

The Ability of Bilirubin in Identifying Smokers with Higher Risk of Lung Cancer: A Large Cohort Study in Conjunction with Global Metabolomic Profiling

Chi-Pang Wen^{1,2}, Fanmao Zhang³, Dong Liang⁴, Christopher Wen⁵, Jian Gu³, Heath Skinner⁶, Wong-Ho Chow³, Yuanqing Ye³, Xia Pu³, Michelle A.T. Hildebrandt³, Maosheng Huang³, Chien-Hua Chen⁷, Chao Agnes Hsiung¹, Min Kuang Tsai¹, Chwen Keng Tsao⁸, Scott M. Lippman⁹, and Xifeng Wu³

Abstract

Purpose: We aimed to identify serum metabolites as potential valuable biomarkers for lung cancer and to improve risk stratification in smokers.

Experimental Design: We performed global metabolomic profiling followed by targeted validation of individual metabolites in a case-control design of 386 lung cancer cases and 193 matched controls. We then validated bilirubin, which consistently showed significant differential levels in cases and controls, as a risk marker for lung cancer incidence and mortality in a large prospective cohort composed of 425,660 participants.

Results: Through global metabolomic profiling and following targeted validation, bilirubin levels consistently showed a statistically significant difference among healthy controls and lung cancer cases. In the prospective cohort, the inverse asso-

ciation was only seen in male smokers, regardless of smoking pack-years and intensity. Compared with male smokers in the highest bilirubin group (>1 mg/dL), those in the lowest bilirubin group (<0.75 mg/dL) had 55% and 66% increase in risks of lung cancer incidence and mortality, respectively. For every 0.1 mg/dL decrease of bilirubin, the risks for lung cancer incidence and mortality increased by 5% and 6% in male smokers, respectively (both $P < 0.001$). There was a significant interaction between low serum bilirubin level and smoking on lung cancer risk ($P_{\text{interaction}} = 0.001$).

Conclusion: Low levels of serum bilirubin are associated with higher risks of lung cancer incidence and mortality in male smokers and can be used to identify higher risk smokers for lung cancer. *Clin Cancer Res*; 1–8. ©2014 AACR.

Introduction

Lung cancer is the second most common cancer and the leading cause of cancer deaths in both men and women in the United States (1). Recent studies by the National Lung Screening Trial (NLST) have showed that low-dose CT (LDCT) can reduce lung cancer mortality by 20% (2). On the basis of these findings, LDCT screening based on NLST selection criteria, that is, current or

former smokers ages 55 to 74 years with at least 30 pack-years of smoking history and no more than 15 years since quitting, has been recommended by the majority of professional organizations in the United States (1, 3–5). Moreover, it has recently been reported that participants with the highest risk for lung cancer deaths accounted for the most screening-prevented lung-cancer deaths and benefitted most from LDCT (6). Although smoking is the predominant risk factor for lung cancer, considering smoking alone is not sufficient to identify the highest-risk individuals for lung cancer (3, 6). Therefore, novel biomarkers for lung cancer incidence and mortality, particularly among smokers, are urgently needed in the clinical setting to improve risk prediction and reduce false positives of LDCT screening.

Metabolomics is the systematic study of the unique chemical fingerprints generated by metabolic processes of an organism (7). Metabolomic profiling, emerging as an important tool to identify biomarkers, provides a functional readout of physiologic and pathologic characteristics (8). An increasing number of studies have utilized metabolomic profiling to reveal metabolic alterations associated with various cancers (8–16), including lung cancer (17–19). However, only a small number of metabolites have been examined and studies to date suffer from a lack of prospective validation (17–19).

To identify serum metabolites as novel biomarkers for lung cancer, we first performed metabolomic profiling followed by targeted metabolite validation in a lung cancer case-control study with three phases to identify top promising metabolites that

¹Institute of Population Health Science, National Health Research Institutes, Zhunan, Taiwan. ²China Medical University Hospital, Taichung, Taiwan. ³Department of Epidemiology, The University of Texas MD Anderson Cancer Center, Houston, Texas. ⁴Department of Pharmaceutical Sciences, College of Pharmacy and Health Sciences, Texas Southern University, Houston, Texas. ⁵Department of Radiological Sciences, University of California at Irvine, Irvine, California. ⁶Department of Radiation Oncology, The University of Texas MD Anderson Cancer Center, Houston, Texas. ⁷Digestive Disease Center, Show-Chwan Memorial Hospital, Changhua, Taiwan. ⁸MJ Health Management Institution, Taipei, Taiwan. ⁹UC San Diego Moores Cancer Center, San Diego, California.

C.-P. Wen and F. Zhang contributed equally to article as first authors.

S.M. Lippman and X. Wu contributed equally to this article as senior authors.

Corresponding Author: Xifeng Wu, Department of Epidemiology, Unit 1340, The University of Texas MD Anderson Cancer Center, 1155 Pressler Blvd., Houston, TX 77030. Phone: 713-745-2485; Fax: 713-745-1165; E-mail: xwu@mdanderson.org

doi: 10.1158/1078-0432.CCR-14-0748

©2014 American Association for Cancer Research.

Translational Relevance

Through this multiphase study composed of global metabolomic profiling and prospective validation in a large cohort, we have identified and validated bilirubin as a risk predictor for lung cancer incidence as well as mortality in smokers. For every 0.1 mg/dL decrease of bilirubin, the risks for lung cancer incidence and mortality increased by 5% and 6% in male smokers, respectively (both $P < 0.001$). Smokers with ≥ 30 pack years had a 4-fold increase in lung cancer risk, but within this group, those with bilirubin of < 0.75 mg/dL compared with > 1 mg/dL had a 31% higher risk. Addition of this variable into National Lung Screening Trial (NLST) selection criteria for low-dose CT (LDCT) screening might help identify higher risk smokers who would benefit more from LDCT screening and reduce false positives.

differentiated lung cancer cases from healthy controls. Bilirubin emerged as the consistently significant metabolite. We then sought to validate bilirubin as a risk marker for lung cancer in a large prospective cohort study. During this validation stage, we prospectively analyzed serum bilirubin levels in a cohort of 425,660 subjects to assess its ability in identifying smokers with particularly high risk for lung cancer.

Materials and Methods

Study population

Stage 1: Case-control studies.

The subjects are participants in an ongoing lung cancer case-control study at the University of Texas MD Anderson Cancer Center (Houston, TX). Details of subject recruitment methods have been reported previously (20). Cases were newly diagnosed, histologically confirmed non-small cell lung cancer (NSCLC) patients previously untreated with chemotherapy or radiotherapy at MD Anderson Cancer Center. There was no restriction of age, sex, or ethnicity at study recruitment. Early-stage NSCLC included stages I and II, whereas late-stage NSCLC included stages III and IV. The healthy controls came from the Kelsey Seybold Clinics, the largest private multispecialty physician group practice in Houston. To control for the confounding effect of ethnicity, we only included Caucasians for our study. Twenty each of controls, early-stage, and late-stage lung cancer cases (hereafter referred to as "trio") were used for metabolomic profiling. Promising metabolites identified from this profiling were examined in two additional sets of case-control samples, consisting of 50 trios and 123 trios, respectively. All participants completed an in-person interview using a structured questionnaire. Demographic characteristics, smoking history, family history of cancer, and exposure data were collected. After the interview, each participant donated 40 mL blood sample for molecular analysis.

Stage 2: Prospective cohort study.

The cohort consisted of 425,660 Taiwanese adults ages 20 years and older who participated in a standard medical screening program between 1994 and 2008. Details of this cohort have been reported (21, 22). Briefly, median follow-up time for the cohort is 8 years (interquartile range: 5–11 years) for male

participants and 9 years (interquartile range: 5–11 years) for female participants. All participants completed a self-administered health history questionnaire and underwent a series of medical tests for blood, urine, physical examination, body measurements, and functional tests. Overnight fasting blood was analyzed for a standard panel of markers, including serum bilirubin. The cohort members were followed through 2008 for cancer and vital status, which were assessed by linkage of the individual ID to the National Cancer Registry and National Death file.

The studies were approved by the Institutional Review Boards of the University of Texas MD Anderson Cancer Center and Kelsey Seybold Clinics, as well as the National Health Research Institutes, Zhunan, Taiwan. All participants provided written informed consent.

Metabolomic profiling and quantification of individual metabolites

The metabolomic profiling analysis was carried out by Metabolon Inc, as described previously (23). Internal controls included injection, process, and alignment standards for quality assurance/quality control (QA/QC) procedures to control for experimental variability. Samples were kept at -80°C until assays were performed. For the series of validation studies, standard powders for two metabolites, that is, λ -glutamylalanine and bilirubin, were purchased from Sigma-Aldrich. Quantification of individual metabolite in serum was determined by LC/MS-MS methods using a 3200 QTRAP MS/MS coupled by an Agilent 1200 Series HPLC system at Dr. Dong Liang's laboratory. Standard curves for each compound were constructed by spiking known amount of the standard to a series of control plasma (Gulf Coast Blood Bank). Serum bilirubin levels measured by both metabolomic profiling and LC/MS-MS were levels of unconjugated bilirubin in serum.

Statistical analysis

In the case-control analysis, the Pearson χ^2 test was used to examine the differences in sex and smoking status between cases and controls. Student t test was used to test for differences in age and pack-years of smoking as continuous variables. For the metabolomic profiling analysis, missing metabolite measurements were imputed with the compound minimum (minimum value imputation). Only metabolites with detectable expression in at least 80% of the samples were kept for further analysis. For both metabolomics profiling and individual metabolite quantification, the nonparametric trend test was used to analyze the trend across the trios. Bonferroni correction was used to account for multiple comparisons from metabolomic profiling results, and a P value $< 0.05/n$ (n = number of comparisons) was considered as the significance level to take into account multiple comparisons. Spearman correlation was used to assess the correlation between the two values measured by metabolomic profiling and individual metabolite quantification using LC/MS-MS.

For the prospective cohort validation study, lung cancer cases diagnosed within one year of recruitment into the cohort were excluded to minimize potential reverse causality. For lung cancer incidence, the event time was from the date of recruitment to the end of follow-up, or the date of lung cancer identification if earlier. For lung cancer mortality, the event time was from the date of recruitment to the end of follow-up, or the date

of death due to lung cancer if earlier. Serum total bilirubin levels were divided into three groups with equal tertile. Cox proportional hazards models were used to assess the association of serum total bilirubin levels with lung cancer incidence and mortality. HRs were adjusted for age, educational level (middle school or lower, high school, junior college, or college or higher), body mass index (BMI), and pack-years of smoking in a multivariable model with continuous variables whenever appropriate. The proportional hazards assumption was assessed by plotting Schoenfeld residuals versus time and examining their correlation. Interaction between smoking and serum total bilirubin level on lung cancer risk was assessed by introducing the product of smoking and serum bilirubin level in the multivariable Cox regression model. All statistical tests were two sided with the threshold for significance set at 0.05. Statistical analyses were performed using Stata 10.0 (StataCorp).

Results

Characteristics of the study populations

In the case-control study, all three phases of lung cancer cases and healthy controls were Caucasians, matched on age and gender (Supplementary Table S1). In the prospective cohort study, there were 202,902 men and 222,758 women ages 20 years and older. Selected demographic characteristics and exposures of the cohort participants are shown in Table 1, presented by gender and tertiles of bilirubin level (<0.75, 0.75–1, and >1 mg/dL for men and <0.61, 0.61–0.82, and >0.82 mg/dL for women). Distribution of serum total bilirubin levels among the participants in the cohort is shown in Supplementary Fig. S1. Among male participants in the cohort, over half (52.1%) were smokers, with 25% of them being heavy smokers of ≥ 30 pack-years. In contrast, only 17,123 (8.3%) female participants were smokers, with 1,327 (8.3%) of them being heavy smokers. During the follow-up, there were 809

incident lung cancer cases and 614 lung cancer deaths among the males, and 524 lung cancer cases and 330 deaths among the females.

Global metabolomic profiling of lung cancer

Serum global metabolomic profiles of 40 lung cancer cases and 20 healthy controls (20 trios) were assessed in the initial case-control study and a total of 403 metabolites were identified. After exclusion of metabolites detected in less than 80% of samples, 306 (76%) metabolites remained. These metabolites were mapped to eight super-pathways and 61 sub-pathways (Supplementary Table S2). Among these, 29 metabolites exhibited a significant trend of expression when comparing normal controls, early-, and late-stage cases, 12 of which had P_{trend} values < 0.01 (Supplementary Table S3). After taking into account multiple comparisons, λ -glutamylalanine remained as the only metabolite meeting statistical significance after Bonferroni correction [$P_{\text{trend}} < 1.63 \times 10^{-4}$ (0.05/306)].

Target validation of individual metabolites

Metabolites exhibiting a significant trend in levels from normal individuals to early- and late-stage patients are also potential biomarkers for the detection and prognosis of lung cancer. Of the 29 metabolites with significant trends, bilirubin caught our most interest given its potent endogenous cytoprotective properties and more importantly, its inverse association with cardiovascular disease and respiratory disease in previous reports (24–27). Therefore, we selected bilirubin and λ -glutamylalanine, which showed the most significant trend from metabolomic profiling and after Bonferroni correction for further validation. We developed standard LC/MS-MS assays for these metabolites and used these assays to measure their levels in the 20 trios of cases and controls from phase I of the case-control study; we found excellent correlation with metabolomic profiling data (Supplementary Tables S4 and S5). We further examined levels of bilirubin and λ -glutamylalanine in

Table 1. Characteristics of the participants in the prospective cohort by gender and serum total bilirubin levels^a

Characteristics	Men (N = 202,902), N (%)				Women (N = 222,758), N (%)			
	Total	Total bilirubin level (mg/dL)			Total	Total bilirubin level (mg/dL)		
		>1	0.75–1	<0.75		>0.82	0.61–0.82	<0.61
Total	202,902	67,841 (33.4)	65,540 (32.3)	69,521 (34.3)	222,758	75,189 (33.8)	72,207 (32.4)	75,362 (33.8)
Age (y), mean (SD)	41 (14)	40 (14)	42 (14)	41 (14)	41 (14)	41 (14)	42 (14)	41 (13)
20–39	112,584 (55.5)	39,399 (35.0)	35,270 (31.3)	37,915 (33.7)	119,946 (53.9)	41,854 (34.9)	37,510 (31.3)	40,582 (33.8)
40–59	63,447 (31.3)	19,927 (31.4)	21,201 (33.4)	22,319 (35.2)	76,087 (34.2)	23,908 (31.4)	25,500 (33.5)	26,679 (35.1)
≥ 60	26,871 (13.2)	8,515 (31.7)	9,069 (33.8)	9,287 (34.6)	26,725 (12)	9,427 (35.3)	9,197 (34.4)	8,101 (30.3)
BMI (kg/m ²), mean (SD)	23.9 (3.4)	23.5 (3.3)	23.9 (3.3)	24.2 (3.4)	22.3 (3.6)	21.8 (3.5)	22.3 (3.6)	22.8 (3.7)
<25	134,591 (66.4)	47,499 (35.3)	43,264 (32.1)	43,828 (32.6)	176,567 (79.3)	62,056 (35.2)	57,139 (32.4)	57,372 (32.5)
25–29.9	59,734 (29.5)	18,062 (30.2)	19,565 (32.8)	22,107 (37.0)	38,454 (17.3)	11,040 (28.7)	12,660 (32.9)	14,754 (38.4)
≥ 30	8,516 (4.2)	2,264 (26.6)	2,696 (31.7)	3,556 (41.8)	7,689 (3.5)	2,080 (27.1)	2,394 (31.1)	3,215 (41.8)
Educational levels								
Middle school or lower	40,499 (20.6)	12,823 (31.7)	13,379 (33.0)	14,297 (35.3)	70,385 (32.6)	23,109 (32.8)	23,546 (33.5)	23,730 (33.7)
High school	45,601 (23.2)	14,665 (32.2)	14,693 (32.2)	16,243 (35.6)	54,124 (25.1)	17,238 (31.9)	17,186 (31.8)	19,700 (36.4)
Junior college	45,367 (23.1)	15,705 (34.6)	14,526 (32.0)	15,136 (33.4)	42,941 (19.9)	15,153 (35.3)	13,525 (31.5)	14,263 (33.2)
College or higher	64,987 (33.1)	22,603 (34.8)	20,844 (32.1)	21,540 (33.2)	48,400 (22.4)	17,428 (36.0)	15,692 (32.4)	15,280 (31.6)
Smoking status								
Non-smoker	92,864 (47.9)	35,175 (37.9)	30,188 (32.5)	27,501 (29.6)	188,685 (91.7)	64,488 (34.2)	61,534 (32.6)	62,663 (33.2)
Smoker	101,092 (52.1)	29,632 (29.3)	32,451 (32.1)	39,009 (38.6)	17,123 (8.3)	4,891 (28.6)	5,228 (30.5)	7,004 (40.9)
<30 pack-years	72,153 (74.9)	21,843 (30.3)	23,084 (32.0)	27,226 (37.7)	14,662 (91.7)	4,303 (29.4)	4,403 (30.0)	5,956 (40.6)
≥ 30 pack-years	24,146 (25.1)	6,269 (26.0)	7,777 (32.2)	10,100 (41.8)	1,327 (8.3)	279 (21)	434 (32.7)	614 (46.3)
Lung cancer incidence	809 (0.4)	215 (26.6)	270 (33.4)	324 (40.1)	524 (0.2)	155 (29.6)	187 (35.7)	182 (34.7)
Lung cancer mortality	614 (0.3)	147 (23.9)	214 (34.9)	253 (41.2)	330 (0.2)	107 (32.4)	115 (34.9)	108 (32.7)

^aPercentage may not total 100 because of rounding.

additional 50 trios of serum samples (phase II) and 123 trios of serum samples (phase III) from controls, early-, and late-stage patients (Supplementary Table S5). Through this process, bilirubin emerged as a metabolite that consistently showed a statistically significant trend in all three sets of trio data.

Validation of bilirubin as a lung cancer marker in a large cohort

Because bilirubin is a routine blood test in health examination, we next assessed the association of blood test serum total bilirubin levels with lung cancer incidence and mortality using a large prospective cohort in Taiwan. As expected, there was a strong dose-response relationship between lung cancer risk/mortality and pack-years of smoking or smoking intensity in this cohort (Tables 2 and 3). Furthermore, among males, using nonsmokers with the highest tertile of bilirubin levels (>1 mg/dL) as reference, smokers in the lowest tertile of bilirubin levels (<0.75 mg/dL) had a 2.86-fold increased risk of developing lung cancer (Table 2). Smokers with <30 and ≥30 pack-years of smoking in the lowest tertile of bilirubin levels had HRs of 1.40 and 4.14, respectively (Table 2 and Supplementary Fig. S2). Similarly, smokers in the lowest tertile of bilirubin levels who smoked <10, 10 to 19, and ≥20 cigarettes per day had HRs of 1.85, 2.70, and 4.32, respectively (Table 2 and Supplementary Fig. S2). Similar results were found for lung cancer mortality (Table 3 and Supplementary Fig. S2). In contrast, among females, lower serum bilirubin levels were not significantly associated with lung cancer incidence or mortality overall, in female smokers or in female nonsmokers (Supplementary Table S6). Table 4 presents the rates of lung cancer incidence and mortality stratified by tertiles of serum bilirubin levels and corresponding risk estimates in males. The incidence rate of lung cancer per 10,000 person-years was 6.93 [95% confidence intervals (CI), 6.20–7.75] in the lowest tertile compared with 4.27 (95% CI, 3.71–4.90) in the highest tertile of bilirubin levels, which translated to a 52% increased risk of lung cancer for the low bilirubin group ($P < 0.001$). The corresponding lung cancer-specific mortality rate was 4.88 (95% CI, 4.32–5.52) in the lowest tertile compared with 2.70 (95% CI, 2.30–3.17) in the highest tertile, a 71% increased risk in lung cancer-specific mortality for the low bilirubin group ($P < 0.001$; Table 4). We plotted the lung cancer incidence rates against subgroups of bilirubin levels and introduced a best-fit model. Those with bilirubin levels <0.42 mg/dL showed more than 80% increase in lung cancer incidence rate (6.1 vs. 3.27 per 100,000 person-years; Fig. 1A) and over 2-folds increase in mortality rate (4.09 vs. 1.94 per 100,000 person-years; Fig. 1B) compared with the subgroup with bilirubin levels >1.62 mg/dL.

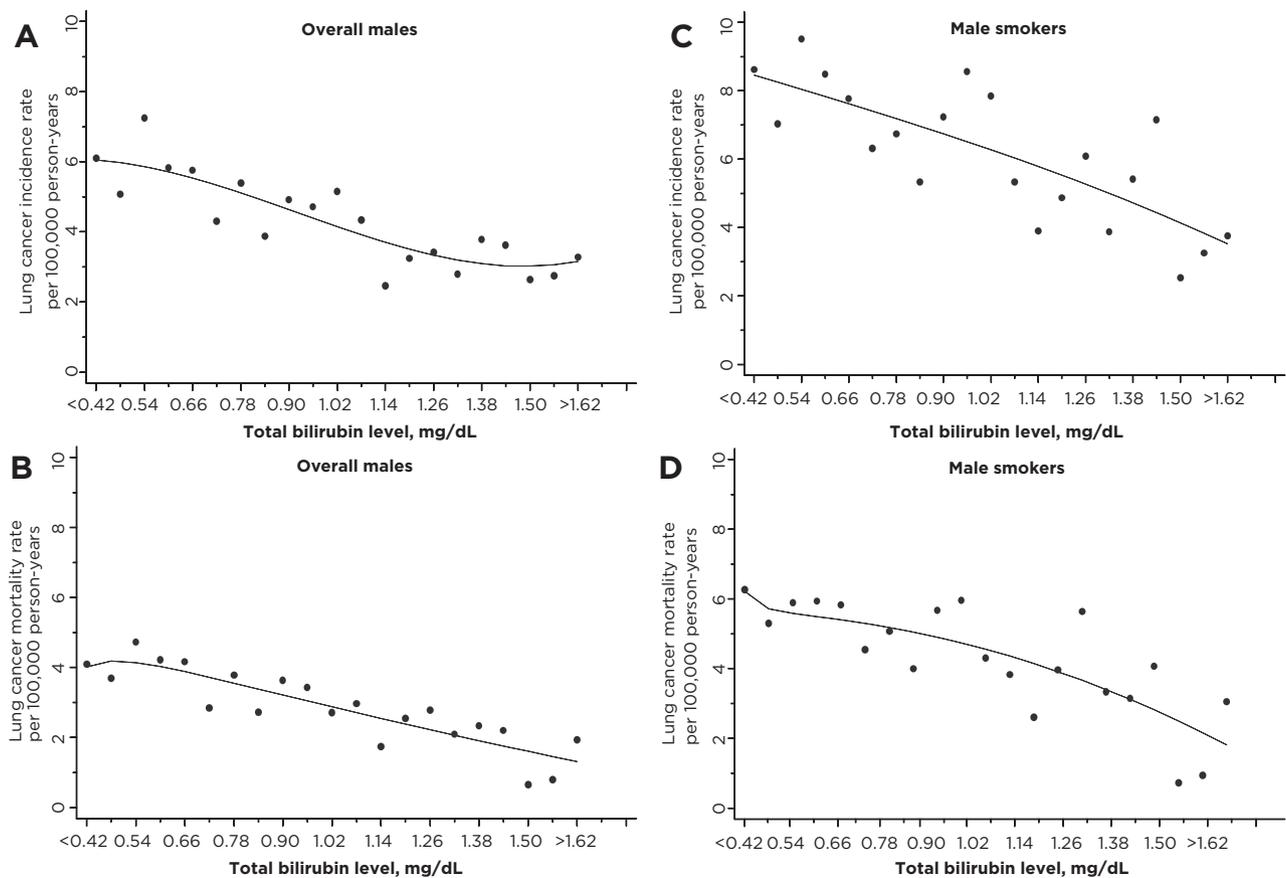
The ability of bilirubin in identifying smokers with higher risk of lung cancer

We then assessed the association between bilirubin levels and lung cancer incidence or mortality rate stratified by smoking status. Among females, neither nonsmokers or smokers showed significant association, as only 17,123 (8.3%) participants were smokers and there were only 37 lung cancer cases among them. Among males, the association was only present in smokers and there was a significant interaction between low serum bilirubin level and smoking on lung cancer risk ($P_{\text{interaction}} = 0.001$). Compared with smokers with bilirubin levels in the highest tertile, smokers with bilirubin levels in the middle and lowest

Table 2. Relationship among smoking quantity, bilirubin levels, and risk for lung cancer incidence in male participants in the prospective cohort study

	Men (N = 202,902)															
	Total				>1				0.75–1				<0.75			
	No.	No. of incidence	HR ^a (95% CI)	No. of incidence	No.	No. of incidence	HR ^a (95% CI)	No.	No. of incidence	HR ^a (95% CI)	No.	No. of incidence	HR ^a (95% CI)			
Nonsmoker	92,864	156	1 (Ref.)	35,175	64	1 (Ref.)	30,188	50	0.87 (0.59–1.27)	27,501	42	0.85 (0.56–1.27)				
Total smokers	101,092	603	2.64 (2.19–3.18)	29,632	139	1.84 (1.35–2.51)	32,451	202	2.38 (1.77–3.19)	39,009	262	2.86 (2.15–3.81)				
Pack-year																
<30 pack-years	72,153	123	1.31 (1.02–1.69)	21,843	27	0.80 (0.50–1.29)	23,084	47	1.37 (0.92–2.03)	27,226	49	1.40 (0.94–2.07)				
≥30 pack-years	24,146	454	4.01 (3.27–4.91)	6,269	108	3.14 (2.25–4.36)	7,777	145	3.48 (2.54–4.77)	10,100	201	4.14 (3.06–5.60)				
# of Cigarettes per day																
<10	31,520	106	1.55 (1.20–2.01)	10,602	23	0.96 (0.58–1.57)	10,270	37	1.39 (0.91–2.13)	10,648	46	1.85 (1.24–2.74)				
10–19	36,866	261	2.71 (2.19–3.34)	11,031	62	1.97 (1.36–2.84)	12,557	91	2.58 (1.84–3.60)	15,278	108	2.70 (1.95–3.72)				
≥20	26,879	221	4.29 (3.46–5.33)	6,759	52	3.39 (2.31–4.96)	8,374	70	3.72 (2.61–5.30)	11,746	99	4.32 (3.11–5.99)				

^aAdjusted for age, educational level, and BMI.

**Figure 1.**

Serum total bilirubin levels and lung cancer incidence rates in overall males (A) and male smokers (C); and lung cancer mortality rates in overall males (B) and male smokers (D) of the prospective cohort study.

tertiles had significantly increased lung cancer risk (HRs, 1.29 and 1.55) and mortality (HRs, 1.37 and 1.66; Table 4). The risk appeared to be stronger in light smokers: the HRs for the lowest tertile of bilirubin compared with the highest tertile were 1.77 for incidence and 2.56 for mortality in smokers of <30 pack-years and 1.31 for incidence and 1.32 for mortality in smokers of ≥ 30 pack years, respectively (Table 4). We also plotted the lung cancer incidence and mortality rates against subgroups of bilirubin levels in smokers and introduced a best fit model, those with bilirubin levels <0.42 mg/dL showed more than 2-folds increase in both lung cancer incidence rate (8.62 vs. 3.76 per 100,000 person-years; Fig. 1C) and mortality rate (6.27 vs. 3.05 per 100,000 person-years; Fig. 1D) compared with the subgroup with bilirubin levels >1.62 mg/dL. The logistic regression model showed a 5% (95% CI, 3%–8%, $P < 0.001$) increase in lung cancer incidence and 6% (95% CI, 3%–9%, $P < 0.001$) increase in lung cancer mortality per 0.1 mg/dL decrease in bilirubin level, after adjusting for age, BMI, and educational level.

Discussion

The purpose of this study is to identify biomarkers among serum metabolites to assist in identifying high-risk individuals for lung cancer development. Through this multistage study,

we have identified and validated serum bilirubin as a risk predictor for lung cancer incidence as well as mortality in male smokers. Although smoking is a strong risk factor for lung cancer and shows a dose–response relationship, the smoking-related risk is particularly high among male smokers with low levels of serum bilirubin, a 55% increase among those with bilirubin <0.75 mg/dL. Among males, smokers with ≥ 30 pack-years had a 4-fold increase in lung cancer risk, and within this group, those with bilirubin level <0.75 mg/dL had a 31% higher risk compared with those with bilirubin level >1 mg/dL. The potential of using serum bilirubin to identify smokers at particularly high-risk for lung cancer, over and above the risk associated with heavy smoking, is an important observation. The inverse relationship between bilirubin levels and lung cancer can be translated into a 5% increase in lung cancer risk and a 6% increase in lung cancer mortality for each 0.1 mg/dL decrease in bilirubin levels. In most clinical settings, emphasis is placed on elevated bilirubin for diagnosis of liver diseases, therefore, low values of bilirubin are generally ignored. Making use of low serum bilirubin values to counsel heavy smokers who are at particularly high risk for lung cancer about smoking cessation can be carried out easily in many clinic settings.

Elevated levels of serum bilirubin have been associated with a lower risk of respiratory diseases and lung cancer (24, 27). The

mechanism of this association was credited to the antioxidant and anti-inflammatory properties of bilirubin. As bilirubin is a commonly ordered laboratory test, uncovering this potentially protective relationship is intriguing. This study, while in line with the reported conclusion, is the first to study the role of bilirubin as a risk factor for lung cancer mortality, to focus on the analysis in smokers in detail, and to quantify the hazards of low bilirubin.

It has been shown that smoking is associated with lower serum bilirubin levels (27–29). In our study, we have also found that serum bilirubin levels were lower in smokers compared with nonsmokers among participants in the cohort. However, the inverse association between serum bilirubin levels and lung cancer incidence/mortality remained significant after we adjusted for smoking status/pack-years among overall male participants in the cohort. We also found that lower bilirubin was associated with higher risks of lung cancer and mortality among male smokers overall, and among male smokers with similar pack-years of smoking through our stratified analyses, suggesting that bilirubin level is associated with lung cancer risk at least partially independent of smoking status/quantity. In addition, we have also found a significant interaction between low serum bilirubin level and smoking on lung cancer risk ($P_{\text{interaction}} = 0.001$), suggesting that bilirubin may exert its function by interacting with smoking and lowering lung cancer risk among smokers who have higher oxidative stress and inflammation (30).

Our findings may also have implications for the LDCT screening for lung cancer. It has been reported that LDCT screening prevented most deaths from lung cancer among participants with the highest risk for lung cancer deaths—60% of participants at the highest risk accounted for 88% of prevented lung-cancer deaths (7). On the basis of our results, male smokers with bilirubin level <0.75 mg/dL have a 66% increased risk for lung cancer mortality compared with those with bilirubin level >1 mg/dL, and for heavy smokers of ≥ 30 pack-years, the HR is smaller, but still significant (HR = 1.32, $P < 0.001$). Consideration of bilirubin levels might improve identifying participants with the highest risk for lung cancer mortality who would benefit the most from the screening, and help improve the specificity of LDCT screening. Furthermore, bilirubin results could be used to target and motivate both light and heavy smokers for smoking cessation. Indeed, the ability of low bilirubin in predicting high risk of lung cancer was not limited to male smokers with ≥ 30 pack-years in our study. The relationship was seen for all male smokers, regardless of pack-years of smoking.

We conducted a series of sensitivity analyses to strengthen our conclusion. We excluded participants with lung cancer diagnosed within 3 years of cohort enrollment. We restricted bilirubin levels within normal range, excluding participants with abnormal liver enzymes or blood counts. Additional variables (drinking status, physical activity, and systolic blood pressure) were adjusted in the multivariable models. Results essentially remained unchanged after all of the above sensitivity analyses were carried out.

Recently, several research groups had applied metabolomic profiling of serum/plasma to unveil metabolic alterations associated with lung cancer, but all were limited by the small number of metabolites detected. Hori and colleagues' study detected a total of only 58 metabolites in serum using gas

chromatography/mass spectrometry and found 23 with differential detection in patients with lung cancer compared with healthy controls in a Japanese population (17). In another Japanese study, Maeda and colleagues studied 21 plasma amino acids in patients with NSCLC by LC/MS and showed that differences in the amino acid profiles may be used for screening NSCLC (19). Jordan and colleagues used nuclear magnetic resonance to measure 21 metabolites and showed the potential of serum metabolomics to differentiate between lung cancer subtypes and between patients and controls (18). These studies were limited by the small number of metabolites detected. Our global unbiased metabolomic profiling approach identified 403 known metabolites from different stages of lung cancer, yielding a comprehensive picture of the metabolic profile changes associated with cancer progression. Validated with two additional study sets, bilirubin was found and confirmed as the consistently significant biomarker for lung cancer, which was further validated prospectively in a large cohort.

A few potential limitations should be considered in the interpretation of our findings. First, although we observed significant inverse associations between serum bilirubin levels and lung cancer in male smokers, the associations were not statistically significant in female smokers, which was most likely due to the lack of power resulting from a small number of female smokers (8.3% of total females) and very few number of lung cancer cases ($n = 37$) among them. Second, although we observed an inverse relationship between bilirubin levels and lung cancer risk, the causality of the association remains unclear. Low bilirubin level could be a consequence of cancer rather than a predisposing factor. It is noteworthy that the significant risk remained after we excluded lung cancer occurring within 3 years of the bilirubin tests. Third, only the bilirubin data at the time of enrollment were analyzed. In a subset of subjects that had two bilirubin tests performed longitudinally, we found highly correlative data, implying the stability of total bilirubin results over time.

In summary, low levels of serum bilirubin are associated with higher risk for lung cancer incidence and mortality in male smokers and can be used to identify higher risk smokers for lung cancer development and mortality. Future prospective studies that incorporate this variable into NLST selection criteria to fully assess its potential use for LDCT screening are warranted.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

Authors' Contributions

Conception and design: C.P. Wen, D. Liang, C.-H. Chen, X. Wu, S.M. Lippman
Development of methodology: C.P. Wen, F. Zhang, D. Liang, H.D. Skinner, C.-H. Chen, M.K. Tsai, X. Wu

Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): D. Liang, C.A. Hsiung, M.K. Tsai, C.K. Tsao, X. Wu, S.M. Lippman

Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): C.P. Wen, F. Zhang, H.D. Skinner, W.-H. Chow, Y. Ye, X. Pu, M.A.T. Hildebrandt, M. Huang, M.K. Tsai, X. Wu, S.M. Lippman

Writing, review, and/or revision of the manuscript: C.P. Wen, F. Zhang, D. Liang, J. Gu, H.D. Skinner, W.-H. Chow, Y. Ye, X. Pu, M.A.T. Hildebrandt, C.A. Hsiung, X. Wu, C. Wen, S.M. Lippman

Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): C.A. Hsiung, M.K. Tsai, C.K. Tsao, X. Wu
Study supervision: C.P. Wen, X. Wu

Grant Support

This work was supported by the National Cancer Institute (P50 CA070907 Project 2 to X. Wu), MD Anderson Research Trust and MD Anderson institutional support for the Center for Translational and Public Health Genomics (to

X. Wu), and Taiwan Department of Health Clinical Trial and Research Center of Excellence (DOH102-TD-B-111-004 to C.P. Wen).

The costs of publication of this article were defrayed in part by the payment of page charges. This article must therefore be hereby marked *advertisement* in accordance with 18 U.S.C. Section 1734 solely to indicate this fact.

Received March 27, 2014; revised October 1, 2014; accepted October 9, 2014; published OnlineFirst October 21, 2014.

References

- Wender R, Fontham ET, Barrera E Jr, Colditz GA, Church TR, Ettinger DS, et al. American Cancer Society lung cancer screening guidelines. *CA Cancer J Clin* 2013;63: 107–17.
- Aberle DR, Adams AM, Berg CD, Black WC, Clapp JD, Fagerstrom RM, et al. Reduced lung-cancer mortality with low-dose computed tomographic screening. *N Engl J Med* 2011;365: 395–409.
- Bach PB, Mirkin JN, Oliver TK, Azzoli CG, Berry DA, Brawley OW, et al. Benefits and harms of CT screening for lung cancer: a systematic review. *JAMA* 2012;307: 2418–29.
- Wood DE, Eapen GA, Ettinger DS, Hou L, Jackman D, Kazerooni E, et al. Lung cancer screening. *J Natl Compr Cancer Netw* 2012;10: 240–65.
- Jaklitsch MT, Jacobson FL, Austin JH, Field JK, Jett JR, Keshavjee S, et al. The American Association for Thoracic Surgery guidelines for lung cancer screening using low-dose computed tomography scans for lung cancer survivors and other high-risk groups. *J Thorac Cardiovasc Surg* 2012;144: 33–8.
- Kovalchik SA, Tammemagi M, Berg CD, Caporaso NE, Riley TL, Korch M, et al. Targeting of low-dose CT screening according to the risk of lung-cancer death. *N Engl J Med* 2013;369: 245–54.
- 2020 visions. *Nature* 2010;463: 26–32.
- Sreekumar A, Poisson LM, Rajendiran TM, Khan AP, Cao Q, Yu J, et al. Metabolomic profiles delineate potential role for sarcosine in prostate cancer progression. *Nature* 2009;457: 910–4.
- Gu H, Pan Z, Xi B, Asiago V, Musselman B, Raftery D. Principal component directed partial least squares analysis for combining nuclear magnetic resonance and mass spectrometry data in metabolomics: application to the detection of breast cancer. *Anal Chim Acta* 2011;686: 57–63.
- Ritchie SA, Ahiaonu PW, Jayasinghe D, Heath D, Liu J, Lu Y, et al. Reduced levels of hydroxylated, polyunsaturated ultra long-chain fatty acids in the serum of colorectal cancer patients: implications for early screening and detection. *BMC Med* 2010;8: 13.
- Zhang J, Liu L, Wei S, Nagana Gowda GA, Hammoud Z, Kesler KA, et al. Metabolomics study of esophageal adenocarcinoma. *J Thorac Cardiovasc Surg* 2011;141: 469–75, 75 e1–4.
- Ikeda A, Nishiumi S, Shinohara M, Yoshie T, Hatano N, Okuno T, et al. Serum metabolomics as a novel diagnostic approach for gastrointestinal cancer. *Biomed Chromatogr* 2012;26: 548–58.
- Xue R, Lin Z, Deng C, Dong L, Liu T, Wang J, et al. A serum metabolomic investigation on hepatocellular carcinoma patients by chemical derivatization followed by gas chromatography/mass spectrometry. *Rapid Commun Mass Spectrom* 2008;22: 3061–8.
- Gao H, Dong B, Liu X, Xuan H, Huang Y, Lin D. Metabonomic profiling of renal cell carcinoma: high-resolution proton nuclear magnetic resonance spectroscopy of human serum with multivariate data analysis. *Anal Chim Acta* 2008;624: 269–77.
- Zhou J, Xu B, Huang J, Jia X, Xue J, Shi X, et al. 1H NMR-based metabolomic and pattern recognition analysis for detection of oral squamous cell carcinoma. *Clin Chim Acta* 2009;401: 8–13.
- Bathe OF, Shaykhtudinov R, Kopciuk K, Weljie AM, McKay A, Sutherland FR, et al. Feasibility of identifying pancreatic cancer based on serum metabolomics. *Cancer Epidemiol Biomarkers Prev* 2011;20: 140–7.
- Hori S, Nishiumi S, Kobayashi K, Shinohara M, Hatakeyama Y, Kotani Y, et al. A metabolomic approach to lung cancer. *Lung Cancer* 2011;74: 284–92.
- Jordan KW, Adkins CB, Su L, Halpern EF, Mark EJ, Christiani DC, et al. Comparison of squamous cell carcinoma and adenocarcinoma of the lung by metabolomic analysis of tissue-serum pairs. *Lung Cancer* 2010;68: 44–50.
- Maeda J, Higashiyama M, Imaizumi A, Nakayama T, Yamamoto H, Daimon T, et al. Possibility of multivariate function composed of plasma amino acid profiles as a novel screening index for non-small cell lung cancer: a case control study. *BMC Cancer* 2010;10: 690.
- Spitz MR, Hong WK, Amos CI, Wu X, Schabath MB, Dong Q, et al. A risk model for prediction of lung cancer. *J Natl Cancer Inst* 2007;99: 715–26.
- Wen CP, Cheng TY, Tsai MK, Chang YC, Chan HT, Tsai SP, et al. All-cause mortality attributable to chronic kidney disease: a prospective cohort study based on 462 293 adults in Taiwan. *Lancet* 2008;371: 2173–82.
- Wen CP, Wai JP, Tsai MK, Yang YC, Cheng TY, Lee MC, et al. Minimum amount of physical activity for reduced mortality and extended life expectancy: a prospective cohort study. *Lancet* 2011;378: 1244–53.
- Lawton KA, Berger A, Mitchell M, Milgram KE, Evans AM, Guo L, et al. Analysis of the adult human plasma metabolome. *Pharmacogenomics* 2008;9: 383–97.
- Ryter SW, Morse D, Choi AM. Carbon monoxide and bilirubin: potential therapies for pulmonary/vascular injury and disease. *Am J Respir Cell Mol Biol* 2007;36: 175–82.
- Lin JP, O'Donnell CJ, Schwaiger JP, Cupples LA, Lingenhel A, Hunt SC, et al. Association between the UGT1A1*28 allele, bilirubin levels, and coronary heart disease in the Framingham Heart Study. *Circulation* 2006;114: 1476–81.
- Novotny L, Vitek L. Inverse relationship between serum bilirubin and atherosclerosis in men: a meta-analysis of published studies. *Exp Biol Med* 2003;228: 568–71.
- Horsfall LJ, Rait G, Walters K, Swallow DM, Pereira SP, Nazareth I, et al. Serum bilirubin and risk of respiratory disease and death. *JAMA* 2011;305: 691–7.
- O'Malley SS, Wu R, Mayne ST, Jatlow PI. Smoking cessation is followed by increases in serum bilirubin, an endogenous antioxidant associated with lower risk of lung cancer and cardiovascular disease. *Nicotine Tob Res* 2014;16: 1145–9.
- Frost-Pineda K, Liang Q, Liu J, Rimmer L, Jin Y, Feng S, et al. Biomarkers of potential harm among adult smokers and nonsmokers in the total exposure study. *Nicotine Tob Res* 2011;13: 182–93.
- How Tobacco Smoke Causes Disease. The biology and behavioral basis for smoking-attributable disease. A Report of the Surgeon General. Atlanta (GA); 2010.

Clinical Cancer Research

The Ability of Bilirubin in Identifying Smokers with Higher Risk of Lung Cancer: A Large Cohort Study in Conjunction with Global Metabolomic Profiling

Chi-Pang Wen, Fanmao Zhang, Dong Liang, et al.

Clin Cancer Res Published OnlineFirst October 21, 2014.

Updated version	Access the most recent version of this article at: doi: 10.1158/1078-0432.CCR-14-0748
Supplementary Material	Access the most recent supplemental material at: http://clincancerres.aacrjournals.org/content/suppl/2014/10/22/1078-0432.CCR-14-0748.DC1

E-mail alerts	Sign up to receive free email-alerts related to this article or journal.
Reprints and Subscriptions	To order reprints of this article or to subscribe to the journal, contact the AACR Publications Department at pubs@aacr.org .
Permissions	To request permission to re-use all or part of this article, use this link http://clincancerres.aacrjournals.org/content/early/2014/12/10/1078-0432.CCR-14-0748 . Click on "Request Permissions" which will take you to the Copyright Clearance Center's (CCC) Rightslink site.