Type 2 diabetes mellitus increases the risk of hepatocellular carcinoma in subjects with chronic hepatitis B virus infection: a meta-analysis and systematic review

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Background: Type 2 diabetes mellitus has been proved to be a risk factor of hepatocellular carcinoma, but how diabetes affects incidence of hepatocellular carcinoma among patients with chronic hepatitis B virus infection remains controversial.

Methods: A comprehensive search of Medline and Embase was performed. Incidence of hepatocellular carcinoma in chronic hepatitis B patients was the primary outcome. Pooled HRs and 95% CIs were calculated to assess the correlation between diabetes and incidence of hepatocellular carcinoma.

Results: Five cohort studies and two case–control studies were identified, with a total of 21,842 chronic hepatitis B patients. The diabetes mellitus cohort was found to have increased incidence of hepatocellular carcinoma (pooled HR 1.77, 95% CI 1.28–2.47; fixed effect) and worse overall mortality (pooled RR 1.93, 95% CI 1.64–2.27; fixed effect) in comparison with those without diabetes. In case–control studies, hepatocellular carcinoma cases were found to have an insignificantly elevated diabetes mellitus rate in comparison with the control group.

Conclusion: Type 2 diabetes mellitus is significantly associated with increased risk of hepatocellular carcinoma among patients with chronic hepatitis B virus infection, and aggressive management of diabetes mellitus is strongly suggested.

Keywords: type 2 diabetes mellitus, hepatocellular carcinoma risk, HBV-infected

Introduction

Hepatocellular carcinoma (HCC) is the fifth-most common cancer worldwide, leads to nearly 1 million deaths every year,¹ and is the third-most frequent cause of cancer-related death. The incidence of HCC is particularly high in Asia (over 20 in 100,000 men and over ten in 100,000 women) and in Africa, intermediate in southern Europe, and much lower in most developed countries.² Hepatitis virus infection, mainly hepatitis B virus (HBV) and hepatitis C virus (HCV), has been widely accepted as the major recognized risk factor of HCC globally, accounting for over three-quarters of primary HCC cases.³,⁴ However, when HBV or HCV is not involved, the etiologic factor of HCC varies, of which diabetes mellitus (DM),⁴ heavy alcohol drinking,⁵ smoking,⁶ obesity,⁷ and aflatoxin⁸ are relatively important.

DM, which has been proved to be a risk factor of various kinds of malignancies, is strongly associated with nonalcoholic fatty-liver disease and many other metabolic processes.⁹ Insulin resistance¹⁰ was believed to play an important role in hepatocarcinogenesis in HBV patients with type 2 DM or even prediabetes.¹¹ The association
between DM and HCC risk was indicated to be independent of cirrhosis, though most HCC cases presented with cirrhosis. A recent systematic review demonstrated that concurrent DM is strongly associated with increased HCC risk among chronic HCV patients, but scanty evidence is available about the correlation between DM and HCC in chronic HBV (CHB) patients. The clinical landscape of HCV is currently facing a great change, such that its cure would be universal for patients for whomever has access to effective therapy, which will definitely result in a decrease in HCC developments. Therefore, HBV infection, alcohol consumption, and metabolic disorders, such as DM and obesity, are supposed to be the leading etiologic factors of HCC in the coming future. There are mixed results of the few studies on the association between DM and the risk of HCC in patients with CHB. As such, we performed this meta-analysis and systematic review of the literature to achieve further understanding of the impact of DM on the risk of developing HCC in patients with CHB.

Methods

Literature-search strategy
A comprehensive search of Medline and Embase was performed to retrieve studies published in English (cutoff date February 5, 2018) using the keywords “diabetes” or “diabetes mellitus, type 2” or “DM”, “hepatitis B” or “HBV”, and “hepatocellular carcinoma” or “HCC” or “liver cancer”. We also examined the reference lists of eligible studies to identify additional articles, in order to guarantee a systemic search. Figure 1 depicts the search strategy in detail.

Inclusion and exclusion criteria
Inclusion criteria from the literature search were: studies that focused on the relationship between DM and the risk of HCC: HCC incidence and/or related mortality as outcomes; results of HR/RR/OR and their corresponding 95% CIs for DM and incidence of HCC; and if two or more studies were reported on the same cohort and objectives, either the higher-quality publication or more recent publication was included in the analysis. Studies were excluded if they had not presented data on the relationship between DM and incidence of HCC in patients with HBV infection or specific results were unable to extract. Studies reporting on the effect of DM on the prognosis of HCC or where HCC was not the only outcome (eg, including cholangiocarcinoma) or including patients with type 1 DM were not considered. Reviews, case reports, letters, animal or in vitro studies, conference abstracts, and non-peer-reviewed articles were also excluded from the meta-analysis.

Figure 1 Flow diagram showing the search strategy along with the selection and screening processes for the eligible studies. Abbreviations: DM, diabetes mellitus; HCC, hepatocellular carcinoma; HBV, hepatitis C virus.
Assessment of quality of articles according to the Newcastle–Ottawa Scale, and data extraction were independently performed by YT and SW, as well as the literature search. Quality assessment of both cohort studies and case–control studies included three main characteristics: selection of groups, comparability of cohorts/cases and controls, and outcome/exposure. Studies with seven or more scores were defined as highly qualified.

**Data extraction**

Incidence of HCC was the main outcome assessed in this meta-analysis, while overall mortality or HCC-related mortality was the secondary main outcome. HR, RR, and OR values with their corresponding 95% CIs were extracted from each study. Other information included (but was not limited to) study design, population characteristics, source of cases and controls, ascertainment of type 2 DM, treatment for HBV infection, anti-HCV status, variables adjusted for in the multivariate regression models, and the measure of association between DM and HCC incidence.

**Statistical analysis and exploration of heterogeneity**

We used pooled HRs for primary outcomes to assess the relationship between type 2 DM and incidence of HCC in HBV patients. RRs and weighted mean differences, both with corresponding 95% CIs, were computed for binary data and continuous data, respectively. Funnel plots were visually inspected to identify publication bias. Sensitivity analysis was conducted by removing a single study each time and recalculating pooled results of the remaining studies. The $\chi^2$ test was used to explore potential heterogeneity with $P$ and $P$-values. $P>50\%$ or $P<0.10$ was defined as increased heterogeneity and a random-effect model used, while a fixed model was used when $P>0.10$. Results are presented as $P$-values and 95% CIs, where appropriate, and two-sided $P<0.05$ was considered to indicate statistical significance. Meta-analyses were performed using RevMan software (version 5.3.5; Cochrane Collaboration, Copenhagen). Quality of evidence was evaluated using the software GradePro, comprising four levels: high quality, moderate quality, low quality, and very low quality.

**Results**

The online search initially found 1,682 studies, and 1,624 were excluded after screening of titles and abstracts. A total of 58 full-text articles were reviewed. Finally, two case–control studies $^{17,19}$ and five cohort studies $^{14–16,20,21}$ were identified, of which six were conducted in Asia $^{14–17,19,20}$ and one in New Zealand. $^{21}$ In all cohort studies, patients diagnosed with HCC before the inception point were excluded. The number of eligible CHB patients ranged from 223 to 6,545, with a total of 21,842. Diagnosis of type 2 DM was obtained from patient self-report, abnormal fasting/random glucose, positive oral glucose tolerance test, and DM management (oral hyperglycemic agent or insulin injection). HCC cases were confirmed with the combination of increased AFP and imaging findings (ultrasound, enhanced computed tomography, or angiography) in five studies, $^{14,15,17,19,21}$ and positive histology or cytology was also used to define HCC in the two case–control studies. $^{17,19}$ Two cohort studies $^{16,20}$ in Taiwan defined HCC cases according to the national cancer registry alone (Table 1). Only Hsiang et al. $^{21}$ reported details of dropouts and withdrawals. In their study, seven patients were lost to follow-up or moved overseas, resulting in a loss to or unavailability for follow-up rate of 3%. Other studies, however, did not provide data about details of follow up.

No statistical differences were found between subjects with and without type 2 DM for average age (mean difference 2.81, 95% CI –2.91 to 8.52), male sex (RR 0.99, 95% CI 0.91–1.08), years of follow-up (mean difference –0.35, 95% CI –1.02 to 0.32), or HBV-treatment rate (RR 1.07, 95% CI 0.73–1.55).

**Type 2 DM and risk of HCC in CHB subjects**

Of the seven studies included in this meta-analysis, three cohort studies $^{15,20,21}$ and one case–control study $^{17}$ demonstrated a positive association between type 2 DM and risk of HCC in CHB patients, while other two cohort studies $^{14,16}$ and one case–control study $^{19}$ failed to find a statistical difference between the two groups. Four in seven studies reported correlations between DM and HCC with HRs, $^{14,16,20,21}$ two with ORs $^{17,19}$ and another with RRs, $^{15}$ all with corresponding 95% CIs (Table 2). All studies had different variables adjusted in multivariate regression analysis (Table 3).

The incidence of HCC in DM cohorts varied from 3.29% to 26.0% in cohort studies, while the rate was 2.02%–13.3% in the non-DM group. The four cohort studies reporting on HR $^{14,16,20,21}$ found increased incidence of HCC among patients with DM over those without DM among CHB patients (Figure 2), resulting in a pooled HR of 1.77 (95%CI 1.28–2.47, heterogeneity $P=0$; fixed effect). In case–control studies, $^{17,19}$ an elevated DM rate was indicated in HCC cases in comparison with control groups (12.35% vs 6.53%), but not statistically, with an RR of 2.10 (95% CI 0.84–5.25; random-effect).
<table>
<thead>
<tr>
<th>Study</th>
<th>Period, location</th>
<th>Study design</th>
<th>Group</th>
<th>n</th>
<th>Participants</th>
<th>DM ascertainment</th>
<th>HCC diagnosis</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lai et al</td>
<td>1999–2002, Taiwan</td>
<td>Cohort</td>
<td>DM NDM</td>
<td>63</td>
<td>A total of 54,916 subjects aged ≥30 years from a community-based program; HBV-positive patients (n=6,545)</td>
<td>FBG &gt;126 mg/dL, self-reported</td>
<td>AFP &gt;400 mg/mL, positive imaging finding (enhanced CT, angiography), and histopathological confirmation</td>
</tr>
<tr>
<td>Chen et al</td>
<td>1991–1992, Taiwan</td>
<td>Cohort</td>
<td>DM NDM</td>
<td>62</td>
<td>23,567 residents from a multiple township-based cancer-screening program; seropositive for HBsAg only (n=3,924)</td>
<td>Self-reported</td>
<td>1) National cancer registry; 2) histopathological confirmation; 3) two imaging findings (ultrasonography, angiography, or CT). one imaging diagnosis and AFP &gt;400 ng/mL</td>
</tr>
<tr>
<td>Wang et al</td>
<td>1997–2004, Taiwan</td>
<td>Cohort</td>
<td>DM NDM</td>
<td>47</td>
<td>5,929 residents &gt;35 years old with hepatitis from a population-based screening program; seropositive for HBV only (n=696)</td>
<td>FBG ≥126 mg/dL, nonfasting glucose ≥200 mg/dL, or using hypoglycemic drugs</td>
<td>National cancer registry</td>
</tr>
<tr>
<td>Ko et al</td>
<td>2004–2005, Taiwan</td>
<td>Case–control</td>
<td>DM NDM</td>
<td>421</td>
<td>Hospitalized inpatients first diagnosed with HCC (n=359) and non-HCC controls (n=1,536) randomly selected from a health-checkup program</td>
<td>FBG ≥126 mg/dL, current use of oral hyperglycemic agent or insulin injection at the time of recruitment</td>
<td>1) Histopathological verification; 2) AFP &gt;400 mg/mL and at least one imaging study (ultrasonography, enhanced CT, or angiography)</td>
</tr>
<tr>
<td>Li et al</td>
<td>2004–2008, Chinese mainland</td>
<td>Case–control</td>
<td>DM NDM</td>
<td>421</td>
<td>Patients aged ≥30 years hospitalized for HCC (n=1,105) or CHB (n=5,170) without HBV treatment</td>
<td>FBG ≥7 mmol/L, self-reported</td>
<td>1) Histopathological confirmation; 2) two imaging findings (ultrasonography, enhanced CT, or MRI); 3) AFP &gt;400 mg/mL and one positive image finding</td>
</tr>
<tr>
<td>Fu et al</td>
<td>1997–2009, Taiwan</td>
<td>Cohort</td>
<td>DM NDM</td>
<td>2,099</td>
<td>Chronic HBV patients with (n=2,099) or without DM (n=2,080) randomly selected from the national health-research database (1:1 ratio)</td>
<td>NA</td>
<td>National insurance-program registry (for catastrophic illness patient database)</td>
</tr>
<tr>
<td>Hsiang et al</td>
<td>2000–2012, New Zealand</td>
<td>Cohort</td>
<td>DM NDM</td>
<td>50</td>
<td>223 HBV patients with transient elastography or radiological features of established cirrhosis with (n=50) or without DM (n=173)</td>
<td>FBG ≥7.0 mmol/L, random glucose level ≥11.1 mmol/L, or positive OGTT and/or HbA1c ≥6.5%, and active DM follow-up*</td>
<td>Combination of elevated AFP and positive imaging finding</td>
</tr>
</tbody>
</table>

Note: *Under primary care with diabetes disease-management program or under hospital diabetes-service follow-up.

Abbreviations: CT, computed tomography; HbA1c, glycosylated hemoglobin; HBV, hepatitis B virus; HCC, hepatocellular carcinoma; HBsAg, HBV surface antigen; DM, diabetes mellitus; NDM, non-DM; MRI, magnetic resonance imaging; FBG, fasting blood glucose; OGTT, oral glucose-tolerance test; NA, not applicable.
Increased HCC risk from T2DM + HBV

Among the three studies reporting RRs or ORs, Chen et al\(^\text{15}\) and Ko et al\(^\text{17}\) also found significant associations between type 2 DM and risk of HCC in HBV subjects, with an RR of 2.41 (95% CI 1.17–4.95) and an OR of 4.32 (95% CI 1.92–9.70), respectively, but Li et al\(^\text{19}\) found no significant relationships between DM and risk of HCC when comparing all HCC cases with cross-sectional controls, with an OR of 0.9 (95% CI 0.7–1.2). The publication bias of studies assessing the relation between DM and risk of HCC is shown in Figure 3.

Analysis of sensibility was carried out by excluding one article at a time to guarantee stability of the meta-analysis. As a result, pooled results remained statistically positive when any of the cohort studies was excluded from the meta-analysis. For example, when Wang et al was excluded, the pooled HR of remaining studies remained at 1.80 (95%CI 1.20–2.72). When it came to heterogeneity, I\(^2\) remained at 0 when any study was excluded.

### DM, overall mortality, and HCC-related mortality

Two cohort studies\(^\text{20,21}\) reported overall mortality between two groups, revealing that subjects with type 2 DM suffered

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**Table 2 DM and incidence of HCC risk**

<table>
<thead>
<tr>
<th>Study</th>
<th>Group</th>
<th>n</th>
<th>HCC cases</th>
<th>HCC, %</th>
<th>OR/HR/RR</th>
<th>95% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lai et al(^\text{14})</td>
<td>DM</td>
<td>NA</td>
<td>NA</td>
<td>NA</td>
<td>HR</td>
<td>1.04</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>NA</td>
<td>NA</td>
<td>NA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Chen et al(^\text{15})</td>
<td>DM</td>
<td>62</td>
<td>8</td>
<td>12.9</td>
<td>RR</td>
<td>2.41</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>3,862</td>
<td>179</td>
<td>4.6</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Wang et al(^\text{16})</td>
<td>DM</td>
<td>47</td>
<td>NA</td>
<td>NA</td>
<td>HR</td>
<td>1.3</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>649</td>
<td>NA</td>
<td>NA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Ko et al(^\text{17})</td>
<td>DM</td>
<td>NA</td>
<td>NA</td>
<td>NA</td>
<td>OR</td>
<td>4.32</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>NA</td>
<td>NA</td>
<td>NA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Li et al(^\text{19})</td>
<td>DM</td>
<td>421</td>
<td>93</td>
<td>NA</td>
<td>OR</td>
<td>0.9</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>5,854</td>
<td>1,012</td>
<td>NA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fu et al(^\text{20})</td>
<td>DM</td>
<td>2,099</td>
<td>69</td>
<td>3.29</td>
<td>HR</td>
<td>1.798</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>2,080</td>
<td>42</td>
<td>2.02</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Hsiang et al(^\text{21})</td>
<td>DM</td>
<td>50</td>
<td>13</td>
<td>26</td>
<td>HR</td>
<td>2.36</td>
</tr>
<tr>
<td></td>
<td>NDM</td>
<td>173</td>
<td>23</td>
<td>13.29</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

**Abbreviations:** DM, diabetes mellitus; HCC, hepatocellular carcinoma; NDM, non-DM; NA, not applicable.

**Table 3 Factors adjusted in the multivariate regression analysis**

<table>
<thead>
<tr>
<th>Study</th>
<th>Adjusted confounders</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lai et al(^\text{14})</td>
<td>Age, gender, HCV status, smoking and cumulative consumption of alcohol</td>
</tr>
<tr>
<td>Chen et al(^\text{15})</td>
<td>Age, sex, cigarette smoking, habitual alcohol consumption, and education levels</td>
</tr>
<tr>
<td>Wang et al(^\text{16})</td>
<td>Age, sex, smoking habit, alcohol consumption, BMI, and diabetes status before the study</td>
</tr>
<tr>
<td>Ko et al(^\text{17})</td>
<td>Age, sex, and other viral hepatitis infection</td>
</tr>
<tr>
<td>Li et al(^\text{19})</td>
<td>Age, sex, city of residence, family history of liver cancer, HBeAg status and cirrhosis</td>
</tr>
<tr>
<td>Fu et al(^\text{20})</td>
<td>Age, sex, hyperlipidemia, HBV treatment, statin therapy, cirrhosis, comorbidity index, and obesity</td>
</tr>
<tr>
<td>Hsiang et al(^\text{21})</td>
<td>Age, sex, antiviral therapy, sustained viral suppression, and MELD score</td>
</tr>
</tbody>
</table>

**Abbreviations:** HCV, hepatitis C Virus; BMI, body-mass index, MELD, model for end-stage liver disease.

**Figure 2** Forest plot of meta-analysis results comparing the incidence of HCC between patients with DM and those without DM. **Abbreviations:** HCC, hepatocellular carcinoma; DM, diabetes mellitus.
significantly higher overall mortality in comparison with those without DM (Figure 4), with a pooled RR of 1.93 (95% CI 1.64–2.27, $P=18\%$; fixed effect). Only Hsiang et al\textsuperscript{21} reported on HCC-related mortality in patients with CHB, and patients assigned to the DM group had significantly higher HCC-related mortality than those in the non-DM group (27.9 vs 8.8 per 1,000 patient-years, $P=0.02$). They also demonstrated increased liver-related mortality or orthotopic liver-transplantation rate in the DM group, at 23.4%, compared to the non-DM group, at 9.4% ($P=0.009$).

**Risk of bias and quality evaluation of evidence**

We conducted a quality evaluation on all five cohort studies and two case–control studies included, based on the coding manual for cohort-studies and for case–control studies. All studies scored at least 6, and five studies scored 7 or more. The main outcomes and some other results are summarized in Table 4. The evidence was considered to be of low quality.

**Discussion**

Chronic hepatitis virus infection, including HBV and HCV,\textsuperscript{22} has always been widely acknowledged as a major risk factor of primary HCC. DM, as well as other metabolic abnormalities, has been proved to be associated with quite a few kinds of malignancies,\textsuperscript{23,24} and this association was suggested not to be mediated by body-mass index.\textsuperscript{25} A recent meta-analysis\textsuperscript{26} and a systemic review\textsuperscript{13} demonstrated a strong associations between concurrent DM and risk of HCC among chronic HCV patients. While well studied in anti-HCV-positive
subjects, the potential relationship of type 2 DM and risk of HCC in the HBV-infected population remains unclear. In addition, with effective treatment for HCV infection, HCV-related HCC cases are expected to decrease worldwide. Therefore, HBV infection and metabolic factors, including obesity, DM, and hyperlipidemia, are predicted to account for most of the increase in HCC.

Five cohort studies and two case–control studies were included in the current meta-analysis, with >20,000 individuals enrolled. A recent meta-analysis by Chen et al summarized three studies (two reporting with HRs and the other with RRs), revealing a similar risk of HCC in diabetics without DM among CHB patients. However, in the current meta-analysis, we found a significant association between type 2 DM and increasing incidence of HCC among CHB patients, with a pooled HR of 1.77 (95% CI 1.28–2.47) and no heterogeneity detected ($I^2=0$). Previous meta-analyses have failed to find statistical differences between the two groups, possibly due to the small number of studies included and statistical issues, as not all results were presented with HRs. The biological mechanism for how type 2 DM influences HCC development is not well understood and remains controversial. Most researchers have suggested that type 2 DM contributes to HCC development independently or synergistically with other risk factors, such as HBV or HCV infection and alcohol consumption.

Though relatively uncommon in the setting of HCC, DM is of growing importance, because of its rapidly increasing incidence among adults, as well as nonalcoholic fatty-liver disease, especially in developed countries. It has been reported that type 2 DM and/or obesity had the greatest population-attributable fractions (36.6%) and proportion of cases that could be attributed to specific risk factors: to HCC, higher than alcohol-related disorders (23.5%) or HCV (22.4%). Insulin resistance and hyperinsulinemia are believed to be key factors of HCC oncogenesis in diabetics, mostly through the process of inflammation and cellular injury. While playing an important role in glucose and lipid metabolism, insulin also has pleiotropic effects on regulation of inflammation and cell proliferation. IGF1, the most powerful activator of cellular proliferation, and IRS1 are crucial downstream targets of insulin. HCC cells have been found to overexpress IGF1 and IRS1, revealing that they are of great importance in the process of HCC development. On the other hand, the inflammatory milieu caused by insulin resistance and nonalcoholic steatohepatitis leads to multiple pathways of inflammatory processes, which in return activated oncogenic signaling pathways, such as PI3K–PTEN–Akt and JAK–STAT.9 Insulin resistance also causes hepatic inflammation and fibrosis by the accumulation of fat within hepatocytes, which produces oxidative stress and results in hepatosteatosis. All these activations of pathways are associated with cellular proliferation promotion, increased angiogenesis, and decreased apoptosis, which finally lead to HCC development.

Table 4 Quality evaluation of evidence

<table>
<thead>
<tr>
<th>Studies, n</th>
<th>Design</th>
<th>Patients, n</th>
<th>Effect</th>
<th>95% CI</th>
<th>Quality</th>
<th>Level</th>
</tr>
</thead>
<tbody>
<tr>
<td>Incidence</td>
<td>4</td>
<td>Observational</td>
<td>11,643</td>
<td>1.88</td>
<td>1.28–2.47</td>
<td>ââââ ââââ</td>
</tr>
<tr>
<td>Overall mortality</td>
<td>2</td>
<td>Observational</td>
<td>4,402</td>
<td>1.93</td>
<td>1.64–2.27</td>
<td>ââââ ââââ</td>
</tr>
<tr>
<td>Age</td>
<td>2</td>
<td>Observational</td>
<td>4,402</td>
<td>2.81</td>
<td>–2.91 to 8.92</td>
<td>ââââ ââââ</td>
</tr>
<tr>
<td>Sex</td>
<td>2</td>
<td>Observational</td>
<td>4,402</td>
<td>0.99</td>
<td>0.91–1.08</td>
<td>ââââ ââââ</td>
</tr>
<tr>
<td>HBV treatment</td>
<td>2</td>
<td>Observational</td>
<td>4,402</td>
<td>1.07</td>
<td>0.73–1.55</td>
<td>ââââ ââââ</td>
</tr>
<tr>
<td>Follow-up, years</td>
<td>2</td>
<td>Observational</td>
<td>4,402</td>
<td>–0.35</td>
<td>–102 to 0.32</td>
<td>ââââ ââââ</td>
</tr>
<tr>
<td>% Diabetes mellitus</td>
<td>2</td>
<td>Observational</td>
<td>8,180</td>
<td>2.18</td>
<td>0.82–5.83</td>
<td>ââââ ââââ</td>
</tr>
</tbody>
</table>
In our meta-analysis, five studies were conducted in Taiwan. Although they were conducted using different databases in different periods (Table 1), it is still difficult to confirm whether the cohort pool used in each study was completely different. Besides, methods used to identify HCC and DM were different in the studies included. Therefore, we think this may be a limitation of our study, and our results should be interpreted with caution.

The major limitation involves the variability of the adjustments within the studies. As is well known, HCC is related to quite a few independent risk factors, of which HBV DNA levels, antiviral therapy, and liver cirrhosis are of great importance. However, specific data about these effecting factors were not available in most of the studies included in this meta-analysis. Three studies19–21 clearly adjusted for cirrhosis in the multivariate analysis, two20,21 adjusted for antiviral therapy, and one19 just excluded patients who accepted antiviral therapy, while only one21 adjusted for cirrhosis in the multivariate analysis, two20,21 adjusted for antiviral therapy, and one19 just excluded patients who accepted antiviral therapy, while only one accepted antiviral therapy.

Consequently, the findings of type 2 DM significantly related to HCC risk may shed some light on the prevention of HCC. Better management of DM and correlated metabolic factors is strongly recommended, and HBV patients with DM should be tested more frequently, in order to improve early detection of HCC or other malignancies.

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References


