Keywords: diabetes mellitus; hepatocellular carcinoma; viral hepatitis

Diabetes mellitus and risk of hepatocellular carcinoma: findings from the Singapore Chinese Health Study

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Background: The increasing prevalence of diabetes may contribute to the rising incidence of hepatocellular carcinoma (HCC) in the US and other developed countries where HCC incidence is relatively low. Data from prospective studies on diabetes and risk of HCC in at-risk populations due to high prevalence of viral hepatitis in southeast Asia are sparse.

Methods: The Singapore Chinese Health Study is a prospective cohort of 63 257 middle-aged and older Chinese men and women enrolled in 1993–1998. Besides an in-person interview administered to all participants at baseline, testing of serologic markers of hepatitis B or C infections were performed on a subset of cohort subjects. After a mean follow-up of 14 years, 499 cohort participants developed HCC.

Results: A history of diabetes at baseline was associated with a hazard ratio of 2.14 (95% confidence interval, 1.69–2.71). This statistically significant association was comparable in magnitude between men and women, and remained equally strong across strata of subjects defined by the number of years between their first clinical diagnosis of diabetes and time of enrolment in this cohort. Within a nested case-control set of cohort subjects tested for serological markers of hepatitis B or C infections, the diabetes–HCC association was found to be present mainly among those devoid of any markers.

Conclusion: A history of diabetes at baseline is highly associated with non-viral HCC. Future studies are warranted to elucidate the biological mechanism underpinning the role of diabetes in nonviral-related hepatocarcinogenesis.

Diabetes mellitus is a major cause of morbidity and mortality worldwide, and the influence of this metabolic disease on the overall health of populations is expected to increase as the overall prevalence of diabetes increases in both developed and developing countries. According to data by the International Diabetes Federation that tracks diabetes prevalence worldwide, the expected prevalence of diabetes is expected to increase from 6.6–7.8% by 2030, with the total number of patients living with diabetes expected to rise from 285 million to 438 million over that time span (Federation, 2009). Besides cardiovascular disease (Bjornholt *et al*,

1999), kidney disease (Foundation, 2002) and blindness (Tapp *et al*, 2003), diabetes has been associated with an increased risk of cancer including that of the pancreas (Huxley *et al*, 2005), endometrium (Friberg *et al*, 2007), colon/rectum (Larsson *et al*, 2005), bladder (Larsson *et al*, 2006) and breast (Larsson *et al*, 2007).

The association between diabetes and hepatocellular carcinoma (HCC) has been investigated for several decades in epidemiological studies. We first reported a positive association between history of diabetes and risk of HCC in a non-Asian population in Los Angeles County, California in 1991 (Yu *et al*, 1991). This

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Received 5 October 2012; revised 21 December 2012; accepted 5 January 2013; published online 31 January 2013

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association was confirmed in our later analysis with a larger study sample size (Yuan *et al*, 2004). Similar findings were reported in other populations in the US (Atchison *et al*, 2011; Coughlin *et al*, 2004; El-Serag *et al*, 2004), Japan (Fujino *et al*, 2001; Tazawa *et al*, 2002; Ohata *et al*, 2003; Shibata *et al*, 2003; Inoue *et al*, 2006), Taiwan (Lai *et al*, 2006; Wang *et al*, 2009; Lai *et al*, 2012) and in western Europe (Adami *et al*, 1996; Wideroff *et al*, 1997; Veldt *et al*, 2008; Ogunleye *et al*, 2009). The rising prevalence of obesity and diabetes could be one of the reasons contributing to the increasing incidence of HCC in the United States in the past three decades (El-Serag *et al*, 2003). Given the growing prevalence of obesity and diabetes in developing or recently developed countries where hepatitis B or C infections are endemic (Kelly *et al*, 2008), the establishment of a direct relationship between diabetes and HCC risk in such high-risk populations bears important public health implications.

In the present study, using a population-based prospective cohort, we evaluated the association between diabetes and HCC in Singaporean Chinese, a population with relatively high prevalence of hepatitis B infection and who are at relatively high risk of HCC (Koh *et al*, 2011). The cohort's extensive period of follow-up (average of 14 years per participant) allows us to meaningfully examine the temporality of the diabetes–HCC association. The high prevalence of viral hepatitis in this target population also allows for a meaningful evaluation of the potential modifying role of viral hepatitis on the diabetes–HCC association.

MATERIALS AND METHODS

Study population. The Singapore Chinese Health Study is a population-based prospective cohort established between April 1993 and December 1998 through the recruitment of 63 257 Chinese men ($n = 27\,959$) and women ($n = 35\,298$), who were aged 45–74 years and residing in public housing estates, where 86% of Singapore resided at that time (Hankin *et al*, 2001). All participants belonged to one of the two major Chinese dialect groups in Singapore, the Hokkians or the Cantonese, who originated from two contiguous prefectures in southern China. The study was approved by the Institutional Review Boards of the various sponsoring institutions in the US and Singapore.

Baseline exposure assessment. At recruitment, an in-person interview was conducted in the subject's home by a trained interviewer using a structured questionnaire that covered demographics, lifetime use of tobacco, current physical activity, menstrual/reproductive history (for women), occupational exposure, medical history and family history of cancer. The subjects were asked if they had a history of physician-diagnosed diabetes mellitus (fasting plasma glucose level $\geq 126\text{ mg dl}^{-1}$); positive subjects were then asked for their ages at the time of diagnosis. Using standard protocols, a separate study of 1651 cohort subjects who self-reported a history of physician-diagnosed diabetes mellitus at baseline (Odegaard *et al*, 2008) has validated the accuracy (at 98.8%) of the self-reported diabetes in this cohort.

Between April 1994 and December 1999, blood and single-void urine specimens were collected from a random 3% sample of study enrollees (Koh *et al*, 2003). Starting in January 2000, the biospecimen collection was extended to all surviving cohort members. By April 2005, when all subjects had been contacted, biospecimens were collected from 32 543 participants, representing a ~60% consent rate. Various components (plasma, red blood cell, serum and white blood cell) of blood were separated and have been stored continuously at -80°C .

Case ascertainment. Incident cases of cancers and deaths among cohort members were identified through linkage of the cohort master files with databases of the population-based Singapore

Cancer Registry and Singapore Registry of Births and Deaths. The nationwide cancer registry has been in place since 1968 and has been shown to be comprehensive in its recording of incident cancer cases (Parkin *et al*, 2002). As of December 31, 2010, 47 cohort subjects were lost to follow-up, mainly due to migration out of Singapore, and 499 cohort participants who were free of cancer at baseline had developed primary liver cancer. Among them, 18 (3.7%) were diagnosed with intrahepatic bile carcinoma or sarcomas. For 362 HCC cases diagnosed before December 31, 2006, their diagnoses were confirmed via manual review of pathology reports by a medically trained research staff. Among these HCC cases, 31.5% (114 cases) were diagnosed histologically, 65.0% (235 cases) were diagnosed on the basis of elevation in serum α -fetoprotein in conjunction with clinical and radiologic evidence consistent with HCC, and 3.5% (13 cases) were identified through death certificates.

Hepatitis B and C serology. Beginning in April 1994, a random 3% sample of cohort participants were asked to provide blood or buccal cells, and spot urine samples. Eligibility for this biospecimen subcohort was extended to all surviving cohort participants starting in January 2000. By April 2005, all surviving cohort subjects had been contacted for biospecimen donation. Samples were obtained from 32 535 subjects, representing a consent rate of about 60%.

We conducted a nested case-control study within the biospecimen subcohort for serological biomarkers of hepatitis B and C infection. All incident HCC cases with a baseline blood sample and the diagnosis of HCC before December 31, 2006 were eligible for this nested case-control study. Ninety-two HCC cases met these study criteria (Koh *et al*, 2011). For each case, three control subjects individually matched to the index case by gender, dialect group (Hokkien, Cantonese), age at enrolment (± 2 years), date of baseline interview (± 6 months) were randomly selected from the biospecimen subcohort who were alive and free of cancer on the date of cancer diagnosis of the index case. We assayed serological markers of hepatitis B virus (HBV) and hepatitis C virus (HCV) infections in baseline serum samples of 92 HCC cases and 274 matched control subjects. Briefly, we tested for the presence of HBsAg (Ausria, Abbott Laboratories, North Chicago, IL, USA), and negative samples were further tested for the presence of anti-HBc and anti-HBs (Corab and Ausab, Abbott Laboratories, North Chicago, IL, USA, respectively). All baseline serum samples also were tested for the presence of anti-HCV using the ELISA version 2.0 kit manufactured by Ortho Diagnostic Systems (Ortho, Raritan, NJ, USA), with confirmation of positive samples using RIBA version 2.0 (Chiron, Emeryville, CA, USA).

Statistical analysis. We excluded 1936 individuals with a history of invasive cancer (except non-melanoma skin cancer) at baseline interview. The present analysis included 61 321 cohort subjects. The distributions of demographic and lifestyle factors were compared between diabetics and non-diabetics, and between HCC cases and non-cases. We used the χ^2 test for categorical variables, and the t -test or the analysis of variance method for continuous variables.

For each subject, person-year of follow-up was counted from the date of enrolment to the date of diagnosis of HCC, being lost to follow-up, death, or 31 December 2010, whichever occurred first. Proportional hazards (Cox) regression was used to evaluate the associations between diabetes status and the risk of developing HCC. The strength of the association was measured by the hazard ratio (HR) and its 95% confidence interval (CI) and P -value. All Cox regression models have the following covariates: age at recruitment (years), gender, year of cohort enrolment, dialect group (Hokkien, Cantonese), body mass index (BMI) (kg m^{-2}), level of education (no formal schooling, primary school, secondary school or higher), cigarette smoking status (never, former or current smoker), consumption of alcoholic beverages (non-drinker,

<7 or 7+ drinks per week), frequency of black tea or green tea consumption (none, monthly, weekly or daily) and coffee consumption (number of cups per week) (Johnson *et al.*, 2011).

Using the nested case-control set of cohort subjects with HBV/HCV serology measurements, we first examined the relationships between markers of HBV and HCV infections and HCC risk using conditional logistic regression methods. We then examined the diabetes–HCC association stratified by the subjects' status on viral serology. These latter analyses were conducted using the unconditional logistic regression methods.

We used SAS statistical software version 9.2 (SAS Institute, Cary, NC, USA). All *P*-values quoted are two-sided. *P*-values <0.05 were considered statistically significant.

RESULTS

A total of 5469 (8.9%) subjects reported a history of diabetes at baseline. The mean age of initial diagnosis of diabetes was 51.8 (s.d. 9.4) years and >95% were diagnosed with diabetes after 35 years of age. The diabetics were older, less educated, had higher BMI, consumed less alcohol and drank less coffee, but more tea than non-diabetics. Diabetics who smoked were more likely to quit smoking than their non-diabetic counterparts (Table 1).

After an average of 14.0 (s.d. 3.7) years of follow-up, there were 499 incident HCC cases in this cohort. Men comprised close to three-fourths of the HCC case group (Table 1). The HCC cases were older, slightly higher in BMI, more likely to smoke cigarettes and consumed more alcoholic beverages than cohort subjects free of HCC (Table 1).

A history of diabetes was associated with an approximately two-fold risk of developing HCC (HR 2.64, 95% CI 2.09–3.33) (Table 2). Adjustment for cigarette smoking, alcohol intake and other potential confounders did not materially change the HR for HCC associated with diabetes (HR 2.14, 95% CI 1.69–2.71). The HRs for HCC in men (HR 2.11, 95% CI 1.58–2.81) and in women (HR 2.14, 95% CI 1.41–3.25) were comparable, though the women had only one-third the incidence rate of HCC in men (28.2 vs 98.3 per 100 000 person-years).

Table 2 also shows the diabetes–HCC association across strata of subjects defined by their time interval (years) between first clinical diagnosis of diabetes and enrolment in the cohort. There was no material differences in risk across the three strata, with subjects having the longest history of diabetes being those diagnosed ≥10 years prior to their cohort enrolment, and subjects having the shortest history being those diagnosed within 5 years of cohort enrolment. The ORs of HCC for subjects with different length of diabetes history compared with non-diabetics were

Table 1. Distributions of selected characteristics in subjects with or without a history of diabetes at baseline, and those who developed hepatocellular carcinoma (HCC) or who remained free of HCC (non-HCC), Singapore Chinese Health Study 1993–2010

	Non-diabetics	Diabetics	P-value ^a	Non-HCC	HCC	P-value ^a
Number of subjects	55 852	5469		60 822	499	
Age at interview, mean (s.d.)	56.0 (7.9)	60.0 (7.7)	<0.0001	56.4 (8.0)	60.4 (7.5)	<0.0001
Body mass index (kg m ⁻²), mean(s.d.)	23.0 (3.2)	24.1 (3.3)	<0.0001	23.1 (3.3)	23.9 (3.5)	<0.0001
Gender, %						
Male	44.6	43.1	0.035	44.3	73.3	<0.001
Female	55.4	56.9		55.7	26.7	
Dialect group, %						
Cantonese	46.3	45.6	0.375	46.2	38.5	0.0005
Hokkien	53.8	54.4		53.8	61.5	
Level of education, %						
No formal schooling	26.4	35.0	<0.0001	27.2	28.1	0.052
Primary school	44.4	44.1		44.4	48.3	
Secondary school or higher	29.2	21.0		28.5	23.6	
Alcohol (drinks per week), %						
Non-drinkers	80.2	88.6	<0.0001	81.0	78.4	<0.0001
<7	14.9	8.5		14.3	12.2	
≥7	4.9	2.9		4.7	9.4	
Tea consumption, %						
Never-drinkers	41.5	38.8	<0.0001	41.3	35.9	0.018
Monthly drinkers	12.1	11.8		12.0	12.8	
Weekly drinkers	24.3	24.7		24.3	23.6	
Daily drinkers	22.2	24.7		22.4	27.7	
Smoking status, %						
Never-smokers	69.6	67.8	<0.0001	69.6	46.3	<0.0001
Former smokers	10.4	15.9		10.8	19.6	
Current smokers	20.0	16.3		19.6	34.1	
Coffee intake (cups per week), mean (s.d.)	9.8 (8.5)	8.2 (7.9)	<0.0001	9.7 (8.5)	9.1 (8.1)	0.149

^aTwo-sided P-values were derived from χ^2 (for frequencies) or t-test (for means).

^aTwo-sided *P*-values were derived from χ^2 (for frequencies) or *t*-test (for means).

constant (P for heterogeneity = 0.58). We further examined the diabetes–HCC association, stratified by established and probable risk factors of HCC including BMI, cigarette smoking status and alcohol consumption status at baseline. No material differences across the subgroups were observed (data not shown).

We have previously published the relationships between markers of HBV and HCV infections and HCC risk in a case-control set nested within the cohort of The Singapore Chinese Health Study (Koh *et al*, 2011) (reproduced in Table 3). Chronic carriers of HBV (HBsAg-positive subjects) exhibited the highest risk of HCC (odds ratio (OR) = 24.79; 95% confidence interval (CI) = 8.61–71.34). Subjects with a history of primary infection of HBV (anti-HBc positive) without subsequent acquisition of immunity (anti-HBs negative) also were at an increased risk of HCC, though the risk level was an order of magnitude lower than those positive for HBsAg (OR = 2.01; 95% CI = 0.92–4.39). Conversely, subjects showing the presence of serum anti-HBs were not at an increased risk of HCC. Hepatitis C virus infection was rare in this population; only 5 HCC cases (5%) and 3 control

subjects (1%) were positive for anti-HCV. Nonetheless, HCV infection was statistically significantly related to HCC (OR = 10.12; 95% CI = 2.19–46.80) (Table 3).

Table 4 shows the diabetes–HCC association within the nested case-control set of HCC as described in Table 3. The diabetes–HCC association was mainly noted in the serology-negative subgroup (HR = 5.15; 95%CI = 2.08–12.73). The corresponding HR in the serology-positive subgroup was 1.01 (95% CI = 0.30–3.39). The interaction between diabetes and serological markers for hepatitis B or C on HCC risk was statistically significant (P for interaction = 0.012).

DISCUSSION

The present study shows that diabetics experienced a two-fold risk of HCC in this high-risk population with a relatively high prevalence of hepatitis B viral infection. The observed elevation in HCC risk among diabetics was statistically significant in the

Table 2. Diabetes in relation to hepatocellular carcinoma (HCC) in total subjects, males and females separately, and by duration of diabetes prior to baseline interview, Singapore Chinese Health Study 1993–2010

	Total number <i>N</i> (%)	HCC cases	HR (95% CI) ^a	HR (95% CI) ^b
All				
Non-diabetics	55 852 (91.1)	412	1.00	1.00
Diabetics	5469 (8.9)	87	2.64 (2.09–3.33)	2.14 (1.69–2.71)
Men				
Non-diabetics	24 933 (91.4)	303	1.00	1.00
Diabetics	2360 (8.6)	58	2.51 (1.89–3.32)	2.11 (1.58–2.81)
Women				
Non-diabetics	30 919 (90.9)	109	1.00	1.00
Diabetics	3109 (9.1)	29	3.27 (2.17–4.92)	2.14 (1.41–3.25)
Number of years of diabetes diagnosed prior to baseline interview				
Non-diabetics	55 852 (91.1)	412	1.00	1.00
< 5 years	2139 (3.5)	39	2.80 (2.01–3.89)	2.41 (1.73–3.35)
5–10 years	1341 (2.2)	21	2.53 (1.63–3.92)	2.14 (1.38–3.33)
≥ 10 years	1989 (3.2)	27	2.52 (1.7–3.71)	1.84 (1.24–2.72)

^aCrude hazard ratio.

^bAdjusted for age, year of recruitment, gender, dialect group, level of education, cigarette smoking status, alcohol intake frequency, body mass index, and consumption of coffee and tea; HR, hazard ratio; CI, confidence interval.

Table 3. HBV and HCV serology in relation to risk of hepatocellular carcinoma, Singapore Chinese Health Study (reproduced with permission from British Journal of Cancer)

HBV/HCV serology	Cases ^a (<i>n</i> = 92)	Controls ^a (<i>n</i> = 274)	Odds ratios ^b (95%CI)
Negative on all four markers	17	95	1.00
Anti-HBs positive	22	128	1.10 (0.54–2.22)
HBsAg positive	36	8	24.79 (8.61–71.34)
Anti-HBc positive, but anti-HBs negative	16	43	2.01 (0.92–4.39)
HBsAg positive or anti-HBc positive, but anti-HBs negative (HBV positive)	52	51	5.34 (2.44–11.67)
Anti-HCV positive	5	3	10.12 (2.19–46.80)

Reproduced with the kind permission of NPG from Koh *et al*, (2011). Abbreviation: anti-HBc, antibodies to hepatitis B core antigen; anti-HBs, antibodies to hepatitis B surface antigen; anti-HCV, antibodies to hepatitis C virus; HBV, hepatitis B virus; HBsAg, hepatitis B surface antigen; HCV, hepatitis C virus.

^aThe sum of cases and controls across all categories of HBV/HCV serology was greater than the total number of subjects as these serology groups were not mutually exclusive.

^bOdds ratios were calculated using conditional logistic regression models with further adjustment for the level of education (no formal education, primary, secondary or higher); CI, confidence interval.

Table 4. Diabetes in relation to the risk of hepatocellular carcinoma (HCC) according to the subjects' viral serology status, Singapore Chinese Health Study 1993–2010

	All subjects		Negative for all HBV or HCV		Positive for HBV or HCV ^a	
	Ca/Co ^b	OR (95% CI) ^c	Ca/Co ^b	OR (95% CI) ^c	Ca/Co ^b	OR (95% CI) ^c
Non-diabetics	71/249	1.00	26/204	1.00	45/45	1.00
Diabetics	21/25	2.55 (1.31–4.95)	12/17	5.15 (2.08–12.73) ^d	9/8	1.01 (0.30–3.39) ^d

^aPositive serologic markers including hepatitis B surface antigen (HBsAg), antibodies to hepatitis B core antigen (anti-HBc) or antibodies to hepatitis C virus (anti-HCV); HBV, hepatitis B virus.
^bNumber of cases/number of controls.
^cOdds ratios (ORs) were calculated using unconditional logistic regression models that also included age, year of recruitment, gender, dialect group, level of education, cigarette smoking status, alcohol intake frequency, body mass index and consumption of coffee and tea; CI, confidence interval.
^dP for the difference in the two odds ratios (or the interaction between diabetes and positive/negative serological markers of hepatitis B or C was 0.012.

subgroup of subjects totally devoid of viral HBV/HCV serology. Chronic carriers of viral hepatitis are at risk for diabetes (Naing *et al*, 2012; Schillie *et al*, 2012). Thus, our observation of a clear and highly significant association between diabetes and HCC risk among subjects completely devoid of viral hepatitis serology greatly strengthens the notion that the diabetes–HCC association is direct and causal in nature. The magnitude of risk for HCC among diabetics did not vary materially by the duration of diabetes among at-risk individuals. Diabetics who were diagnosed ≥ 10 years prior to cohort enrolment continued to exhibit a two-fold risk of HCC that was statistically significant and comparable to the risk of HCC among patients with the diagnosis of diabetes within 5 years prior to study enrolment. These latter findings strongly argue against the possibility of diabetes present in HCC patients as a consequence of late-stage liver disease. These results also suggest that medication treatment for diabetes did not have much impact on the risk of developing HCC.

Our results are consistent with the literature. Previous population-based studies representing observations from varied geographical locations, including Japan (Fujino *et al*, 2001; Tazawa *et al*, 2002; Shibata *et al*, 2003; Inoue *et al*, 2006), Korea (Jee *et al*, 2005), the US (Coughlin *et al*, 2004; El-Serag *et al*, 2004; Atchison *et al*, 2011) and European countries (Adami *et al*, 1996; Wideroff *et al*, 1997; Batty *et al*, 2004; Ogunleye *et al*, 2009; Zhou *et al*, 2010) have recorded a statistically significant increased incidence of HCC among patients with diabetes. The present study represents the first analysis from a southeast Asian population at relatively high risk for HCC. Overall, the epidemiologic evidence strongly supports a role for diabetes in the development of liver cancer across diverse populations with different levels of risk for HCC.

The biological mechanism of diabetes and its associated condition, obesity, in hepatocarcinogenesis is not well understood. The effect of increased serum insulin levels remains the most well studied and plausible mechanism for the association between diabetes and cancer (Ma *et al*, 1999), though increased insulin level alone may not be sufficient in causing HCC. Levels of insulin-like growth factor-1 (IGF-1) has been associated with the risk of colorectal (Ma *et al*, 1999; Grimberg and Cohen, 2000) and pancreatic cancer (Ohmura *et al*, 1990). Recent studies have suggested that circulating IGF-1 levels were associated with the increased risk of HCC (Elsammak *et al*, 2006; Su *et al*, 2010), and IGF-1 levels have been shown to promote liver tumour cell growth in experimental studies (Dunn *et al*, 1997; Pollak, 2000).

The development of primary liver cancer undergoes a long process in which chronic hepatic insult results in increased tissue turnover that leads to elevated risk of hepatocarcinogenesis. The intersecting effects of alcohol intake, chronic infection with hepatitis B and/or C viruses, obesity and the development of insulin resistance makes understanding the exact nature of the association between diabetes and HCC difficult, though the effects of elevated insulin levels due to insulin resistance remains the most

well studied effect of diabetes on HCC development. Increased levels of insulin in the body due to insulin resistance results in compensatory increases in the level of growth hormone (Le Roith, 1997), potentially leading to the downstream promotion of carcinogenesis. This effect has been shown to promote cell proliferation in the pancreas (Ohmura *et al*, 1990), and it is possible that similar effects would be observed in the liver.

In the present study, the effect of diabetes on HCC risk was confined to individuals negative for all HBV and HCV serological markers. Our results are consistent with those reported from a recent prospective study of Chinese in Taiwan. History of diabetes was associated with a statistically significant 5.4-fold risk of HCC among study participants negative for chronic HBV or HCV infections (Wang *et al*, 2009). Unlike the Singapore Chinese in whom hepatitis C infection has a negligible role in HCC, both HBV and HCV have been shown to contribute to the HCC burden in Taiwan Chinese (Lai *et al*, 2012). Significantly, the Taiwan study reports a 3.1-fold HCC risk for diabetes among subjects positive for anti-HCV but no increased HCC risk for diabetes among subjects positive for HBsAg, a marker of chronic HBV infection (Wang *et al*, 2009).

The issue of temporality has been raised for the association between diabetes and HCC, given the critical roles of the liver in the metabolism of glucose. Glucose is absorbed from the intestinal tract and transported via the portal vein to the liver, where glycogen is made and stored. Hepatocytes have specific cell membrane insulin receptors, where insulin secreted by the pancreatic β -cells can bind and facilitate the uptake and utilisation of glucose. The utilisation and storage of glucose as a fuel in humans is promoted by insulin. Excessive glycogen accumulation in the liver is seen in 80% of diabetic patients (Stone and Van Thiel, 1985). Patients showing solely excessive glycogen deposition may exhibit hepatomegaly and liver enzyme abnormalities (Chatila and West, 1996), an indication of liver damage. Diabetes increases the risk of steatohepatitis, which can progress to cirrhosis. Obesity is a comorbidity of diabetes and the underlying cause of nonalcoholic steatohepatitis (NASH). Clinical studies have shown that 40–100% patients with NASH are obese and that 20–75% of them have a history of adult-onset diabetes (Reid, 2001). Furthermore, the severity of fibrosis among NASH patients is positively associated with obesity and diabetes (Angulo *et al*, 1999). It is known that a high percentage (up to 70%) of patients with cryptogenic cirrhosis were obese and/or diabetic (Caldwell *et al*, 1999). These clinical data suggest that obesity/diabetes are risk factors for hepatic fibrosis and progression to cirrhosis, a recognised predisposing factor for HCC regardless of the underlying cause of hepatic cirrhosis (Zaman *et al*, 1985).

Conversely, individuals with cirrhosis have elevated insulin levels, suggesting insulin resistance or reduced degradation of insulin by the cirrhotic liver. Impaired insulin secretion from the pancreatic β -cells has been proposed as a cause for the hyperglycaemia. Hepatogenous diabetes, characterised by glucose

intolerance due to extensive liver damage, is estimated to be present in 30–60% of patients with cirrhosis (Garcia-Compean *et al*, 2009). Glucose intolerance in patients with cirrhosis has been found to be associated with low insulin secretion (Shah *et al*, 1995). These clinical findings suggest that elevated levels of circulating glucose or diabetes could be the result of a chronic liver disease that is aetiologically linked to the patient's HCC development. The long period between the onset of diabetes and diagnosis of HCC allowed us to address this issue. The equally strong elevation in HCC risk, we noted in diabetics diagnosed 10 or more years prior to cohort enrolment, supports a causal interpretation of the association rather than the occurrence of diabetes as a consequence of late-stage liver disease.

The strengths of this study include the prospective study design, long-term follow-up, (up to 15 years) and complete ascertainment of incident HCC cases among cohort participants. A comprehensive questionnaire for the collection of data on many potentially confounding factors allowed for their adjustment in the examination of the association between diabetes and HCC risk. The available data on HBV/HCV serology on a subset of the present study population also allowed us to rule out the possibility of confounding by HBV/HCV infection on the diabetes–HCC risk association.

However, there are a number of potential limitations in our study. First, we only use the baseline status of diabetes in our analysis, and with increasing age, there would be an increase in incidence of diabetes in the non-diabetic subpopulation. Such misclassification in diabetes based on the baseline interview only can potentially bias the magnitude of the diabetes–HCC risk association towards the null. We did not differentiate between type 1 and type 2 diabetes when we interviewed the participants. However, >95% of patients were diagnosed with diabetes after 35 years of age. Hence, virtually all diabetes cases in the present study would be considered as type 2 diabetes. We did not have data on the treatment for diabetes, including the use of medication, and the adequacy of diabetes control in our study population. The lack of a dose–response relationship for duration of diabetes prior to the baseline interview (Table 2) did not support a modifying effect of the use of diabetes medication or diabetes severity on HCC risk. Furthermore, diabetic patients using statins or metformin experienced reduced, but not increased risk of HCC (El-Serag *et al*, 2009; Chen *et al*, 2012). Therefore, the observed positive association between diabetes and the risk of developing HCC in this prospective study would be less likely due to the use of medication for diabetes. Finally, we did not measure serological markers of HBV/HCV infections on all cohort participants nor did we assess the history of cirrhosis in our baseline interviews. Therefore, we were unable to examine these confounders in a whole cohort statistical analysis of the diabetes–HCC risk association. However, an appropriately designed nested case-control study within a whole cohort is universally recognised to yield valid findings, and we have conducted such a study to examine the potential modifying role of viral hepatitis in the diabetes–HCC association. We noted that the diabetes–HCC association was seen only among subjects negative for all tested HBV/HCV serology, indicating a lack of influence by viral hepatitis or its clinical sequelae (cirrhosis) on the diabetes–HCC association.

In conclusion, the present study demonstrates a statistically significant, positive association between diabetes status at baseline and elevated risk of developing HCC among Singaporean Chinese, a population with relatively high prevalence of HBV infection and HCC incidence. This positive diabetes–HCC risk association present in individuals without chronic infection with HBV or HCV virus suggests an independent role of diabetes in HCC development, which has an important implication in public health, given the worldwide increasing incidence of type 2 diabetes.

ACKNOWLEDGEMENTS

We thank Siew-Hong Low of the National University of Singapore for supervising the field work of the Singapore Chinese Health Study. The Singapore Cancer Registry assisted with the identification of cancer and mortality outcomes via database linkages. This study was supported by National Institutes of Health (NCI R01 CA55069, R35 CA53890 and R01 CA80205, R01 CA144034).

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