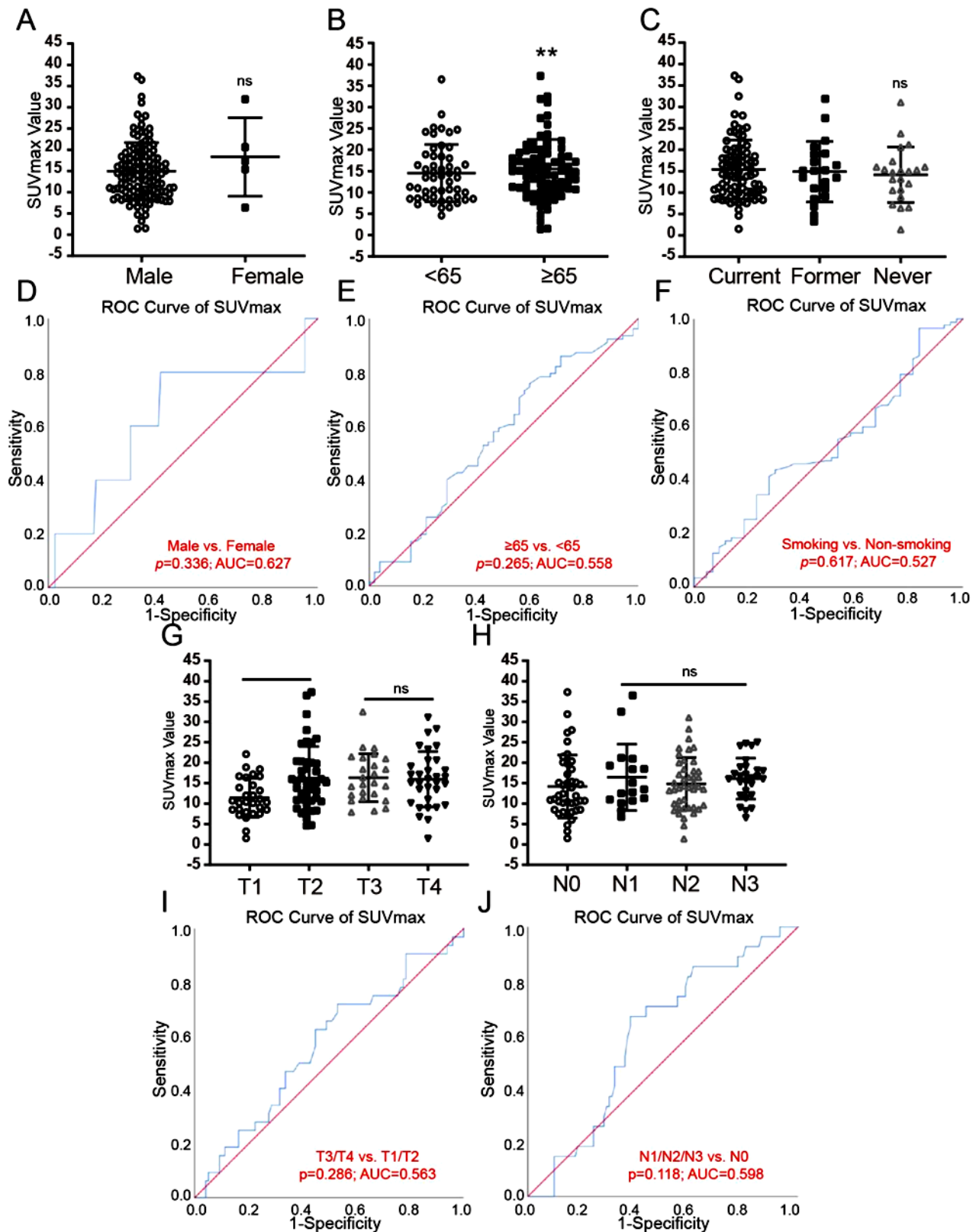


Figure 4. ROC curve analysis establishing the cut off value of SUVmax in LUSC. (A&D) The distribution and ROC curve of SUVmax in male and female; (B&E) The distribution and ROC curve of SUVmax in <65 years and ≥ 65 years; (C&F) The distribution and ROC curve of SUVmax in current smoker, former smoker and never smoker; (G&I) The distribution and ROC curve of SUVmax in T1/T2/T3/T4; (H&J) The distribution and ROC curve of SUVmax in N0/N1/N2/N3.

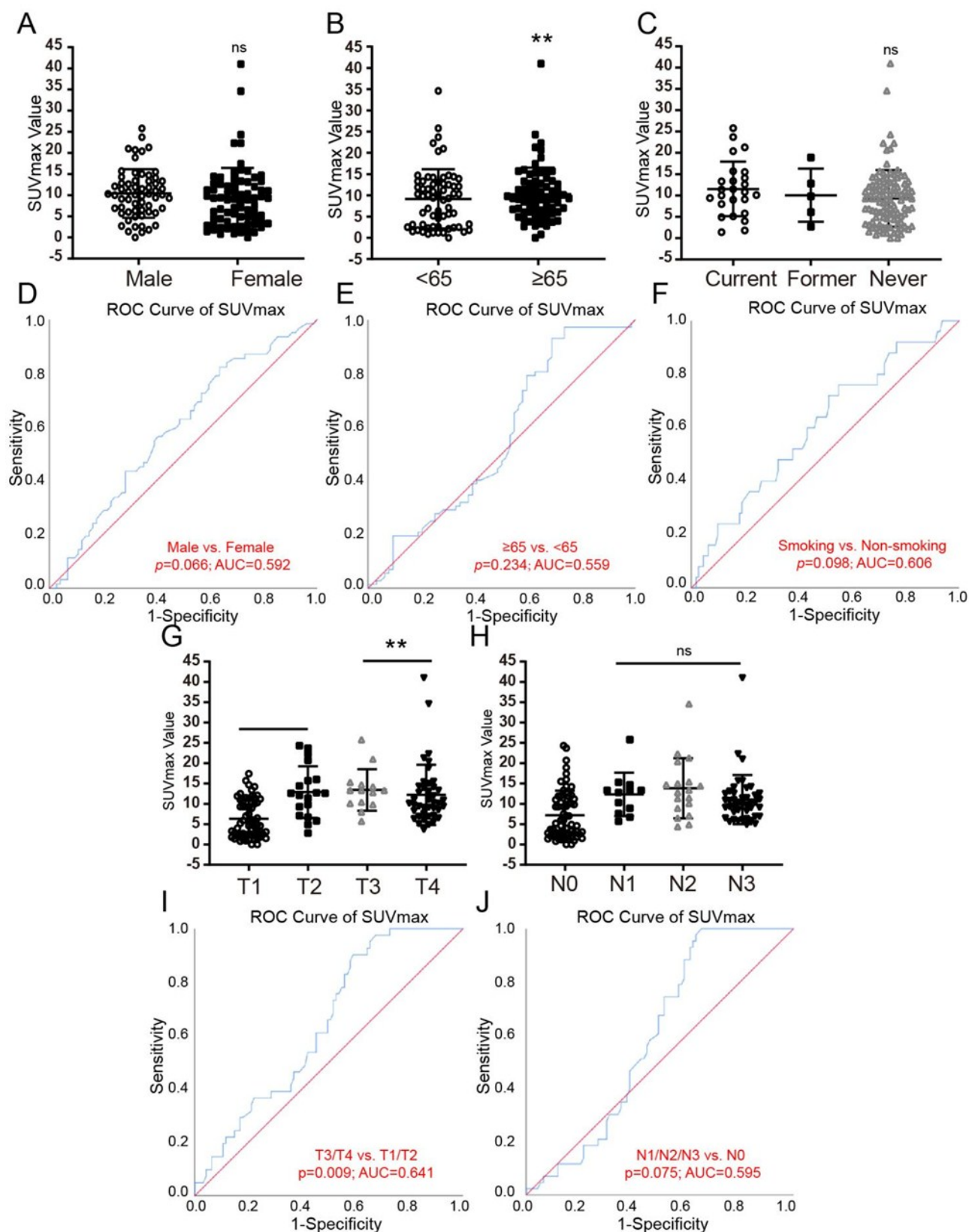


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Table 5. The results of correlation and regression analysis

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Gender		2.796/0.094		0.122/0.727
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N3	0 (0%); 43 (35.54%)		0 (0%); 27 (21.09%)	

DISCUSSION

The increasing role of radiomics in clinical is nothing less than its convenient realization of non-invasive acquisition of *in vivo* lesions characteristics (including prognostic data and treatment response) (24, 25). PET/CT has the advantage of achieving co-integration of images from both functional and anatomical morphological perspectives, but it still suffers from deficiencies such as too small lesions that cannot be identified, and poor display of anatomical structures around lesions due to tissue specificity of tracers, and difficult image comprehension (26, 27). Thus, Biopsy is necessary for histological classification of disease. In our study, the relations between [18F] FDG PET features, CT features, and histological types have been evaluated. After nearly 300 NSCLC samples were grouped according to SUVmax, statistical analysis confirmed the correlation between age, gender and SUVmax group; and between smoking history, T stage, N stage and histological features and SUVmax group.

We know that SUVmax can reflect the degree of tumor uptake of FDG, that is, an indicator that reflects the level of tumor glucose metabolism. Metabolic volume MTV that reflects the tumor burden of solid tumors can be obtained by PET/CT imaging (28-30). Therefore, we next used the median values of 40% SUVmax, 50% SUVmax, 60% SUVmax and 70% SUVmax to count the correlations with clinical indicators, respectively. Statistical analysis of our results revealed that SUVmax correlated with tumor clinical stage, lymph node involvement, and histological features. Previous studies have shown that patients with higher SUVmax values have a poorer prognosis compared with those with lower SUVmax values, indicating higher glycolytic activity, which predicts adverse responses to subsequent therapy (31). Also, Pyka *et al.* uncovered that tumor

heterogeneity obtained by FDG-PET texture analysis correlated with the response of NSCLC patients to radiotherapy efficacy, suggesting that imaging and correspondence analysis data can be useful for individualizing treatment and analyzing the population benefiting from different treatment options (32).

Studies have confirmed that the sensitivity, specificity, positive predictive rate and negative predictive rate of PET/CT were 96%, 92%, 92%, and 95%, respectively (33). In our study, we combined PET/CT results and clinicopathological data and performed correlation analysis by grouping according to SUVmax values. It revealed a correlation between age, gender, smoking history, and stage with SUVmax groups in NSCLC. It also confirmed that age, T-stage and N-stage were correlated with SUVmax values in LUAD by lung adenocarcinoma or lung squamous carcinoma, while only age was correlated with SUVmax values in LUSC. Similar studies are scarce, with one similar study of only 119 cases by Christian *et al.* (34). Our results are a strong addition to the clinical data.

LUAD and LUSC, the two main histological subtypes of NSCLC, differ in terms of prognosis and recurrence. Therefore, accurate diagnosis and follow-up can avoid unnecessary side effects (35). Clinical patients will benefit from the combination of pathological staging and non-invasive imaging, which is used to further adjust therapeutic plans by tracking pathological progression. Therefore, our study enrolled a nearly 300 cases sample size (LUAD: LUSC close to 1:1), and try to discover the patterns and correlations and provide data support for clinical application. The median of SUVmax was used as the cutoff value for gender, age, smoking history, TNM stage, and histological grouping. ROC curve analysis revealed significant differences in four indicators: gender, smoking history, T stage, and histological

characteristics. The current study is not without its limitations: on the one hand, the size of the sample, which may produce some unexpected results, and there is no systematic statistics between CT features and overall survival, which is not helpful for the analysis of survival. It is worth mentioning that, by calculating the AUC value, we carried out ROC analysis to determine the relationship between SUVmax and pathological indicators ⁽³⁶⁾.

CONCLUSION

In this study, the diagnostic accuracy and correlation of SUVmax with pathological indicators of patients with NSCLC were evaluated. These data should be fully utilized and organically combined for individualized treatment and prognostic assessment. This study analyzed the diagnostic value of SUV under different subgroups, which can provide research ideas and data support for pathological staging analysis of NSCLC.

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