

Association of Lung Cancer Risk With the Presence of Both Lung Nodules and Emphysema in a Lung Cancer Screening Trial

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Abstract

Background: The coexistence of emphysema and lung nodules could interact with each other and then lead to potential higher lung cancer risk. The study aimed to explore the association between emphysema combined with lung nodules and lung cancer risk.

Methods: A total of 21,949 participants from the National Lung Screening Trial (NLST) who underwent low-dose computed tomography (LDCT) examination were included. Participants were categorized into four groups (NENN group (non-emphysema and non-nodules), E group (emphysema without nodules), N group (nodules without emphysema), and E + N group (nodules with emphysema)) according to whether there were lung nodules and emphysema. Multivariable Cox regression and stratified analyses were performed to estimate the association between the four groups and lung cancer risk.

Results: Among the 21,949 participants, there were 9,040 (41.2%), 5,819 (26.5%), 4,737 (21.6%), and 2,353 (10.7%) participants in the NENN group, E group, N group, and E + N group. The risk of lung cancer incidence increased in turn in NENN group, E group, N group and E + N group. Compared with NENN group, the ageadjusted hazard ratios (HRs) (95% confidence intervals (CIs)) of lung cancer incidence were 2.07 (1.69 - 2.54) for E group, 4.13 (3.47 - 5.05) for N group, and 6.26 (5.14 - 7.62) for E + N group. The association was robust to adjustment for potential confounders (1.83 (1.47 - 2.27) for E group, 3.97 (3.24 - 4.86) for N group, and

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5.23 (4.28 - 6.48) for E + N group). Comparable results as the lung cancer incidence were observed for lung cancer mortality, whether in age-adjusted model (E group: 1.85 (1.39 - 2.46), N group: 2.49 (1.89 - 3.29), E + N group: 4.27 (3.21 - 5.68)) or fully adjusted model (E group: 1.56 (1.15 - 2.11), N group: 2.43 (1.81 - 3.26), E + N group: 3.39 (2.50 - 4.61)). However, the trend of all-cause mortality risk among the four groups was somewhat different from that of lung cancer risk, whether in age-adjusted model (1.37 (1.21 - 1.54) for E group, 1.06 (0.92 - 1.21) for N group, and 1.75 (1.51 - 2.02) for E + N group) or fully adjusted model (1.26 (1.10 - 1.44) for E group, 1.09 (0.94 - 1.27) for N group, and 1.52 (1.30 - 1.79) for E + N group).

Conclusion: Based on a large-scale lung cancer screening trial in the United States, this study demonstrated that either emphysema or lung nodules can increase lung cancer risk, and lung nodules combined with emphysema can further increase the lung cancer risk and all-cause mortality. The significance of these findings for lung cancer screening should be evaluated.

Keywords: Lung nodules; Emphysema; Lung cancer; Incidence; Mortality

Introduction

Lung cancer has been the most common cancer and the main cause of cancer death for many years. In 2020, it is estimated that there were 2.2 million new lung cancer cases and 1.8 million lung cancer deaths, accounting for 11.4% of all new cancer cases and 18.0% of all cancer deaths [1]. Despite the evolving systemic treatment scenario for lung cancer [2-4], resulting in a 5-year survival rate of 17.4% [5], the increasing burden of lung cancer combined with poor prognosis is a challenge to cancer prevention. In 2011, the National Lung Screening Trial (NLST) proved for the first time that lowdose computed tomography (LDCT) screening in high-risk population can reduce lung cancer mortality by 20-33% [6]. Therefore, LDCT screening has been introduced and is now recommended as a strategy for early detection of lung cancer all over the world [7].

Lung nodule is a common finding of LDCT in lung cancer screening. Although about 95% of nodules are benign, patients

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This article is distributed under the terms of the Creative Commons Attribution Non-Commercial 4.0 International License, which permits unrestricted non-commercial use, distribution, and reproduction in any medium, provided the original work is properly cited and clinicians are still worried about the diagnosis of lung cancer [8-11]. Recently, the Dutch-Belgian Randomized Lung Cancer Screening Trial (Nederlands-Leuvens Longkanker Screenings Onderzoek (NELSON)) reported new solid nodules discovered at two post-baseline screens of approximately 8,000 subjects [12]. Several other studies have also reported on the presence of risks associated with new nodules, including the Mayo Clinic study and the International Early Lung Cancer Action Program [13, 14]. These studies reported rates of 5-13% for the detection of new nodules per annual postbaseline screen [12-14]. Overall lung cancer risks for new nodules were generally approximately 5%.

LDCT can not only detect lung nodules in early lung cancer, but also reveal other important lung findings, such as emphysema. There are a lot of studies on emphysema as a risk factor for lung cancer [15, 16]. More and more evidence confirmed that there is a correlation between the existence of emphysema and lung cancer incidence and cancer-related mortality, regardless of smoking history and nodule size [17-23].

Some studies on lung cancer screening show that 48-58% of patients with emphysema also have lung nodules [24], and the coexistence of emphysema and lung nodules will affect carcinogenesis, which may lead to higher lung cancer mortality because of increased sensitivity to biological damage [25]. There is also evidence that almost 80% of patients screened by LDCT have solitary lung nodules coexisting with chronic obstructive pulmonary disease (COPD) or emphysema, which is related to the unfavorable outcome [26-28].

In view of the above, several studies have explored the relationship between the solitary existence of lung nodules or emphysema and lung cancer risk. But there is no independent study on the risk of emphysema combined with pulmonary nodules. The purpose of this study was to explore the association of lung cancer risk with the presence of both lung nodules and emphysema by using the first round of LDCT screening data in NLST database.

Materials and Methods

Institutional Review Board approval

Consent to access data from the NLST was obtained from the National Cancer Institute's Cancer Data Access System through a data transfer agreement among us, the authors, and the National Cancer Institute (NCI). Nonidentifiable patient data were used in this study for secondary data analysis, and its use was approved by the institutional review boards of our institutions.

Ethical statement

This human study was approved by NCI and their local institutional review board. Each participating center's institutional review board approved the protocols and all participants provided written informed consent. The trial registration number (on ClinicalTrials.gov) of NLST is NCT00047385.

Ethical compliance with human/animal study

This study was conducted in compliance with the ethical standards of the responsible institution on human subjects as well as with the Helsinki Declaration.

Source of population

The designs of the NLST cancer screening trials have been previously published. Briefly, from 2002 to 2004, the NLST cancer screening trial randomized 53,452 smokers aged 55 - 74 years with at least 30 pack-years of smoking history and at most 15 years smoking cessation history to receive three annual LDCT screening (the intervention arm) or posterior-anterior chest X-ray screening (the control arm) in 1:1 ratio [6].

Selection of participants

In the NLST, we first excluded 26,730 participants in the control group who received chest X-ray screening, and 26,722 participants in the intervention group who received LDCT screening were initially included in this study. After further excluding the participants who did not meet the enrollment conditions, did not have LDCT in the first round (T0) or had no information about the results of LDCT, a total of 21,949 participants were finally included in this study (Fig. 1). According to the LDCT results of the first round (T0) on whether there is emphysema, the participants were divided into two groups (emphysema group and non-emphysema group), and further according to the presence or absence of lung nodules, the participants were divided into four groups: NENN group (nonemphysema and non-nodules), E group (emphysema without nodules), N group (nodules without emphysema), and E + Ngroup (nodules with emphysema).

Information of baseline variables

After informed consent, a baseline questionnaire was provided to all participants in the NLST to collect the information about cancer-related demographics and potential risk factors, such as demographics, smoking history, family history of cancer, medical history, etc. In the end, these potential confounders were included in this analysis based on previous literature and availability of data in the NLST: age (55 - 59, 60 - 64, 65 - 69, 70 - 74 years), sex (female, male), race/ ethnicity (non-Hispanic White, others), education level (< senior high school, senior high school, college and above), marital status (married/living as married, widowed/divorced/ separated, never married), smoking status (former smoking, current smoking), smoking pack-year (30 - 59, 60 - 89, \geq 90 pack-year), body mass index (BMI) (< 25, 25 - 29.9, \geq 30



Figure 1. Flow chart of participants' selection. NENN group: non-emphysema and non-nodules; E group: emphysema without nodules; N group: nodules without emphysema; E + N group: nodules with emphysema.

 kg/m^2) and family history of lung cancer (no, yes). BMI is calculated by dividing the weight in kilograms by the square of the height in meters (kg/m^2).

Nodules characteristics

We defined lung nodules as participants with nodules ≥ 4 mm, otherwise as having no lung nodules. According to whether the participants were diagnosed as emphysema in the first round (T0), we divided the participants diagnosed as lung nodules into two groups: N group (nodules without emphysema), and E + N group (nodules with emphysema). The characteristics of lung nodules include size (4 - 6, 7 - 10, 11 - 20, 21 - 30, > 30 mm), localization (right upper, right middle, right lower, left upper, lingula, right lower), and component (soft tissue, ground glass, or mixed). If participants had multiple nodules, we would analyze the largest nodule as the main nodule.

Ascertainment of endpoints

The primary endpoint events of this study were lung cancer incidence and mortality. The NLST confirmed diagnosis of lung cancer through medical record abstraction (MRA), which was triggered by annual or semi-annual study update form, positive computed tomography (CT) or chest X-ray screening exam, direct report by relatives or physicians, and supplemented by NDI Plus searches. Active follow-up data were collected on cancer diagnoses and deaths that occurred through December 31, 2009. Extended follow-up data were collected for deaths through December 31, 2015. Furthermore, in this study, the primary outcomes were censored at the date of the lung cancer diagnosis, death, loss of follow-up, or end of the follow-up period, whichever came first.

Statistical analysis

The main purpose of this study was to determine the association of lung cancer risk with the presence of both lung nodules and emphysema. According to the diagnosis of emphysema and lung nodules based on the LDCT results of the first round (T0), we defined four groups in advance (NENN group (nonemphysema and non-nodules), E group (emphysema without nodules), N group (nodules without emphysema), and E + Ngroup (nodules with emphysema)) to evaluate the association between lung nodules combined with emphysema and lung cancer risk. The NENN group was used as the reference category. Descriptive analysis (variance and Chi-square test) was conducted to evaluate the distribution of baseline characteristics in different groups and the distribution of lung nodules in participants with or without emphysema. Log-rank test was originally used to compare the lung cancer incidence, lung cancer mortality and all-cause mortality of four groups. Multivariate Cox proportional hazard regression model was used to analyze the associations between the four groups on lung cancer incidence, lung cancer mortality and all-cause mortality. These associations were measured by hazard ratio (HR) and 95% confidence interval (CI). We constructed two models, namely, the age-adjusted model (model 1) only adjusted to age, and the fully adjusted model (model 2) adjusted to age, gender, race/ethnicity, education level, marital status, smoking status, smoking pack-year, BMI and family history of lung cancer.

Parameter	Overall (N = 21,949)	NENN group (n = 9,040)	E group (n = 5,819)	N group (n = 4,737)	E + N group (n = 2,353)	P value
Age (years)						< 0.001
≤ 59	8,920 (40.6)	3,981 (44.0)	2,243 (38.5)	1,904 (40.2)	792 (33.7)	
60 - 64	6,816 (31.1)	2,809 (31.1)	1,819 (31.3)	1,488 (31.4)	700 (29.7)	
65 - 69	4,130 (18.8)	1,545 (17.1)	1,149 (19.7)	902 (19.0)	534 (22.7)	
70 - 74	2,083 (9.5)	705 (7.8)	608 (10.4)	443 (9.4)	327 (13.9)	
Sex						< 0.001
Women	8,996 (41.0)	3,764 (41.6)	2,269 (39.0)	2,079 (43.9)	884 (37.6)	
Men	12,953 (59.0)	5,276 (58.4)	3,550 (61.0)	2,658 (56.1)	1,469 (62.4)	
Race						< 0.001
White	20,099 (91.6)	8,167 (90.3)	5,319 (91.4)	4,407 (93.0)	2,206 (93.8)	
Other	1,850 (8.4)	873 (9.7)	500 (8.6)	330 (7.0)	147 (6.2)	
Education						< 0.001
< high school	1,416 (7.7)	557 (7.5)	397 (7.9)	279 (7.0)	183 (9.1)	
Senior high school	8,336 (45.2)	3,230 (43.3)	2,290 (45.8)	1,859 (46.8)	957 (47.6)	
College or above	8,691 (47.1)	3,672 (49.2)	2,311 (46.2)	1,836 (48.2)	872 (43.3)	
Marital status						0.037
Married/living as married	14,741 (67.3)	6,028 (66.9)	3,879 (66.9)	3,226 (68.2)	1,608 (68.4)	
Widowed/divorced/separated	6,153 (28.1)	2,525 (28.0)	1,677 (28.9)	1,301 (27.5)	650 (27.7)	
Never married	995 (4.5)	457 (5.1)	242 (4.2)	204 (4.3)	92 (3.9)	
BMI (kg/m ²)						< 0.001
< 25	7,322 (33.5)	2,518 (28.0)	2,358 (40.6)	1,404 (29.7)	1,042 (33.5)	
25 - 29.9	8,864 (40.5)	3,724 (41.4)	2,319 (39.9)	1,909 (40.4)	912 (38.9)	
\geq 30	5,688 (26.0)	2,760 (30.7)	1,129 (19.4)	1,411 (29.9)	388 (16.6)	
Smoking status						< 0.001
Former smoking	11,246 (51.2)	5,020 (55.5)	2,643 (45.4)	2,572 (54.3)	1,011 (43.0)	
Current smoking	10,703 (48.8)	4,020 (44.5)	3,176 (54.6)	2,165 (45.7)	1,342 (57.0)	
Smoking, pack-years						< 0.001
30 - 59	14,352 (65.4)	6,276 (69.4)	3,476 (59.7)	3,222 (68.0)	1,378 (58.6)	
60 - 89	5,494 (25.0)	2,034 (22.5)	1,668 (28.7)	1,102 (23.3)	690 (29.3)	
≥ 90	2,103 (9.6)	730 (8.1)	675 (11.6)	413 (8.7)	285 (12.1)	
Family history of lung cancer						0.047
No	17,607 (80.2)	7,312 (80.9)	4,623 (79.4)	3,817 (80.6)	1,855 (78.8)	
Yes	4,342 (19.8)	1,728 (19.1)	1,196 (20.6)	920 (19.4)	498 (21.2)	

 Table 1. Baseline Characteristics of the Study Group

P values from the χ^2 test. BMI: body mass index; NENN group: non-emphysema and non-nodules; E group: emphysema without nodules; N group: nodules without emphysema; E + N group: nodules with emphysema.

We conducted subgroup analysis according to covariables (age, sex, smoking status, smoking pack-year, and family history of lung cancer) to identify the potential subgroups significantly related to the lung cancer risk among the four groups and confirm the robustness of the results.

All statistical analyses were performed using SPSS 23.0 software and R version 3.4.3. P value < 0.05 was considered statistically significant.

Results

Baseline characteristics of the participants according to study group

A total of 21,949 participants were included in this analysis (Fig. 1). There were 9,040 (41.2%) participants in the NENN

Parameter	All nodules (N = 7,090)	N group (n = 4,737)	E + N group (n = 2,353)	P value
Size, mm				< 0.001
4 - 6	3,707 (52.4)	2,557 (54.1)	1,150 (49.0)	
7 - 10	2,120 (30.0)	1,391 (29.4)	729 (31.1)	
11 - 20	937 (13.3)	599 (12.7)	338 (14.4)	
21 - 30	197 (2.8)	114 (2.4)	83 (3.5)	
> 30	109 (1.5)	64 (1.4)	45 (1.9)	
Localization				0.013
Right upper	1,733 (24.4)	1,137 (24.0)	596 (25.3)	
Right middle	971 (13.7)	669 (14.1)	302 (12.8)	
Right lower	1,681 (23.7)	1,145 (24.2)	536 (22.8)	
Left upper	1,019 (14.4)	638 (13.5)	381 (16.2)	
Lingula	304 (4.3)	218 (4.6)	86 (3.7)	
Left lower	1,347 (19.0)	908 (19.2)	439 (18.7)	
Other	35 (0.5)	22 (0.5)	13 (0.6)	
Margins				< 0.001
Spiculated	865 (12.2)	462 (9.8)	403 (17.1)	
Smooth	4,407 (62.2)	3,000 (63.3)	1,407 (59.8)	
Poorly defined	1,467 (20.7)	983 (20.8)	484 (20.6)	
Unable to determine	351 (5.0)	292 (6.2)	59 (2.5)	
Component				0.004
Soft tissue	5,182 (73.1)	3,417 (72.1)	1,765 (75.1)	
Ground glass	1,093 (15.4)	782 (16.5)	311 (13.2)	
Mixed	389 (5.5)	259 (5.5)	130 (5.5)	
Other	423 (6.0)	278 (5.9)	145 (6.2)	

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P values from the χ^2 test. N group: nodules without emphysema; E + N group: nodules with emphysema.

group, 5,819 (26.5%) participants in the E group, 4,737 (21.6%) participants in the N group, and 2,353 (10.7%) participants in the E + N group (Table 1). The average age of the participants was 61.66 (6.04) years old, 12,953 (59.0%) were male and 10,703 (48.8%) were current smokers. Of the 21,949 participants, 7,090 (32.3%) had lung nodules and 8,172 (37.2%) had emphysema.

Characteristics of nodules, overall and stratified by with and without emphysema

As shown in Table 2, of the 7,090 participants with lung nodules, 4,737 (66.8%) participants did not have emphysema and 2,353 (33.2%) participants had emphysema. Compared with the participants without emphysema, the participants with emphysema had larger nodules (7 - 10 mm: 29.4% vs. 31.1%, 11 - 20 mm: 12.7% vs. 14.4%, 21 - 30 mm: 2.4% vs. 3.5%, > 30 mm: 1.4% vs. 1.9%, P < 0.001). Participants with emphysema had a higher proportion of nodules in the upper lung than those without emphysema (right upper: 24.0% vs. 25.3%, left upper: 13.5% vs.

16.2%, P = 0.013). Participants with emphysema were more likely to have spiculated nodules than those without emphysema (9.8% vs. 17.1%, P < 0.001). Most participants with nodules had soft tissue nodules (73.1%, 5,128/7,090), followed by ground glass nodules (15.4%, 1,093/7,090) and partial solid nodules (5.5%, 389/7,090). This remained true when participants were divided into those with and without emphysema.

Kaplan-Meier survival curves of cumulative lung cancer incidence, lung cancer mortality, and all-cause mortality

Throughout follow-up, the cumulative incidence of lung cancer and cumulative mortality of lung cancer in the four groups had the same trend. The cumulative incidence of lung cancer (Fig. 2a) and the cumulative mortality of lung cancer (Fig. 2b) were the highest in the E + N group, followed by the N group, then the E group and the lowest in the NENN group. All-cause cumulative mortality was the highest in the E + Ngroup, followed by the E group, then N group and the lowest in the NENN group (Fig. 2c).



Figure 2. Kaplan-Meier survival curves of lung cancer incidence (a), lung cancer mortality (b) and all-cause mortality (c). NENN group: non-emphysema and non-nodules; E group: emphysema without nodules; N group: nodules without emphysema; E + N group: nodules with emphysema.

Forest plot of the adjusted HRs for the association between lung nodules combined with emphysema and lung cancer incidence, lung cancer mortality, and all-cause mortality

HR from the Cox proportional hazard regression model was used to evaluate the relationship between the incidence of lung cancer, mortality of lung cancer and all-cause mortality (Fig. 3). We provided two models: age-adjusted model (model 1) and fully adjusted model (model 2). It was found that the risk of lung cancer incidence increased in turn in NENN group, E group, N group and E + N group (Fig. 3). In the age-adjusted model (model 1), compared with NENN group, the HRs (95% CIs) of lung cancer incidence were 2.07 (1.69 - 2.54) for E group, 4.13 (3.47 - 5.05) for N group, and 6.26 (5.14 - 7.62) for E + N group. The association was robust to adjustment for

the age, sex, race, marital status, education level, BMI, smoking status, smoking package years and family history of lung cancer. In the fully adjusted model (model 2), compared with NENN group, adjusted HRs (95%CIs) of lung cancer incidence were 1.83 (1.47 - 2.27) for E group, 3.97 (3.24 - 4.86) for N group, and 5.23 (4.28 - 6.48) for E + N group.

Comparable results as the lung cancer incidence were observed for lung cancer mortality (Fig. 3). Compared with NENN group, the risk of lung cancer mortality increased in turn in E group (age-adjusted HR (model 1): 1.85 (1.39 - 2.46), fully adjusted HR (model 2): 1.56 (1.15 - 2.11)), N group (age-adjusted HR (model 1): 2.49 (1.89 - 3.29), fully adjusted HR (model 2): 2.43 (1.81 - 3.26)) and E + N group (age-adjusted HR (model 1): 4.27 (3.21 - 5.68), fully adjusted HR (model 2): 3.39 (2.50 - 4.61)), whether in age-adjusted model (model 1) or fully adjusted model (model 2).

Group	Events/Total (%)	Model1 HR (95	% CI) ^a	Model2 HR (95	%СІ) ^ь
Lung cancer incidence					
NENN group	159/9040(1.8)	1.00	•	1.00	+
E group	223/5819(3.8)	2.07(1.69-2.54)	I∳I	1.83(1.47,2.27)	II
N group	349/4737(7.4)	4.13(3.47-5.05)	⊢⊷⊣	3.97(3.24,4.86)	⊢∙-1
E + N group	272/2353(11.6)	6.26(5.14-7.62)	⊢← ⊣	5.23(4.28,6.48)	⊢← →
Lung cancer mortality					
NENN group	85/9040(0.9)	1.00	•	1.00	+
E group	109/5819(1.9)	1.85(1.39-2.46)	₩	1.56(1.15,2.11)	H o H
N group	120/4737(2.5)	2.49(1.89-3.29)	⊢⊷⊣	2.43(1.81,3.26)	⊢⊷⊣
E + N group	110/2353(4.7)	4.27(3.21-5.68)	⊢← -i	3.39(2.50,4.61)	
All-cause mortality					
NENN group	543/9040(6.0)	1.00	•	1.00	+
E group	512/5819(8.8)	1.37(1.21-1.54)	*	1.26(1.10,1.44)	•
N group	325/4737(6.9)	1.06(0.92-1.21)	+	1.09(0.94,1.27)	•
E + N group	287/2353(12.2)	1.75(1.51-2.02)	●	1.52(1.30,1.79)	lei
			12345678		1234567

Figure 3. Forest plot of associations of four groups participants with lung cancer incidence, lung cancer mortality and all-cause mortality. Model 1a: adjusted for age. Model 2b: adjusted for age, gender, race/ethnicity, education level, marital status, smoking status, smoking pack-year, BMI and family history of lung cancer. NENN group: non-emphysema and non-nodules; E group: emphysema without nodules; N group: nodules without emphysema; E + N group: nodules with emphysema; BMI: body mass index.

However, the trend of all-cause mortality risk among the four groups was somewhat different from that of lung cancer risk (Fig. 3). Compared with NENN group, E + N group (age-adjusted HR (model 1): 1.75 (1.51 - 2.02), fully adjusted HR (model 2): 1.52 (1.30 - 1.79)) had the highest risk of all-cause mortality, followed by E group (age-adjusted HR (model 1): 1.37 (1.21 - 1.54), fully adjusted HR (model 2): 1.26 (1.10 - 1.44)) and N group (age-adjusted HR (model 1): 1.06 (0.92 - 1.21), fully adjusted HR (model 2): 1.09 (0.94 - 1.27)). The association between lung nodules (N group) and all-cause mortality was not significant and had no statistical significance.

Subgroup analysis

Supplementary Materials 1 and 2 (www.wjon.org) reveal the subgroup analysis of the relationship between the incidence of lung cancer and the mortality of lung cancer in the four groups, respectively. We found that the trend of lung cancer risk among the four groups remained unchanged after stratification to age, sex, smoking status, smoking package year, and family history of lung cancer. That is to say, compared with NENN group, it was found that the risk of lung cancer in E group, N group and E + N group increased in turn, whether it is age-adjusted model or fully adjusted model. Subgroup analysis of all-cause mortality relationship of the four groups is shown in Supplementary Material 3 (www.wjon.org). The result of the highest all-cause mortality rate in the E + N group is reliable.

Discussion

In this study, we used the lung cancer screening LDCT data of NLST in the United States to focus on the association of lung cancer risk with the presence of both lung nodules and emphysema. This study revealed that the size, location and margins shape of lung nodules were different between participants with emphysema and those without emphysema. Among participants with lung nodules, participants with emphysema have larger lung nodules, and have a higher proportion of locating in the upper lobe of the lung and spiculated margins. What's more, compared with NENN group, lung cancer risk in the E group, N group and E + N group increased gradually. Lung cancer risk in NENN group was the lowest and that in E + N group is the highest, even if stratified according to age, sex, smoking status, smoking package year, and family history of lung cancer. However, the trend of all-cause mortality risk was different.

Many previous studies showed that lung nodules with the above characteristics are more likely to be malignant nodules or developing into lung cancer [29-34]. Gomez-Saez et al found that cancer risk was associated with smoking habit, nodule size and spiculated edge which was nearly significant and there was a linear relationship between nodule size and risk of lung cancer [35]. The recent Fleischer guidelines mention that nodules located in the (right) upper lobe of lung have been identified as independent risk factors for lung cancer, and have been added to the definition of "high-risk" nodules, even in low-risk individuals [34]. In a systematic literature review by Wahidi et al, irregular, spiculated, and lobulated margin were found to be predictive of malignancy [36]. Hence, our research results show that participants with lung nodules coexisting with emphysema have a higher lung cancer risk because of the more malignant nodule characteristics.

We found that there is a correlation between the existence of emphysema and the incidence and mortality of lung cancer, independent of age, sex, smoking history and lung nodules. Consistent with our findings, a systematic review and metaanalysis [37] revealed that emphysema detected visually at CT is associated with significantly increased odds of lung cancer. However, not all studies report a positive correlation between emphysema and lung cancer. Maldonado et al did not confirm the influence of emphysema on the diagnostic frequency of malignant nodules [38].

The role of lung nodules as a risk factor for lung cancer has been widely studied [39-41]. Beyond that, our study also found that the existence of lung nodules combined with emphysema can further increase the lung cancer risk. Similarly, a study mentioned that solitary lung nodules may have a greater probability of malignancy in the presence of emphysema [29].

As mentioned above, emphysema and lung nodules do increase the lung cancer risk. Interestingly, though the group with lung nodules (N group) has a higher lung cancer risk than the group with emphysema (E group), there is no such trend in all-cause mortality risk. Suffering from emphysema can increase the risk of all-cause mortality, and lung nodules have no obvious significance for all-cause mortality. Similarly, Lee's study [42] also revealed that emphysema was associated with long-term all-cause mortality; suspicious nodules were not identified as an independent risk factor for all-cause mortality in the multivariable Cox proportional hazards regression analysis. The mechanism that emphysema and lung nodules have different effects on lung cancer risk and all-cause mortality risk needs more research to explore.

Our study explored association of lung cancer risk with the presence of both lung nodules and emphysema, which provided a more solid foundation for the traditional risk prediction method of lung cancer. Currently, there are also more and more emerging methods to predict the lung cancer risk. Radiomic features, calculated based on LDCT images, and clinical data are frequently used for lung cancer screening and risk prediction, and some models [43, 44] have been independently validated. Developing alternative methods based on deep learning to perform feature extraction similar to radiology is also the main research focus of many teams. However, the interpretability of deep learning models should be taken into account during model development, and further research should be conducted in this field. In addition, biomolecular markers are also a hot method to predict the risk of lung cancer. At present, promising candidate molecules include autoantibodies, microRNAs, circulating tumor DNA and so on [45]. The emergence of genome-wide association studies (GWAS) may provide powerful evidence of genetic susceptibility genes for the lung cancer, and these genes may be included in the lung cancer risk prediction model [46]. Although no molecular biomarker has been routinely used for lung cancer screening, the existing candidate methods have great potential and represent a very promising method for early lung cancer screening in the future. In the future, if more related research can be done in the above field, we believe that this will provide more new insights for the occurrence and development mechanism and risk prediction of lung cancer.

Additionally, several limitations of this study should also be considered. First of all, the subtypes and severity of emphysema are not measured and classified in our data. Secondly, we have not explored the mechanism of the result that lung nodules have a higher lung cancer risk than emphysema. Thirdly, we did not explore the etiological relationship between emphysema and lung nodules. Fourth, the participants in this study are heavy smokers, and the generalizability of our results to populations outside the NLST eligibility criteria is indeterminate. Fifth, this study enrolled mainly (> 90%) Caucasian patients, so the results may not be applicable to all populations at risk for lung cancer. Finally, at present, radiomics and deep learning algorithms are significantly improving the sensitivity of predicting lung cancer risk from lung nodule imaging, but the radiographic descriptions of the lung nodules are also fairly limited in this study. Though they have limitations in population promotion, they are currently recommended high-risk screening groups, and the results of this study have targeted guiding significance for screening.

Conclusion

Based on a large-scale lung cancer screening trial in the United States, this study demonstrated that either emphysema or lung nodules can increase lung cancer risk, and lung nodules combined with emphysema can further increase the lung cancer risk and all-cause mortality. Compared with NENN group, lung cancer risk in the E group, N group and E + N group increased gradually. In addition, among participants with lung nodules, participants with emphysema have more malignant characteristics of lung nodules than those without emphysema. The significance of these findings for lung cancer screening should be evaluated.

Supplementary Material

Suppl 1. Subgroup Analysis Between Four Groups Participants and Lung Cancer Incidence.

Suppl 2. Subgroup Analysis Between Four Groups Participants and Lung Cancer Mortality.

Suppl 3. Subgroup Analysis Between Four Groups Participants and All-Cause Mortality.

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Conflict of Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Informed Consent

Each participating center's institutional review board approved the protocols and all participants provided written informed consent.

Author Contributions

Ya Liu: conceptualization, methodology, data curation, formal analysis, investigation, writing - original draft, writing - review and editing. Zhuo Wei Feng: data curation, investigation, writing - review and editing. Xiao Min Liu: data curation, investigation, writing - review and editing. Hong Yuan Duan: data curation, investigation, writing - review and editing. Zhang Yan Lyu: data curation, writing - review and editing. Yu Bei Huang: data curation, investigation, writing - review and editing, supervision. Fang Fang Song: data curation, writing - review and editing. Feng Ju Song: conceptualization, data curation, investigation, writing - review and editing, supervision, funding acquisition.

Data Availability

The data are available on application to the NLST trial (https:// cdas.cancer.gov/nlst/).

Abbreviations

BMI: body mass index; CI: confidence interval; COPD: chronic obstructive pulmonary disease; HR: hazard ratio; LDCT: low-dose computed tomography; NLST: National Lung Screening Trial

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