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 Zhang\(^3\), Dong Liang\(^4\), Christopher Wen\(^5\), Jian Gu\(^3\), Heath Skinner\(^6\), Wong-Ho Chow\(^3\), Yuanqing Ye\(^5\), Xia Pu\(^5\), Michelle A.T. Hildebrandt\(^3\), Maosheng Huang\(^5\), Chien-Hua Chen\(^7\), Chao Agnes Hsiung\(^1\), Min Kuang Tsai\(^1\), Chwen Keng Tsao\(^8\), Scott M. Lippman\(^9\), and Xifeng Wu\(^3\)

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 association 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 benefited 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
differenced 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 ~8°C until assays were performed. For the series of validation studies, standard powders for two metabolites, that is, L-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 amounts 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 χ² 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.

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

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Men (N = 202,902), N (%)</th>
<th>Women (N = 222,758), N (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total</td>
<td>202,902</td>
<td>222,758</td>
</tr>
<tr>
<td>Total bilirubin level (mg/dL)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;0.75</td>
<td>67,841 (33.4)</td>
<td>75,189 (33.8)</td>
</tr>
<tr>
<td>0.75–1</td>
<td>65,540 (32.3)</td>
<td>72,207 (32.4)</td>
</tr>
<tr>
<td>&gt;1</td>
<td>69,521 (34.3)</td>
<td>72,356 (33.8)</td>
</tr>
<tr>
<td>Age, (years, mean (SD))</td>
<td></td>
<td></td>
</tr>
<tr>
<td>20-39</td>
<td>112,584 (55.5)</td>
<td>119,946 (53.9)</td>
</tr>
<tr>
<td>40-59</td>
<td>63,447 (31.3)</td>
<td>76,087 (34.2)</td>
</tr>
<tr>
<td>BMI (kg/m2), mean (SD)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;25</td>
<td>134,591 (66.4)</td>
<td>176,567 (79.3)</td>
</tr>
<tr>
<td>25-29</td>
<td>59,734 (29.5)</td>
<td>84,454 (37.3)</td>
</tr>
<tr>
<td>30-</td>
<td>8,516 (4.2)</td>
<td>7,689 (3.5)</td>
</tr>
<tr>
<td>Educational levels</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Middle school or lower</td>
<td>40,499 (20.6)</td>
<td>70,385 (32.6)</td>
</tr>
<tr>
<td>High school</td>
<td>45,601 (23.2)</td>
<td>54,124 (25.1)</td>
</tr>
<tr>
<td>Junior college</td>
<td>45,367 (23.1)</td>
<td>42,941 (19.9)</td>
</tr>
<tr>
<td>College or higher</td>
<td>64,987 (33.1)</td>
<td>48,400 (22.4)</td>
</tr>
<tr>
<td>Smoking status</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Non-smoker</td>
<td>92,864 (47.9)</td>
<td>128,232 (57.3)</td>
</tr>
<tr>
<td>Smoker</td>
<td>101,092 (52.1)</td>
<td>124,518 (42.7)</td>
</tr>
<tr>
<td>&lt;30 pack-years</td>
<td>72,153 (35.2)</td>
<td>97,226 (43.5)</td>
</tr>
<tr>
<td>≥30 pack-years</td>
<td>72,153 (35.2)</td>
<td>97,226 (43.5)</td>
</tr>
<tr>
<td>Lung cancer incidence</td>
<td>809 (0.4)</td>
<td>4,241 (0.8)</td>
</tr>
<tr>
<td>Lung cancer mortality</td>
<td>614 (0.3)</td>
<td>330 (0.2)</td>
</tr>
</tbody>
</table>

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 \( P_{\text{trend}} \) values < 0.01 (Supplementary Table S3). After taking into account multiple comparisons, L-glutamylalanine remained as the only metabolite meeting statistical significance after Bonferroni correction \( P_{\text{trend}} < 1.63 \times 10^{-6} \) (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 L-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 L-glutamylalanine in
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, the association was only present in smokers and there were only 37 lung cancer cases among them. 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 tertile with bilirubin levels <1 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 3. Relationship among smoking quantity, bilirubin levels, and risk for lung cancer mortality in male participants in the prospective cohort study

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Total bilirubin level (mg/dL)</th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>No. of mortality</td>
<td>HR* (95% CI)</td>
<td>No. of mortality</td>
<td>HR* (95% CI)</td>
</tr>
<tr>
<td>Non-smoker</td>
<td>92,864</td>
<td>98</td>
<td>1 (Ref.)</td>
<td>35,175</td>
</tr>
<tr>
<td>Total smokers</td>
<td>101,092</td>
<td>478</td>
<td>3.24 (2.60–4.05)</td>
<td>29,632</td>
</tr>
<tr>
<td>Pack-year</td>
<td>&lt;30 pack-years</td>
<td>72,153</td>
<td>90</td>
<td>1.62 (1.20–2.18)</td>
</tr>
<tr>
<td>10–19</td>
<td>24,146</td>
<td>370</td>
<td>4.78 (3.77–6.05)</td>
<td>6,269</td>
</tr>
<tr>
<td>≥20</td>
<td>28,789</td>
<td>172</td>
<td>5.16 (4.01–6.65)</td>
<td>6,759</td>
</tr>
</tbody>
</table>

*Adjusted for age, educational level, and BMI.

### Table 4. Lung cancer incidence and mortality rates and adjusted HR per tertile of serum total bilirubin level among the male participants in the prospective cohort study by smoking status and smoking intensity

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>No. of lung cancer incidence</th>
<th>Adjusted HR* (95% CI)</th>
<th>Incidence Rate Per 10 000 Person-years (95% CI)</th>
<th>No. of lung cancer mortality</th>
<th>Adjusted HR* (95% CI)</th>
<th>Mortality Rate Per 10 000 Person-years (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Total bilirubin level (mg/dL)</td>
<td>No. of mortality</td>
<td>HR* (95% CI)</td>
<td>No. of mortality</td>
<td>HR* (95% CI)</td>
<td>No. of mortality</td>
</tr>
<tr>
<td>Total</td>
<td>215</td>
<td>270</td>
<td>324</td>
<td>1 (Ref.)</td>
<td>1.24 (1.03–1.51)</td>
<td>1.52 (1.26–1.82)</td>
</tr>
<tr>
<td>Nonsmoker</td>
<td>64</td>
<td>50</td>
<td>42</td>
<td>1 (Ref.)</td>
<td>0.86 (0.59–1.27)</td>
<td>0.84 (0.56–1.26)</td>
</tr>
<tr>
<td>Total smokers</td>
<td>159</td>
<td>202</td>
<td>262</td>
<td>1 (Ref.)</td>
<td>1.29 (0.93–1.76)</td>
<td>1.55 (1.25–1.92)</td>
</tr>
<tr>
<td>&lt;30 pack-years</td>
<td>27</td>
<td>47</td>
<td>49</td>
<td>1 (Ref.)</td>
<td>1.71 (1.04–2.79)</td>
<td>1.77 (1.09–2.89)</td>
</tr>
<tr>
<td>≥30 pack-years</td>
<td>88</td>
<td>154</td>
<td>201</td>
<td>1 (Ref.)</td>
<td>1.50 (0.85–1.53)</td>
<td>1.31 (0.63–1.67)</td>
</tr>
</tbody>
</table>

*Adjusted for age, educational level, and BMI.
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 ($P_{interaction} = 0.001$), 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
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
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

www.aacrjournals.org
Clin Cancer Res; 2015
OF7

Published OnlineFirst October 21, 2014; DOI: 10.1158/1078-0432.CCR-14-0748

Downloaded from clinicancreas.aacrjournals.org on July 19, 2017. © 2014 American Association for Cancer Research.
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 (R55 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


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, contact the AACR Publications Department at permissions@aacr.org.