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*Observational Study***Total cholesterol to high-density lipoprotein ratio and nonalcoholic fatty liver disease in a population with chronic hepatitis B**

Correlation between TC/HDL-C and NAFLD in the CHB population

Yu-Ge Zhou, Ning Tian, Wei-Ning Xie

**Abstract****BACKGROUND**

Nonalcoholic fatty liver disease (NAFLD) is characterized by hypertriglyceridemia, increased low-density lipoprotein cholesterol levels, and reduced high-density lipoprotein cholesterol (HDL-C) particles. Previous studies have shown that the total cholesterol to high-density lipoprotein cholesterol ratio (TC/HDL-C) was superior to other lipid metabolism biomarkers for predicting NAFLD risk, and could be a new indicator of NAFLD. However, the association between TC/HDL-C and NAFLD in patients with hepatitis B virus (HBV) has not yet been determined.

**AIM**

To investigate the association between TC/HDL-C and NAFLD in a population with chronic hepatitis B (CHB).

**METHODS**

In this study, 183 HBV-infected patients were enrolled. All participants underwent blood chemistry examinations and abdominal ultrasound. Univariate and multivariate

logistic regression models, curve fitting analysis, and threshold calculation were used to assess the relationship between TC/HDL-C and NAFLD.

## RESULTS

The overall prevalence of NAFLD was 17.49% ( $n = 32$ ) in the 183 CHB participants. The TC/HDL-C of non-NAFLD and NAFLD patients were  $3.83 \pm 0.75$  and  $4.44 \pm 0.77$ , respectively ( $P < 0.01$ ). Logistic regression analysis showed that TC/HDL-C was not associated with NAFLD after adjusting for other pertinent clinical variables. However, at an optimal cutoff point of 4.9, a non-linear correlation between TC/HDL-C and NAFLD was detected. The effect size of the left and right sides of the inflection point were 5.4 (95%CI: 2.3-12.6,  $P < 0.01$ ) and 0.5 (95%CI: 0.1-2.2,  $P = 0.39$ ), respectively. On the left side of the inflection point, TC/HDL-C was positively associated with NAFLD. However, no significant association was observed on the right side of the inflection point.

## CONCLUSION

This study demonstrated a non-linear correlation between TC/HDL-C and NAFLD in a population with CHB. TC/HDL-C was positively associated with NAFLD when TC/HDL-C was less than 4.9 but not when TC/HDL-C was more than 4.9.

**Key Words:** Cholesterol; Lipoprotein cholesterol ratio; Nonalcoholic fatty liver disease; Chronic hepatitis B population; Correlation

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**Core Tip:** Nonalcoholic fatty liver disease (NAFLD) and chronic hepatitis B (CHB) are both common chronic liver diseases. In this observational cross-sectional study, we

explored the association between NAFLD and a lipid metabolism biomarker [the total cholesterol to high-density lipoprotein cholesterol ratio, total cholesterol to high-density lipoprotein cholesterol ratio (TC/HDL-C)] in a population with CHB. Our findings showed a non-linear correlation between TC/HDL-C and NAFLD. TC/HDL-C was positively associated with NAFLD when TC/HDL-C was less than 4.9 but not when TC/HDL-C was more than 4.9.

## **INTRODUCTION**

Chronic hepatitis B (CHB) is a common disease threatening public health and a leading cause of multitudinous liver-related morbidity and mortality<sup>[1]</sup>. In 2016, about 257 million people were affected by hepatitis B virus (HBV) infection worldwide, with an estimated prevalence of 3.5%<sup>[2]</sup>. Over the past decades, with the implementation of nucleoside analogs (NAs) and hepatitis B vaccine, the risk of liver cirrhosis complications and hepatocellular carcinoma (HCC) have been substantially reduced in CHB patients<sup>[3]</sup>. However, since approximately 25% of the CHB population with non-alcoholic fatty liver disease (NAFLD) have hepatic steatosis, the effects of CHB on hepatosteatosis have recently been garnering attention<sup>[4]</sup>.

NAFLD is a common chronic hepatic disease worldwide that is closely associated with cardiovascular disease, metabolic disorders and end-stage liver diseases such as cirrhosis and HCC<sup>[5]</sup>. The prevalence of NAFLD has been increasing and in the past few years it has reached alarming proportions (29.1%) in China due to changes in lifestyle habits and rapid socio-economic growth<sup>[6]</sup>; thereby, urging increasing awareness to recognize NAFLD as a chronic liver disease.

Due to the growing prevalence of NAFLD, the coexistence of HBV infection and NAFLD is commonly observed around the world. However, a clear association between these two diseases remains questionable. Previous studies indicated that NAFLD could be inversely associated with the levels of HBV seromarkers<sup>[7,8]</sup>, but interestingly, there is substantial evidence indicating an association between HBV infection and reduced incidence of hyperlipidemia or NAFLD<sup>[9,10]</sup>.

Alterations in lipid metabolism are central drivers of disease progression, for instance, the progression of hepatic steatosis to non-alcoholic steatohepatitis (NASH) and hepatic fibrosis<sup>[11]</sup>. Therefore, deciphering the lipid metabolism characteristic of NAFLD is the crucial for disease treatment and prevention. NAFLD is characterized by hypertriglyceridemia, increased low-density lipoprotein cholesterol (LDL-C) levels and reduced high-density lipoprotein cholesterol (HDL-C) particles<sup>[12,13]</sup>. A recent study showed that the total cholesterol to high-density lipoprotein cholesterol ratio (TC/HDL-C) was better at predicting NAFLD risk than other markers such as total cholesterol (TC), HDL-C and the ratio of Apolipoprotein B to Apolipoprotein A1, and might be a new indicator of NAFLD<sup>[14]</sup>. However, the association between TC/HDL-C and NAFLD in an HBV-infected population has not yet been investigated. Therefore, the objective of this study was to assess the correlation between TC/HDL-C and NAFLD in a population with CHB.

## **MATERIALS AND METHODS**

### ***Study population and criteria***

This was a retrospective, observational study comprising of HBV-infected patients who were treated at the Integrated Traditional Chinese and Western Medicine Hospital of Foshan (Guangdong, China) from January 2019 to December 2020. The study flow chart is illustrated in Figure 1. Chronic HBV infection was defined by hepatitis B surface antigen (HBsAg) positive for more than 6 mo<sup>[1]</sup>. All participants underwent abdominal ultrasonography for NAFLD, and blood tests for assessing lipid metabolism and hepatic and renal function. Each participant completed a detailed questionnaire concerning information on their gender, age, alcohol consumption history, disease history and medication history. Patients were excluded if they had (1) a daily alcohol intake  $\geq 30$ g (for men) or 20 g (for women)<sup>[15]</sup>; (2) history of cancer; (3) history of chronic renal insufficiency; (4) history of hepatobiliary surgery, and; (5) missing data on the key clinical variables required for study analysis.

### ***Laboratory investigations***

Blood samples were collected from all patients after an overnight fasting of at least 8 h. Peripheral venous blood was drawn from their cubital vein. Blood test parameters, including aspartate aminotransferase (AST), alanine aminotransferase (ALT),  $\gamma$ -glutamyl transpeptidase ( $\gamma$ -GT), creatinine (CRE), uric acid (UA), triglyceride (TG), TC, HDL-C, LDL-C, apolipoprotein A1 (ApoA1) and apolipoprotein B (ApoB), were measured using an Olympus AU-640 autoanalyzer (Olympus, Japan). Platelets (PLT) were measured using the Sysmex 2100 whole blood cell analyzer (Sysmex, Japan). Hepatitis B serum examinations included the detection of HBV-DNA level, HBsAg and hepatitis B e antigen (HBeAg) using polymerase chain reaction. HBV-DNA(+) was defined as a level of serum hepatitis B virus DNA over 100 IU/mL. TC/HDL-C was defined as TC divided by HDL-C. The patient's height and weight were measured while wearing light clothing without shoes. Body mass index (BMI) was calculated by dividing a person's weight in kilograms by the square of their height in meters. Abdominal ultrasound was used to detect the presence of NAFLD. NAs therapy was defined as the use of oral NAs antiviral drugs for more than 3 mo. All the above data were obtained from the Clinical Laboratory of the Guangdong Provincial Hospital of Integrated Traditional Chinese and Western Medicine.

### ***Ethics and consent***

The research project was submitted to and approved by the Ethics Committee of Guangdong Provincial Hospital of Integrated Traditional Chinese and Western Medicine (Approval number: 2018-1254). No informed consent was required because this was a retrospective observational study.

### ***Definition of NAFLD based on abdominal ultrasonography***

Ultrasonography was the most commonly used examination for fatty liver screening due to its noninvasiveness, low cost, and easily operability<sup>[16]</sup>. Abdominal ultrasonography was performed on all enrollees by two trained ultrasound physicians

using the ACUSON X150 ultrasound system (Siemens, Japan). The presence of at least two of the following criteria was required for considering fatty liver: (1) More than 5% of hepatocytes had excessive hepatic fat accumulation and steatosis<sup>[17]</sup>; (2) Diffuse echo enhancement of the liver relative to the kidney; (3) Occurrence of ultrasonic beam attenuation; and (4) Poor visualization of intrahepatic structures. After excluding alcohol abuse and other hepatic diseases, NAFLD was then formally diagnosed by abdominal ultrasonography<sup>[18,19]</sup>.

### *Statistical analysis*

All statistical analyses were conducted using SPSS (version 23.0, IBM, United States) and EmpowerStats (<http://www.empowerstats.com>, X&Y solutions, Inc. Boston MA) software. Normal distribution continuous variables are expressed as mean  $\pm$  standard deviation, and *t*-test was used for group comparison. Non-normal distribution variables are described as median [interquartile range (IQR)], and the Mann-Whitney *U* test was used to compare the groups. Categorical variables were presented as their corresponding number and percentage (*n*, %) and compared using the chi-squared test. All enrollees were stratified into two groups based on the presence or absence of NAFLD on ultrasonography. Then, the demographic characteristics of the study participants of the two groups were assessed. Univariate analyses of all variables were conducted using a binary logistic regression analysis model. Based on their TC/HDL-C, the patients were also divided into three groups according to TC/HDL-C tertiles: TC/HDL-C  $\leq 3.5$ ,  $3.5 < \text{TC/HDL-C} \leq 5$ , and TC/HDL-C  $> 5$ . Multivariable models were constructed as follows: Model 1 was not adjusted for other pertinent clinical variables; Model 2 was adjusted for gender and age; Model 3 was adjusted for gender, age, BMI, AST, ALT,  $\gamma$ -GT, PLT, HBsAg, CRE, UA, TG, TC, HDL-C, LDL-C, ApoA1, ApoB, HBV-DNA (+), HBeAg(+) and NAs. Lastly, a non-linear relationship between TC/HDL-C and NAFLD was investigated and smooth curve fitting was also used. *P* values (two-tailed) less than 0.05 were considered statistically significant.

## RESULTS

### Demographic characteristics

The demographic characteristics of the study participants with and without NAFLD are shown in Table 1. In the whole study population, the overall prevalence of NAFLD was 17.49% ( $n = 32$ ). A total of 183 patients (70.5% males) were included in this study. Their mean age was  $45.41 \pm 11.59$  years and their average BMI was  $23.14 \pm 2.63$  kg/m<sup>2</sup>. The TC/HDL-C of the non-NAFLD and NAFLD groups were  $3.83 \pm 0.75$  and  $4.44 \pm 0.77$ , respectively ( $P < 0.01$ ). Compared with the non-NAFLD group, patients from the NAFLD group had higher levels of BMI, ALT,  $\gamma$ -GT, PLT, UA, TG, TC, LDL-C, ApoB and TC/HDL-C ( $P < 0.05$ ). Conversely, the age, HBV-DNA (+) levels and usage of NAs were significantly lower in the NAFLD group than in the non-NAFLD group ( $P < 0.01$ ). However, there was no statistically significant difference between the two groups in terms of gender and levels of AST, HBsAg, CRE, HDL-C, ApoA1 and HBeAg (+) ( $P > 0.05$ ).

### Univariate analysis

Binary logistic regression of independent risk factors of NAFLD is shown in Table 2. Univariate analysis indicated that age, BMI, ALT,  $\gamma$ -GT, PLT, UA, TG, LDL-C, ApoB, TC/HDL-C and HBV-DNA (+) were significantly positively correlated with NAFLD ( $P < 0.05$ ), whereas age and NAs were negatively correlated with NAFLD. However, no significant association between NAFLD and gender, AST, HBsAg, CRE, TC, HDL-C, ApoA1 and HBeAg(+) were observed ( $P > 0.05$ ).

### Association between TC/HDL-C and NAFLD using logistic regression model

Logistic regression model was used to evaluate the association between TC/HDL-C and NAFLD. The unadjusted and adjusted models are shown in Table 3. In model 1, TC/HDL-C was positively correlated with NAFLD (OR = 0.94, 95%CI: 1.55-4.19,  $P < 0.01$ ). In model 2 (adjusted for gender and age), the relationship between TC/HDL-C and NAFLD were significant (OR = 0.96, 95%CI: 1.52-4.51,  $P < 0.01$ ). However, this



association was not detected in model 3 (OR = -2.27, 95%CI: 0.0001-79.91,  $P = 0.50$ ). The same trend was observed for TC/HDL-C from 3.5 to 5 in model 1 and model 2 ( $P < 0.05$ ).

#### *Analysis of non-linear relationships between TC/HDL-C and NAFLD*

Since TC/HDL-C was a continuous variable, we analyzed the non-linear relationship with NAFLD. After adjusting for all variables and conducting smooth curve fitting, we found that the relationship between TC/HDL-C and NAFLD was non-linear (Figure 2). The inflection point was calculated as 4.9 by a piecewise linear regression model. On the left side of the inflection point, TC/HDL-C was found to be positively correlated with NAFLD ( $\beta = 5.4$ , 95%CI: 2.3-12.6,  $P < 0.01$ ). However, no significant correlation was observed between TC/HDL-C and NAFLD on the right side of the inflection point ( $\beta = 0.5$ , 95%CI: 0.1-2.2,  $P = 0.39$ ) (Table 4).

## **DISCUSSION**

In epidemiologic studies, the highest incidences of NAFLD were reported in the Middle East (32%) and South America (31%), followed by Asia (27%) and the United States of America (24%)<sup>[20]</sup>. It was shown that NAFLD affected 15% to 20% of the Chinese population and that this was increasing at alarming rates<sup>[21]</sup>. However, in a study involving 810 northern Japanese children, the prevalence of fatty liver was observed to be only 2.6% based on ultrasonographic criteria<sup>[22]</sup>, but the prevalence of NAFLD was increased to 77% among obese children<sup>[23]</sup>. In our previous study, we found that the incidence rate of NAFLD in the general population was 35.92%<sup>[24]</sup>. The above studies only focused on the prevalence of NAFLD in the general population. In this present study, the overall prevalence of NAFLD in the investigated CHB population was 17.49%. Consistent with our findings, previous studies indicated that the prevalence of NAFLD with and without HBsAg positivity was 14.3% and 28.6%, respectively ( $P < 0.01$ )<sup>[25]</sup>. A prior study observed a low incidence of NAFLD in their investigated CHB

population and hypothesized that such could be mainly because HBV infection influences the secretion of a variety of adipokines and alterations in lipid profiles<sup>[26]</sup>.

Substantial evidence indicated an association between HBV infection and reduced incidence of hyperlipidemia or NAFLD risk<sup>[9,10]</sup>. In a large cross-sectional study, the researchers observed that HBsAg-positive subjects had a significantly lower risk of NAFLD (OR = 0.42)<sup>[25]</sup>. Additionally, a large prospective cohort study found that TC levels decreased over time among CHB individuals<sup>[27]</sup>. Adiponectin may be central to this observed association. Adipokine may attenuate hepatic steatosis and the degree of its decline was shown to correlate with the severity of NAFLD<sup>[27]</sup>. Moreover, adiponectin levels were also shown to be positively correlated with HBV-DNA viral load in CHB patients<sup>[28, 29]</sup>.

However, the cross-talk between CHB and NAFLD remained controversial. There are studies indicating that NAFLD was inversely associated with the levels of HBV seromarkers<sup>[7,8]</sup>. In this present study, the proportion of HBV-DNA positivity in the NAFLD group ( $n = 15$ , 46.9%) was significantly lower than the non-NAFLD group ( $n = 116$ , 76.8%). Further, a large cohort study demonstrated that HBsAg clearance was significantly higher in CHB patients with hepatic steatosis than in those without<sup>[30]</sup> and these results were in agreement with animal experiments<sup>[31,32]</sup>.

In regards to treatment, long-term oral use of NAs drugs such as entecavir and tenofovir were the main anti-HBV treatment as they are simple and safe to use, which is recognized all over the world<sup>[1]</sup>. Thus, oral NAs therapy alone was the first option for patients with CHB. However, CHB patients with NAFLD needed additional treatment besides antiviral drugs. Lifestyle intervention was a basic method for losing weight. For severe cases, pharmacological treatment was required to regulate the patients' lipid metabolism disorders<sup>[33]</sup>.

Metabolic alterations in NAFLD may directly or indirectly affect the HBV-DNA levels of CHB patients<sup>[34]</sup>. Due to the common immune pathways of NAFLD and CHB, NAFLD-related metabolic stress may activate the suppressed innate immunity to

restores the production of antiviral substances, which ultimately accelerates the clearance of HBV-DNA and HBsAg<sup>[35,36]</sup>.

Metabolic syndrome is a highly prevalent concern in patients with NAFLD<sup>[18,37]</sup>. The typical characteristics of NAFLD are abnormal lipid accumulation in hepatocytes, hypertriglyceridemia, increased LDL-C levels and reduced HDL-C particles. Metabolic perturbations promote liver injury and inflammation, which can lead to increased risk for hepatic fibrosis<sup>[38]</sup>. A cohort study of Chinese people with normal lipid metabolism indicated that a low-density lipoprotein to high-density lipoprotein (LDL/HDL) ratio was superior to other lipoproteins in identifying people at risk of NAFLD<sup>[39]</sup>. Studies from the Framingham Cardiovascular Institute also showed that a ratio of TC/HDL-C greater than 4 was a major risk factor for cardiovascular thrombosis<sup>[40]</sup>. In this present study, our results showed that TG, TC, LDL-C, ApoB, TC/HDL-C had a significant increment in CHB patients combined with NAFLD. Concordant with the results of previous studies, we observed that although the levels of HDL-C and ApoA1 were decreased, no significant statistical difference was observed<sup>[12,41]</sup>.

In our study, TC/HDL-C was a positive risk factor for NAFLD ( $P < 0.01$ ) in univariate analysis. Previous studies suggested that there was a linear relationship between TC/HDL-C and NAFLD in the general population<sup>[14]</sup>. However, in this study, curve fitting analysis **model showed that the association between TC/HDL-C and NAFLD was non-linear in the CHB population for an inflection point of 4.9. Thus, we speculated that TC/HDL-C was positively associated with NAFLD when the ratio of TC/HDL-C was less than 4.9 in the CHB population.**

There were some limitations observed in this study. First, the investigated population was relatively small and therefore, large-scale studies are needed to validate our findings. Second, the assessment of NAFLD was based on hepatic ultrasonography rather than liver biopsy, which was the traditional gold standard for the assessment of NAFLD<sup>[42]</sup>. Patients could be reluctant to undergo liver biopsy because of its high cost, invasiveness and risk of complications<sup>[43]</sup>. Furthermore, fibrosis indices such as hyaluronic acid, laminin, procollagen III peptide, collagen type IV and transient

elastography were not included in the analyses due to missing data on fibrosis indices and could have been conducive to evaluating the relationship between NAFLD and different stages of CHB. In future studies, we will assess the relationship between fibrosis indices and TC/HDL-C. Lastly, this cross-sectional study only explored the relationship between the TC/HDL-C and NAFLD and was unable to reveal the causal and effect relationship between them.

## **CONCLUSION**

In conclusion, the study demonstrated that the relationship between TC/HDL-C and NAFLD was non-linear in the CHB population. TC/HDL-C was positively correlated with NAFLD when TC/HDL-C was less than 4.9 but no such trend could be observed when the ratio of TC/HDL-C was more than 4.9.

## **ARTICLE HIGHLIGHTS**

### ***Research background***

Due to the growing prevalence of non-alcoholic fatty liver disease (NAFLD), the coexistence of hepatitis B virus (HBV) infection and NAFLD is commonly observed around the world. However, the cross-talk between these two diseases remained questionable.

### ***Research motivation***

Previous studies showed that the <sup>3</sup> total cholesterol to high-density lipoprotein cholesterol ratio (TC/HDL-C) was a better predictor of NAFLD than other lipid metabolism biomarkers and might be a new indicator of NAFLD. However, the association between TC/HDL-C and NAFLD in an HBV-infected population has not been previously investigated.

### ***Research objectives***

To investigate the association between TC/HDL-C and NAFLD in a CHB population.

### *Research methods*

Univariate and multivariate logistic regression models, curve fitting analysis and threshold calculations were used to assess the relationship between TC/HDL-C and NAFLD.

### *Research results*

A non-linear association was detected between TC/HDL-C and NAFLD in the CHB population at an inflection point of 4.9. The effect size on the left and right sides of inflection point were 5.4 (95%CI: 2.3-12.6,  $P < 0.01$ ) and 0.5 (95%CI: 0.1-2.2,  $P = 0.39$ ), respectively.

### *Research conclusions*

In the CHB population, the relationship between TC/HDL-C and NAFLD was non-linear. TC/HDL-C was positively correlated with NAFLD when TC/HDL-C was less than 4.9.

### *Research perspectives*

Further large-scale cohort studies are needed to validate whether TC/HDL-C is indeed a better predictor of NAFLD than other lipid metabolism biomarkers in the CHB population.

### **ACKNOWLEDGEMENTS**

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SIMILARITY INDEX

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