The clinical behavior and survival of patients with hepatocellular carcinoma and a family history of the disease

Jihyun An | Seheon Chang | Ha Il Kim | Gi-Won Song | Ju Hyun Shim

1Gastroenterology and Hepatology, Hanyang University College of Medicine, Guri, Korea
2Internal Medicine, Myongji Saint Mary’s Hospital, Seoul, Korea
3Gastroenterology, Asan Medical Center, University of Ulsan College of Medicine, Seoul, Korea
4Asan Liver Center, Asan Medical Center, University of Ulsan College of Medicine, Seoul, Korea
5Surgery, Asan Medical Center, University of Ulsan College of Medicine, Seoul, Korea

Correspondence
Ju Hyun Shim, Department of Gastroenterology, Asan Medical Center, University of Ulsan College of Medicine, 88 Olympic-ro 43-gil, Songpa-gu, Seoul, 05505, Korea.
Email: s5854@amc.seoul.kr

Funding information
This study was supported by Basic Science Research Program through the National Research Foundation of Korea funded by the Ministry of Science and ICT (NRF-2017R1E1A1A01074298) and grant 2015-660 from the Asan Institute for Life Sciences of Asan Medical Center.

Abstract
Purpose: Familial clustering is a common feature of hepatocellular carcinoma (HCC) as well as a risk factor for the disease. We aimed to assess whether such a family history affected prognostic outcomes in patients with HCC diagnosed at different stages of the disease.

Materials/Methods: This hospital registry-based cohort study included 5484 patients initially diagnosed with HCC. Individual family histories of cancer were obtained by interview and reported by trained nurses who constructed three-generation pedigrees. Overall survival data were compared between cases with and without first-degree relatives affected by HCC, with adjustment for other potential predictors.

Results: Of 5484 patients, 845 (15.4%) had first-degree relatives with a history of HCC. Family history was associated with longer survival in the entire cohort (adjusted hazard ratio [HR] 0.89, 95% confidence interval [CI] 0.80-0.98, \( P = .025 \)). A significant trend for reduced risk of death with increasing number of affected family members was also observed (\( P \) for trend = 0.018). The stage-stratified analysis showed that the presence of family history was especially associated with a reduced risk of death in the subset of patients with HCC at a (very) early stage (adjusted HR 0.83, 95% CI 0.69-0.99; \( P = .042 \)). The proportion of cases receiving curative treatment was also higher in early-stage patients with a family history (72.6% vs 63.3%; \( P < .001 \)).

Conclusions: A first-degree family history of the disease is a prognostic factor for improved survival in patients with HCC, especially in those whose tumors can be cured by radical treatments.

KEYWORDS
clustering, family, liver cancer, prognosis, treatment

Dr. An and Chang contributed equally to this work.

This is an open access article under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited.

© 2019 The Authors. Cancer Medicine published by John Wiley & Sons Ltd.

Cancer Medicine. 2019;8:6624-6633.
1 | INTRODUCTION

Evidence has accumulated over many years of a relationship between the risk of developing a specific cancer and a family history of the disease. In addition, numerous studies have reported positive or negative effects of a family history on the prognostic outcomes of patients with different types of cancer. Most attention in this matter has been given to malignancies of the digestive and reproductive systems, which are the most common cancers in both men and women.

Interestingly, hepatocellular carcinoma (HCC), which is the third leading cause of cancer deaths globally despite its lower ranking for incidence, has been observed to cluster within families sharing genes and environments. The familial clustering of the disease was found to be unrelated to a viral etiology of hepatitis B in both Asians and Europeans, but understandably increased in subjects with HCC due to vertically transmitted hepatitis B virus (HBV). Because of this strong familial association of HCC risk, the former American guidelines recommend routine surveillance of hepatitis B carriers of all ages with family histories of HCC, like for high-risk cirrhotic patients. An international study of the prognostic role of family history in HCC patients concluded that the familial cancer group had better survival than its sporadic counterpart, and suggested that this was due to the cancers being detected at an earlier stage of tumor growth and liver damage. However, that (Hong Kong) study gave only unadjusted estimates without detailed information on family membership and generations.

Our aim in this study was to examine the incidence of a family history of cancer at the time of HCC diagnosis, and to investigate the association between familial cancer clustering and survival outcomes over time in a large clinical set of patients first diagnosed with HCC; since the patients were ethnically homogeneous Koreans the data should not be skewed by racial and environmental biases. We also wanted to see whether the presence of affected blood relatives influenced treatment decision-making in clinical practice.

2 | PATIENTS AND METHODS

2.1 | Data sources and collection

Approval of the Institutional Review Board of our center (IRB No. 2016-0683) was obtained for this large registry-based retrospective cohort study, and treatment-naive patients initially diagnosed with HCC by a three-digit diagnostic code specified by the seventh revision of the International Classification of Diseases (ICD-7) were identified from our prospectively constructed hospital-based cancer registry. This registry is a part of the National Cancer Registration Program and has been described in previous studies from our center. Health-related behaviors together with relevant demographic factors and clinical information on the patients were reviewed from their inpatient and outpatient medical records using the anonymized clinical database system of our institution (Asan Biomedical research Environment, ABLE). Demographic and socioeconomic data were collected from computerized admission documents completed by trained nurses during patient interviews employing a structured questionnaire. The nursing charts included information on educational level, substance use (tobacco and alcohol), past and present medical histories, and basic anthropometric data. Medical histories of family members were recorded in detail on each patient’s chart, together with pedigrees containing information including history and sites of cancers, and causes of death of close blood relatives (first- and second-degree relatives). We also examined laboratory data related to liver function and viral hepatitis, and checked radiological results to determine stage of HCC based on the size and number of tumors, vascular invasion, and extrahepatic metastasis; in addition, HCC treatment modalities and the associated survival outcomes were obtained from the ABLE system and database of the National Population Registry of the Korea National Statistical Office using the unique personal identification numbers of the patients.

2.2 | Patient details

Patients over 20 years of age who were diagnosed as having HCC and underwent treatment for the disease for the first time between 2007 and 2011 were included in this study (n = 8246, Figure 1). Of these, 2762 were initially excluded for the following reasons: 2712 had had previous treatment for HCC prior to visiting our center; 36 had concurrent non-HCC malignancies; and 14 of whom did not have complete records of family health histories. The diagnosis of HCC was based on either pathological or radiological findings in accord with international guidelines. Cirrhosis of the liver was also defined either histologically or based on radiographic abnormalities (ie, nodular changes of liver morphology, spleenomegaly, gastrointestinal varices, or ascites). Stages of HCC at diagnosis were classified by the Barcelona Clinic Liver Cancer (BCLC) system. The HCC treatment for each patient was principally decided according to their hierarchy of efficacy in lengthening life.

Surgical resection was based on the anatomical segments of the liver whenever possible. Radiofrequency ablation was performed percutaneously under sonographic or computed tomographic guidance. Transarterial chemoembolization (TACE) was usually carried out using a mixture of iodized oil and cisplatin or adriamycin, and absorbable gelatin sponge particles. Most of transplant cases (97.4%) received grafts from living-related donors.
2.3 | Family histories

Family histories of cancer were routinely taken by a trained nurse using a structured questionnaire based on a three-generation pedigree. For each relative, the study participant was asked about any serious medical conditions and whether the relative was still alive, or, if the relative had died, the date and cause of death. Positive family histories of cancer were recorded according to type of cancer (HCC or cancer of all types other than HCC) and the generation of the affected relatives (first-degree or second-degree). First-degree relatives included parents, siblings, and offspring; and second-degree relatives included aunts, uncles, nieces, nephews, and grandparents. A patient with a family history of cancer in both first- and second-degree relatives was regarded as having a first-degree history. Only histories of cancer in one or more first-degree relatives that were reliably reported were included as established family histories in the final analysis.

2.4 | Statistical analysis

The main aim of the statistical analysis was to compare the overall survival of patients with and without a family history of cancer. The survival analysis was censored on 31 December 2016, and deaths occurring up to that time were considered events. In general, overall survival, rather than progression-free survival, is the most appropriate end-point of studies of HCC patients, most of whom have underlying liver disease, or some other serious disorder, especially as patients receive different types of curative and non-curative anti-cancer treatments. Using multivariate Cox proportional hazards models, we estimated the hazard ratio (HR) for death of the familial group compared with the sporadic group as the reference control. The HRs were adjusted for age, gender, level of education, body mass index (BMI), smoking, alcohol consumption, etiology of liver disease, presence of cirrhosis, laboratory results related to liver function, tumor stage at diagnosis, and serum alpha-fetoprotein (AFP). A backward elimination approach involving candidate variables with P-values < .10 in the univariate analysis was used in the multivariable analysis.

Differences in clinical and pathologic parameters between the familial and sporadic groups were analyzed with the $X^2$ test or Fisher’s exact probability test, as appropriate.

Stratified analyses were also performed by number of affected family members, and cancer stage. A two-sided P-value < .05 was considered statistically significant.

3 | RESULTS

3.1 | Family histories of study subjects

Of 5484 HCC patients included, 1859 (33.9%) had at least one relative with some form of cancer (Table S1); 1823 had family histories in first-degree relatives and 36 in second-degree relatives. When the family history was limited to HCC, 870 (15.9%) had a family history of HCC. 845 (15.4% of the entire cohorts) had family histories in one or more first-degree relatives and 25 in second-degree relatives. Nine of the 870 patients had both first- and second-degree family histories of HCC. A total of 1213 patients (22.1%) had family histories of non-HCC cancers.

3.2 | Demographic and clinical characteristics according to presence or absence of a family history of HCC

Table 1 presents the baseline characteristics of patients and tumors at the time of HCC diagnosis. The median age of the entire subjects was 56 years (interquartile range [IQR], 49-63 years), and the majority of the patients were male (80.7%) and had HBV infections (80.6%). Liver cirrhosis was observed in 4431 patients (80.8%). Females and younger patients, non-diabetics, and never-drinkers were more common among individuals with first-degree family histories of HCC than among those without such histories ($P$’s < .05; Table 1). Those with first-degree family histories of HCC also had a higher proportion of non-HCC cancer histories ($P = .002$). In addition, subjects with HBV or Child-Pugh class A were more common and hepatitis C virus (HCV) carriers were less common in the familial group ($P$’s < .05). Size and multiplicity
### TABLE 1 Demographic and Hepatic Characteristics by Family History of Hepatocellular Carcinoma (n = 5484)

| Variable                          | Family history (n = 845) | No family history (n = 4639) | P value |
|----------------------------------|--------------------------|-----------------------------|---------|
| **Demographic factor**           |                          |                             |         |
| Male sex                         | 656 (77.6%)              | 3768 (81.2%)                | .015    |
| Age, years                       | 54 (49-61)               | 56 (49-64)                  | <.001   |
| Body mass index, kg/m²           | 24.1 (22.1-26.1)         | 24.1 (22.1-26.1)            | .608    |
| Alcohol consumption              |                          |                             | .007    |
| Never                            | 323 (38.2%)              | 1515 (32.7%)                |         |
| Former                           | 360 (42.6%)              | 2140 (46.1%)                |         |
| Current                          | 162 (19.2%)              | 984 (21.2%)                 |         |
| Smoking status                   |                          |                             | .201    |
| Never                            | 348 (41.2%)              | 1762 (38.0%)                |         |
| Former                           | 339 (40.1%)              | 1959 (42.2%)                |         |
| Current                          | 158 (18.7%)              | 914 (19.8%)                 |         |
| Education, years                 |                          |                             | .143    |
| ≤9                               | 310 (36.7%)              | 1857 (40.0%)                |         |
| 10-12                            | 298 (35.3%)              | 1597 (34.4%)                |         |
| >12                              | 237 (28.0%)              | 1185 (25.6%)                |         |
| Diabetes                         | 142 (16.8%)              | 975 (21.1%)                 | .005    |
| Hypertension                     | 201 (23.8%)              | 1250 (27.0%)                | .052    |
| Family history of non-HCC cancers| 219 (25.9%)              | 977 (21.1%)                 | .002    |
| **Liver disease-related factor** |                          |                             |         |
| Etiology of liver disease        |                          |                             |         |
| Hepatitis B virus infection      | 762 (90.2%)              | 3347 (72.1%)                | <.001   |
| Hepatitis C virus infection      | 28 (3.3%)                | 513 (11.1%)                 | <.001   |
| Liver cirrhosis                  | 690 (81.7%)              | 3741 (80.6%)                | .491    |
| Ascites                          | 87 (10.3%)               | 562 (12.1%)                 | .132    |
| Platelet count (x10³/mm³)        | 143 (98-190)             | 137 (94-189)                | .416    |
| Serum albumin (g/dL)             | 3.7 (3.3-4.0)            | 3.6 (3.1-4.0)               | <.001   |
| Serum bilirubin (mg/dL)          | 1.0 (0.8-1.4)            | 1.0 (0.8-1.5)               | .078    |
| International normalized ratio (INR) | 1.08 (1.03-1.17) | 1.09 (1.03-1.19) | .259 |
| Serum creatinine (mg/dl)         | 0.8 (0.7-0.9)            | 0.8 (0.7-1.0)               | .256    |
| Child-Pugh class                 |                          |                             | <.001   |

(Continues)
of tumors were not associated with a family history of HCC; BCLC stages were similar in the two groups \((P = .280)\), and TACE and surgical resection were the most common primary anti-HCC treatments in both groups. There was also no difference in the time interval between diagnosis of HCC and initiation of treatment in the two groups, this interval being generally less than one month in new cases \((P = .306)\). Curative therapies such as resection, transplantation, and local ablation were initially chosen in 49.1% of the patients with a family history, significantly higher than the 44.0% among those without family histories \((P < .001)\), and the converse was true for non-curative options (50.9% vs. 56.0%; \(P < .001\); Table 1).

### 3.3 Effect of a family history of HCC on survival in patients with HCC

During a median observation period of 4.0 years (IQR 1.0-6.6 years), 3228 of the 5484 patients (58.9%) died of any cause. Of those who died, 89.4% \((n = 2886)\) were treated and followed-up in our tertiary center for at least the last 6 months before death, and this proportion did not depend on the presence or absence of a family history of HCC (91.7% vs 89.0%, \(P = .083\)). The 3-, 5-, and 7-year estimated overall survival rates were 55.5%, 46.7%, and 41.1%, respectively in the entire population. Kaplan-Meier log-rank analysis revealed a significant increase of survival time in the patients with a history of HCC (52.1% vs 45.7% at 5 years, \(P < .001\); Figure 2A).

In multivariate Cox models after adjustment for co-predictors (ie, age, gender, family history of non-HCC cancer, smoking and drinking habitus, level of education, BMI, etiology of liver disease, presence of cirrhosis, Model for end-stage liver disease [MELD] score, platelet count, serum AFP levels, BCLC stage, and infiltrative type of tumor), a family history of HCC was independently associated with improved overall survival (adjusted hazard ratios [HRs], 0.89; 95% confidence interval [CI], 0.80 to 0.98; \(P = .025\); Table 2).

The relationship between family history and outcomes according to the number of family members with HCC was also investigated. Although the majority of patients with a family history reported only one affected relative, there was a significant trend for an increased reduction in death risk with increasing number of affected family members after adjustment for demographic and tumoral factors \((P\ for\ trend = .018;\ Figure\ 2B)\).

### 3.4 Survival analysis stratified by stage of HCC

We further analyzed the prognostic effect of a family history of HCC in patients who were at different initial stages of HCC when diagnosed. A family history of HCC was positively correlated with overall survival in patients with BCLC 0 or A stage HCC (adjusted HR 0.83, 95% CI 0.69-0.99; \(P = .042\)), as shown in Table S2 and Figure 3. The proportion of cases receiving curative treatment was also higher in

![Figure 2](image_url)
early-stage patients with a family history (72.6% vs 63.3%;  \( P < .001 \)). In terms of specific anti-cancer treatments, surgical resection was more frequently performed in patients with familial histories than in those without histories (58.5% vs 47.3%, \( P < .001 \); Figure 4). TACE treatment were more common in the latter group (26.7% vs 35.0%, \( P = .001 \)).

When the analysis was restricted to patients with advanced stage HCC, we observed no relationship between survival and family history (HRs [95% CIs] 0.88 [0.68-1.13] for BCLC B stage; 0.98 [0.85-1.12] for BCLC C stage; and 0.77 [0.48-1.24] for BCLC D stage; \( P \)'s > .05). Among the patients with more advanced HCC, the primary treatment pattern was similar in the two groups (curative vs non-curative treatments 25.8% vs 24.5%, \( P = .603 \)).

### DISCUSSION

Several studies have demonstrated that a family history of HCC increases the risk of developing HCC, after adjusting for proven...
risk factors including HBV and HCV infection. Familial aggregation of liver cancer has been frequently reported, although there have been no suggestions of an underlying genetic predisposition for hepatic neoplasms. However, the influence of a family history of HCC on subsequent outcomes in patients with the established disease is controversial. In this large, well-characterized, hospital-based cohort study, the incidence of a family history in a new HCC series was about 16%, and familial clustering of HCC was associated with a reduced risk of overall mortality in patients with the disease. Curative treatment, especially surgical resection, was also more common in patients with a positive family history.

The role of family history has been investigated as a prognostic factor in several types of malignancy, and diverse correlations have been observed. The presence of familial cancer in stomach, breast, prostate, and colon cancer patients had protective effects, as in our HCC series, but no such effects were seen for brain and ovarian cancers. The univariate findings in a study by a group in Hong Kong pointed to better survival in familial HCC patients (who accounted for...
approximately 10% of the total), especially in an early-stage non-metastatic sub-cohort, a result that appears to resemble the present findings based on a more intensive and less confounded analysis. Although another Chinese investigation with 12% familial cases did not find a significant relationship between family history of HCC and survival after resection, the fact that it was restricted to surgical patients limits its generalizability.

There are some possible explanations for the association between familial cancer clusters and prognoses. First, cancer patients with a family history may more often present with early-stage disease, perhaps because they adhere more rigorously to cancer screening through greater awareness of the implications of the disease, as has been found in studies of prostate, breast, and gastric cancer. However, a family history did not influence the initial profile of tumor stages among our new HCC cases, and the familial effect persisted after controlling for differences in stage. Second, genetic differences in inherent tumor biology between patients with and without a family history may influence cancer mortality. A Swedish population-based study found a higher proportion of indolent subtypes in familial leukemia, whereas familial cases of ovarian cancers had a more aggressive course with poorer survival. Functional genetic or immunologic polymorphisms may well determine not only susceptibility to specific diseases, but also individual responses to cancer treatment. Third, health-related behavioral changes including regular physical activity, stopping smoking and drinking, and a healthy diet and nutrition may contribute to the superior disease course, since these factors have been shown to have anti-cancer effects in HCC.

The beneficial effect of a family history in our study appeared greater among patients with early cancers. Since, unlike intermediate or advanced HCC, for which there is a single standard treatment, early-stage HCC can be treated in a variety of ways, from radical resection or liver transplantation with more curative intent, to less potent but more convenient interventional procedures. This association suggests that familial HCC patients are more likely to seek medical attention, a factor which is seldom controllable; they may therefore have a higher probability of, and even a preference for, undergoing more effective, albeit more invasive, treatment. Patients with no family history may have less understanding of the therapeutic options and prognoses, and less opportunity to consult someone with experience of HCC, which may influence doctor-patient decision-making regarding treatment modality. Our results show that the presence of a family history was closely associated with the receipt of formally recommended definitive surgery, rather than palliative TACE, as the initial therapy in equivalent early cases. In addition to accurate medical knowledge of therapeutic risks and benefits, a variety of factors including fear of complications related to surgery, concern about recurrence, and advice from the patient's family are usually involved in the choice of cancer treatment. Our results suggest that such behavioral benefits associated with family history increase with increasing number of affected relatives. The absence of an association...
between presence of a family history and survival in patients who had opted for and received surgical resection in our and prior Chinese studies may indicate that a family history is mainly influential during clinical decision-making.^{19}

Several limitations of this study deserve comment. First, because we relied on self-reported family histories, family history status may have been misclassified. However, self-reported data have repeatedly been shown to be reliable in prior studies.^{43,44} Because the data on family history were collected at the time of first diagnosis of HCC, prior to the initial treatment, any errors in recall should not have influenced the association with patient outcome.^{45} In addition, we tried to minimize ascertainment bias, and thus in the end only tested the familial effect of first-degree relatives, and the latter are presumably reported quite reliably.^{26,46} Second, differences in adherence to medical management could introduce a certain bias. However, the interval between diagnosis and treatment among all the new cases, and the follow-up compliance in our center among the patient who ultimately died, was not affected by a family history of HCC during the period of observation, and indeed patient compliance is likely to be more reliable in a high-volume hospital like ours.^{47}

In conclusion, this investigation revealed that patients with HCC who had a first-degree family history of the disease survived better than those without such a family history. This familial benefit was stronger among early-stage patients, whose attitudes would have a greater impact on treatment decisions and subsequent outcomes than among those at a later stage. The molecular and genetic factors underlying familial-clustered HCC remain to be elucidated.

ACKNOWLEDGMENT

We thank Drs Danbi Lee, Kang Mo Kim, Young-Suk Lim, Han Chu Lee, Young-hwa Chung, and Yung Sang Lee, department of Gastroenterology, Asan Medical Center for data collection. None received compensation for their work. This study was supported by Basic Science Research Program through the National Research Foundation of Korea funded by the Ministry of Science and ICT (NRF-2017R1E1A1A01074298).

AUTHOR CONTRIBUTIONS

J An and S Chang contributed to study concept and design, acquisition, analysis and interpretation of data, statistical analysis, drafting of the manuscript, critical revision of the manuscript for important intellectual content. HI Kim and G-W Song contributed to acquisition of data and critical revision of the manuscript for important intellectual content. JH Shim contributed to study concept and design, interpretation of data, drafting of the manuscript, critical revision of the manuscript for important intellectual content and study supervision.

DATA AVAILABILITY STATEMENT

Data used in this research are available to other research teams upon request to the corresponding author.

ORCID

Jihyun An https://orcid.org/0000-0002-0110-0965
Ju Hyun Shim https://orcid.org/0000-0002-7336-1371

REFERENCES

1. Lu KH, Wood ME, Daniels M, et al. American society of clinical oncology expert statement: collection and use of a cancer family history for oncology providers. *J Clin Oncol*. 2014;32:833-840.
2. Bratt O, Drevin L, Akre O, Garmo H, Stattin P. Family history and probability of prostate cancer, differentiated by risk category: a nationwide population-based study. *J Natl Cancer Inst*. 2016;108:dwj110.
3. Lowery JT, Ahnen DJ, Schroy PC 3rd, et al. Understanding the contribution of family history to colorectal cancer risk and its clinical implications: a state-of-the-science review. *Cancer*. 2016;122:2633-2645.
4. Altieri A, Bermejo JL, Hemminki K. Familial risk for non-Hodgkin lymphoma and other lymphoproliferative malignancies by histopathologic subtype: the Swedish family-cancer database. *Blood*. 2005;106:668-672.
5. Kupelian PA, Kupelian VA, Witte JS, Macklis R, Klein EA. Family history of prostate cancer in patients with localized prostate cancer: an independent predictor of treatment outcome. *J Clin Oncol*. 1997;15:1478-1480.
6. Lee M, Reilly M, Lindstrom LS, Czene K. Differences in survival for patients with familial and sporadic cancer. *Int J Cancer*. 2017;140:581-590.
7. Lindstrom LS, Hall P, Hartman M, Wiklund F, Gronberg H, Czene K. Familial concordance in cancer survival: a Swedish population-based study. *Lancet Oncol*. 2007;8:1001-1006.
8. Chan JA, Meyerhardt JA, Niedzwiecki D, et al. Association of family history with cancer recurrence and survival among patients with stage III colon cancer. *JAMA*. 2008;299:2515-2523.
9. Han MA, Oh MG, Choi JJ, et al. Association of family history with cancer recurrence and survival in patients with gastric cancer. *J Clin Oncol*. 2012;30:701-708.
10. Jobsen JJ, van der Palen J, Brinkhuis M, Ong F, Struikmans H. Long-term effects of first degree family history of breast cancer in young women: recurrences and bilateral breast cancer. *Acta Oncol*. 2016;55:449-454.
11. Stoffel EM, Mercado RC, Kohlmann W, et al. Prevalence and predictors of appropriate colorectal cancer surveillance in Lynch syndrome. *Am J Gastroenterol*. 2010;105:1851-1860.
12. Hemminki K, Li X. Familial liver and gall bladder cancer: a nationwide epidemiological study from Sweden. *Gut*. 2003;52:592-596.
13. Turati F, Edelfonti V, Talamini R, et al. Family history of liver cancer and hepatocellular carcinoma. *Hepatology*. 2012;55:1416-1425.
14. Yang Y, Wu QJ, Xie L, et al. Prospective cohort studies of association between family history of liver cancer and risk of liver cancer. *Int J Cancer*. 2014;135:1605-1614.
15. Loomba R, Liu J, Yang HI, et al. Synergistic effects of family history of hepatocellular carcinoma and hepatitis B virus infection
on risk for incident hepatocellular carcinoma. Clin Gastroenterol Hepatol. 2013;11(1636–1645):e1631–1633.

16. Wan DW, Tzimas D, Smith JA, et al. Risk factors for early-onset and late-onset hepatocellular carcinoma in Asian immigrants with hepatitis B in the United States. Am J Gastroenterol. 2011;106:1994-2000.

17. Hassan MM, Kaseb A, Li D, et al. Association between hypothyroidism and hepatocellular carcinoma: a case-control study in the United States. Hepatology. 2009;49:1563-1570.

18. Bruix J, Sherman M. Management of hepatocellular carcinoma: an update. Hepatology. 2011;53:1020-1022.

19. Dai WC, Fan ST, Cheung TT, et al. The impact of family history of hepatocellular carcinoma on its patients' survival. Hepatobiliary Pancreat Dis Int. 2012;11:160-164.

20. Kim HJ, Cho JH, Lyu Y, Lee SH, Hwang KH, Lee MS. Construction and validation of hospital-based cancer registry using various health records to detect patients with newly diagnosed cancer: experience at Asan Medical Center. J Prev Med Public Health. 2010;43:257-264.

21. Seo HJ, Oh IH, Yoon SJ. A comparison of the cancer incidence rates between the national cancer registry and insurance claims data in Korea. Asian Pac J Cancer Prev. 2012;13:6163-6168.

22. Shin SY, Lyu Y, Shin Y, et al. Lessons learned from development of de-identification system for biomedical research in a Korean tertiary hospital. Healthc Inform Res. 2013;19:102-109.

23. Shin SY, Park YR, Shin Y, et al. A De-identification method for bilingual clinical texts of various note types. J Korean Med Sci. 2015;30:7-15.

24. European Association for the Study of the Liver, European Organisation for Research and Treatment of Cancer. EASL-EORTC clinical practice guidelines: management of hepatocellular carcinoma. J Hepatol. 2012;56:908-943.

25. Shim JH, Han S, Shin YM, et al. Optimal measurement modality and method for evaluation of responses to transarterial chemoembolization of hepatocellular carcinoma based on enhancement criteria. J Vasc Interv Radiol. 2013;24:316-325.

26. Zgas A, Anton-Culver H. Validation of family history data in cancer family registries. Am J Prev Med. 2003;24:190-198.

27. Huang LP, De Sanctis Y, Shan MH, et al. Weak correlation of overall survival and time to progression in advanced hepatocellular carcinoma [abstract]. J Clin Oncol. 2017;35:233.

28. Reig M, Rimola J, Torres F, et al. Postprogression survival of patients with advanced hepatocellular carcinoma: rationale for second-line trial design. Hepatology. 2013;58:2023-2031.

29. Patel T, Harnois D. Assessment of response to therapy in hepatocellular carcinoma. Ann Med. 2014;46:130-137.

30. Liu L, Li L, Zhou S, et al. Familial correlations of onset age of hepatocellular carcinoma: a population-based case-control family study. PLoS ONE. 2014;9:e108391.

31. Huang J, Zhang Y, Chen M, Huang J, Xu L, Chen M. Family history of hepatocellular carcinoma is not associated with its patients' prognosis after hepatectomy. World J Surg Oncol. 2013;11:280.

32. Melvin JC, Wulaningsih W, Hana Z, et al. Family history of breast cancer and its association with disease severity and mortality. Cancer Med. 2016;5:942-949.

33. Williams KP, Reiter P, Mabiso A, Maurer J, Paskett E. Family history of cancer predicts Papanicolaou screening behavior for African American and white women. Cancer. 2009;115:179-189.

34. Wallner LP, Sarma AV, Lieber MM, et al. Psychosocial factors associated with an increased frequency of prostate cancer screening in men ages 40 to 79 years: the Olmsted County study. Cancer Epidemiol Biomarkers Prev. 2008;17:3588-3592.

35. Tracy KA, Quillin JM, Wilson DB, et al. The impact of family history of breast cancer and cancer death on women’s mammography practices and beliefs. Genet Med. 2008;10:621-625.

36. Dragani TA. Risk of HCC: genetic heterogeneity and complex genomics. J Hepatol. 2010;52:252-257.

37. Sampson JN, Wheeler WA, Yeager M, et al. Analysis of heritability and shared heritability based on genome-wide association studies for thirteen cancer types. J Natl Cancer Inst. 2015;107:dvj279.

38. Humpel N, Magee C, Jones SC. The impact of a cancer diagnosis on the health behaviors of cancer survivors and their family and friends. Support Care Cancer. 2007;15:621-630.

39. Patterson F, Wileyto EP, Segal J, Kurz J, Glanz K, Hanlon A. Intention to quit smoking: role of personal and family member cancer diagnosis. Health Educ Res. 2010;25:792-802.

40. Hamed MA, Ali SA. Non-viral factors contributing to hepatocellular carcinoma. World J Hepatol. 2013;5:311-322.

41. Hawley ST, Griggs JJ, Hamilton AS, et al. Decision involvement and receipt of mastectomy among racially and ethnically diverse breast cancer patients. J Natl Cancer Inst. 2009;101:1337-1347.

42. Vornanen M, Konttinen H, Kaariainen H, et al. Family history and perceived risk of diabetes, cardiovascular disease, cancer, and depression. Prev Med. 2016;90:177-183.

43. Kerber RA, Slattery ML. Comparison of self-reported and database-linked family history of cancer data in a case-control study. Am J Epidemiol. 1997;146:244-248.

44. Aitken J, Bain C, Ward M, Siskind V, MacLennan R. How accurate is self-reported family history of colorectal cancer? Am J Epidemiol. 1995;141:863-871.

45. Fiedlerling J, Shams AZ, Haug U. Validity of self-reported family history of cancer: a systematic literature review on selected cancers. Int J Cancer. 2016;139:1449-1460.

46. Mai PL, Garceau AO, Graubard BI, et al. Confirmation of family cancer history reported in a population-based survey. J Natl Cancer Inst. 2011;103:788-797.

47. Hebert-Croteau N, Brisson J, Lemaire J, Latreille J, Pineault R. How to cite this article: An J, Chang S, Kim HI, Song G-W, Shim JH. The clinical behavior and survival of patients with hepatocellular carcinoma and a family history of the disease. Cancer Med. 2019;8:6624–6633. https://doi.org/10.1002/cam4.2543

SUPPORTING INFORMATION
Additional supporting information may be found online in the Supporting Information section at the end of the article.