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Hepatocellular Carcinoma in Non-alcoholic Fatty Liver Disease: Epidemiology, Pathogenesis, and Prevention

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Abstract

Hepatocellular cancer (HCC) is the fifth most prevalent cancer worldwide and the third leading cause of cancer-related deaths. Non-alcoholic fatty liver disease (NAFLD), a spectrum of hepatic disorders associated with obesity and the metabolic syndrome, is a recognized risk factor for HCC. NAFLD that is advanced to cirrhosis carries the highest risk for HCC, but there is increasing concern that NAFLD-associated HCC may also occur in non-cirrhotic liver. As NAFLD is rapidly becoming the most common liver condition, it has been implicated in the worrisome trend of rising HCC incidence in a number of countries, which may offset successful measures in reducing the effect of virus-related liver cancer. Independently or in synergy with cirrhosis, NAFLD may provide a special oncogenic microenvironment through its pathogenic association with chronic nutrient excess and adipose tissue remodeling, characterized by pro-inflammatory adipokine profiles, lipotoxicity, altered hepatocellular bioenergetics, and insulin resistance. Better understanding of this complex process, and development of reliable biomarkers for HCC will be critical for early recognition and risk prediction. Moreover, correcting deranged lipid metabolism and restoring insulin sensitivity by lifestyle measures and targeted pharmacotherapy holds major promise for effective prevention of NAFLD-associated HCC.

Epidemiology of HCC associated with NAFLD

HCC emerging in NAFLD that has advanced to cirrhosis

Autopsy records and imaging data indicate that 70–80% of individuals with obesity have increased liver fat content, suggesting that every third person in the general adult population of the USA and other industrialized countries has some form of NAFLD. Recent reports indicate that NASH can be verified by histological evaluation in up to 47% of all NAFLD cases among obese individuals. Moreover, there are 26 million individuals with manifest diabetes in the USA, and the presence of biopsy-proven NAFLD among people with diabetes has been reported to reach up to 74%, with evidence of NASH in every fourth case. The pool of individuals at risk of NAFLD advancing to NASH and cirrhosis is therefore a substantial one.

There is good evidence that HCC may complicate NAFLD cirrhosis in the absence of other risk factors. In a prospective study conducted over 10 years, HCC developed in 10 out of 149 patients with NAFLD cirrhosis in comparison with 25 out of 147 American patients with HCV cirrhosis.
another analysis from the USA, the yearly cumulative incidence of NAFLD cirrhosis was 2.6%, whereas that of HCV cirrhosis was 4.0%.[25] In a Japanese study, HCC developed in 11.3% of patients with NAFLD cirrhosis compared with 30.5% of patients with HCV cirrhosis over 5 years.[26] The overall contribution of NAFLD to the burden of liver cancer appears therefore comparable with that of major established causes of HCC such as chronic hepatitis B virus, chronic HCV, and alcoholic liver disease (Fig. 1).

Based on the significant worldwide prevalence of obesity and diabetes, there is an increasing probability that NAFLD coexists with chronic liver disease of other etiologies, increasing the risk of HCC. A study in Taiwan found that body mass index (BMI) ≥ 30 kg/m² was independently associated with a four-fold increased risk of HCC among HCV-seropositive subjects, while this risk was only two-fold higher among obese subjects who were HCV-seronegative.[27] In a French study, the combined presence of increased BMI and clinically manifest diabetes predicted an adjusted hazard ratio of 6.0 for HCC among 431 patients with cirrhosis associated with HCV and alcohol.[28]

NAFLD is recognized as a major cause of cryptogenic cirrhosis, a condition in which histological features of IFL or NASH may have disappeared, and only thorough medical history taking can verify this relationship.[29,30] Cryptogenic cirrhosis in developed societies is linked to 30–40% of diagnosed HCC cases without other obvious risk factors.[11] Cryptogenic cirrhosis may be the cause of HCC diagnosed at more advanced stages, as demonstrated in a single-center study in which HCC was primarily identified by clinical symptoms, rather than via surveillance of patients with cryptogenic cirrhosis.[31]

**HCC emerging in non-cirrhotic NAFLD**

An additional concern is that HCC may complicate NAFLD with minimal or no fibrosis.[11,32] Analysis of a healthcare claims database accounting each year for 18 million American lives identified 4,406 cases of HCC between 2002 and 2008, with NAFLD being the most common underlying risk factor (59%) followed by diabetes (36%) and chronic HCV (22%).[33] Intriguingly, the majority of NAFLD-associated HCC cases (54%) had no International Classification of Diseases code for cirrhosis, in comparison with only 22% of HCV-associated HCC documented in the absence of cirrhosis.[34] In a compilation of case reports and case series from various geographical areas, the degree of fibrosis was F2 or lower in 28% of cases with NAFLD-associated HCC.[11] Finally, in a recent study of 17,895 patients with HCC identified from the SEER-Medicare database, non-cirrhotic NAFLD was the only etiology in 1,031 cases (5.8%), of which 186 cases (approximately 1% of the total HCC cohort) developed in isolated fatty liver.[34]

HCC risk remains very low when the analysis includes patients with all stages of NAFLD, not just those with biopsy-proven NASH. In a cohort study conducted on Japanese patients with echogram-based diagnosis of NAFLD, the annual incidence rate of HCC was 0.043%, amounting to a cumulative rate of 0.51% after a period of 12 years.[35] A systematic review found that NAFLD cohorts with few or no cases of cirrhosis had a cumulative HCC mortality of 0–3% for study periods of up to 20 years.[36] Based on large-scale epidemiological and long-term follow-up studies, HCC affects no more than 1 out of 3,000–4,000 individuals who have NAFLD in the absence of other liver disease.[12,13] With the growing prevalence of obesity and diabetes, however, these numbers may increase, and additional studies are warranted to predict and understand the societal impact of NAFLD on HCC.

**Pathogenesis of HCC associated with NAFLD**

**General considerations**

NAFLD has a complex pathogenesis, with cellular and molecular mechanisms providing multiple links to the development of liver cancer.[37,38] Most cases of NAFLD-associated HCC occur in cirrhosis, which provides a unique and strong tumorigenic microenvironment.[39] However, NAFLD usually develops in the context of obesity and diabetes, which are independently associated with increased risk for cancer in a variety of tissues, including the liver.[40,41] Therefore, interaction of oncogenic pathways related to adipose tissue dysfunction and established cirrhosis may provide a particularly conducive setting for the emergence of HCC (Fig. 2).

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**Fig. 1. Disease burden of HCC by major etiologies in the USA.** The number of individuals with chronic liver disease (prevalence shown by light colored squares) including those with cirrhosis (prevalence shown by dark color squares) who develop HCC in any given year (incidence shown by small pie chart in the center) is illustrated by areas corresponding to low estimates from data published for alcoholic liver disease,[11,12] NASH,[4,24,25] chronic hepatitis B,[13,14] and chronic HCV.[24,25,15,16] Accordingly, the highest number of cases of HCC are associated with chronic HCV, with the next highest number being associated with NAFLD. Note that the actual incidence of HCC is less than the sum of these estimates, indicating overlap between groups of individuals with chronic liver disease of various etiologies.
Molecular mechanisms of hepatocarcinogenesis in cirrhosis

There is substantial evidence that cirrhosis is one of the most important risk factors of HCC.39,42 Erratic liver remodeling in cirrhosis creates a microenvironment with repeated cycles of hepatocellular destruction and compensatory regeneration to foster the development of HCC.43,44 This complex process involves many different cell types and oncogenic pathways. Triggered by chronic hepatocellular or cholangiocellular injury, resident liver macrophages (Kupffer cells), hepatic stellate cells, sinusoidal endothelial cells, and intrahepatic lymphocytes gradually change their phenotypes and acquire pro-oncogenic properties. Along with the recruitment of additional players, these cells produce a variety of cytokines, chemokines, growth factors, free radicals, and other bioactive substances that drive the initiation and progression of HCC.45 Through stepwise changes that may take many years, emergence of HCC follows a dysplasia–carcinoma sequence seen in other cancers.43,44 Accordingly, identification of molecular signatures that distinguish dysplastic nodules from early HCC in cirrhotic liver may allow enhanced surveillance, reliable prognostication, and timely intervention.46

Essentially, all major mechanisms of oncogenesis have been implicated in the development of HCC, including structural defects of the genome, epigenetic silencing or activation, and overly active or aberrant signal transduction cascades.47,48 This extraordinary heterogeneity of HCC and the need to correlate molecular alterations with clinical characteristics have been the subject of several recent and excellent reviews.49,50 Initiation and progression of HCC involves numerous oncogenic pathways that may offer pharmacological targets for different subsets of patients.51,52 Thus, hepatocyte growth factor acting on c-met is critical for G1 to S transition in the cell cycle, making this receptor a candidate for targeted chemotherapy.53 Oncogenic signaling that depends on activation of Frizzled receptors by Wnt occurs through β-catenin, c-Jun N-terminal kinases (JNKs), and protein kinase C.54 Single nucleotide polymorphisms in the gene encoding epidermal growth factor, another major cell growth stimulator acting through cell surface receptors such as erbB1 and Her2/neu, have been linked to a variable risk of HCC.55 Recent research suggests that defects in the post-transcriptional regulation of gene expression via microRNA (miRNA) also contribute to development of HCC, and that different miRNA signatures may point to the underlying etiology.56,57

Adipose tissue remodeling and molecular mechanisms of hepatocarcinogenesis

So far, no structural or functional aberrations of liver cells have been specifically associated with the pathogenesis of HCC in NAFLD, although there are multiple candidates that may fulfill this role. Emergence of HCC in non-cirrhotic NAFLD suggests sufficient complexity to initiate hepatocarcinogenesis without the tumorigenic microenvironment of cirrhosis. Chronic nutrient excess results in expansion, redistribution, and remodeling of adipose tissue, which is associated with inflammation, lipotoxicity, and insulin resistance.58,59 These events are linked to numerous oncogenic pathways, and have a major role in the progression of NAFLD. Expansion of adipose tissue is characterized by disproportionate growth of visceral and deep subcutaneous fat depots, with an adverse adipokine secretory profile that includes higher levels of leptin and lower levels of adiponectin.60,61 Leptin...
has pro-inflammatory, pro-fibrogenic, and growth-promoting effects mediated by the Janus kinase/signal transducer and activator of transcription (JAK/STAT), phosphoinositide 3-kinase (PI3K)/Akt, and extracellular signal-regulated kinases (ERKs). By contrast, adiponectin, a potent activator of S'-adenosine monophosphate activated protein kinase (AMPK), has anti-inflammatory, anti-angiogenic, and tumor growth-limiting properties by opposing the actions of leptin and suppressing signal transduction through Akt, STAT3, and mammalian target of rapamycin (mTOR). Thus, coinciding leptin surplus and adiponectin deficiency may activate numerous oncogenic pathways.

Adipose tissue remodeling also results in increased secretion of the pro-inflammatory cytokines interleukin-6 (IL-6) and tumor necrosis factor alpha (TNF-α), which promote cell growth and inhibit apoptosis by targeting multiple signaling effectors such as ERKs, JNKs, nuclear factor-κB (NF-κB), signal transducer and activator of transcription 3 (STAT3), and mTOR. The role of these cytokines in the development of HCC is supported by observations that obesity-associated promotion of malignant liver lesions in mice treated with diethylnitrosamine requires intact TNF and IL-6 signaling.

**Oncogenic pathways associated with lipotoxicity**

When the capacity of expanding adipose tissue is exhausted, lipid molecules are stored in non-adipose tissues (e.g., liver, skeletal muscle, pancreas, and heart) and initiate lipotoxicity, defined as chronic cellular and tissue damage related to ectopic lipid accumulation. Liver lipotoxicity is advanced by de novo lipogenesis resulting from increased activation of steroidal response element binding protein-1c (SREBP-1c) caused by elevated insulin levels. Removal of excess hepatic lipids in NAFLD may be hampered by insufficient mitochondrial β-oxidation (a common cause of microvesicular steatosis) and impaired export of very low-density lipoproteins. Moreover, lysosome-mediated degradation of lipid molecules in intracellular autophagosomes (lipophagy) is inhibited by excess fatty acids, representing a self-amplifying mechanism of lipotoxicity in hepatocytes.

Experimental evidence indicates that lipotoxicity depends on the altered composition rather than the source or amount of ectopically deposited lipids. Thus, lipotoxicity has been attributed primarily to the harmful effect of free fatty acids and free cholesterol rather than triglycerides, which are considered relatively innocuous, and possibly actually protective. The panel of fatty acids implicated in lipotoxicity is broad, and tissue damage may depend on the length (short, medium, long, and very long), saturation (saturated, mono- and polyunsaturated), and isomerism (cis vs. trans) of acyl chains.

A recent lipidomic analysis by thin-layer chromatography of liver tissue obtained from patients with NAFLD indicated that free cholesterol content and the ratio of n-6 to n-3 polyunsaturated fatty acids were significantly higher in NASH compared with IFL, and diminished levels of phosphatidylcholine and arachidonic acid allowed further distinction of NASH. Recent reports indicate that magnetic resonance-based methods may eventually provide similar qualitative information on hepatic lipid content, although this approach is still in the experimental phase. If fully developed, these non-invasive methods hold the promise of predicting disease severity in NAFLD.

Lipotoxicity results in various adverse conditions such as mitochondrial dysfunction, oxidative injury, endoplasmic reticulum stress, and insulin resistance. These mechanisms provide multiple links between lipotoxicity and hepatocarcinogenesis. For example, hepatocellular death induced by lipotoxicity (lipoapoptosis) correlates with the severity of NAFLD, and implicates the effects of liver macrophages activated by damaged and dying hepatocytes. Enhanced lipid peroxidation may activate macrophages by generating ligands (e.g., oxidized LDL) of scavenger receptors. Stimulation of the pivotal danger recognition receptor, toll-like receptor 4, by saturated fatty acids is another mechanism of macrophage activation that may contribute to the liver inflammatory response. Importantly, macrophage-mediated stimulation of surviving hepatocytes via NF-κB and other cell-proliferation pathways is a major component of hepatocarcinogenesis, which has been established in various experimental models.

As a fundamental consequence of obesity-associated adipokine imbalance, chronic inflammation, and lipotoxicity, insulin resistance results from multiple defects of the insulin-mediated cellular signaling network, and leads to compensatory hyperinsulinemia. This process contains important self-amplification loops such as increased lipolysis from peripheral adipose depots due to uninhibited hormone-sensitive lipase, and increased hepatic de novo lipogenesis due to retained insulin responsiveness of SREBP-1c. Elevated insulin levels stimulate the production of insulin-like growth factor (IGF)-binding protein and increase bioavailability of IGF1 and IGF2, further promoting oncogenic pathways such as PI3K/Akt, mitogen-activated protein kinase, and vascular endothelial growth factor.

**Prevention of HCC associated with NAFLD**

**HCC reduction by weight management and physical exercise**

**HCC risk reduction by insulin-sensitizing agents**

Pharmacological therapy for the metabolic derangements associated with NAFLD such as insulin resistance and hyperlipidemia may provide additional opportunities to...
prevent hepatocarcinogenesis. There is evidence that insulin-sensitizing agents reduce the risk of HCC in NAFLD associated with manifest diabetes. Most of these data relate to the effect of metformin, although thiazolidinediones may carry similar benefit. By contrast, use of insulin and sulfonylureas has been associated with increased risk of HCC. Metformin has been reported to have tumor inhibitory properties in a variety of cancers, and this effect was confirmed in a xenograft model of human hepatoma cells.

The anti-proliferative action of metformin is linked to activation of AMPK, a master regulator of cellular energy metabolism, including inhibition of de novo lipogenesis, which has additional molecular targets such as the mTOR pathway and the retinoblastoma protein. A recent meta-analysis encompassing 105,495 patients with diabetes found that the risk of primary liver cancer was reduced by 62% (pooled odds ratio 0.38; 95% CI 0.24–0.59; p < 0.001) in those who were regularly taking metformin. It remains to be seen whether metformin has a similar chemopreventive effect on the risk of HCC in patients with NAFLD who have insulin resistance but no manifest diabetes.

The use of thiazolidinediones to improve insulin resistance has been associated with a reduction in risk for HCC of 44–70% in patients with diabetes. There are conflicting data on the molecular mechanisms by which these agents exert tumor prevention. Peroxisome proliferator active receptor γ (PPARγ), a nuclear hormone receptor and the main target of thiazolidinediones, has been shown to inhibit tumor growth and metastasis formation in HCC, although some findings suggest that the effect of thiazolidinediones on HCC is PPARγ-independent. Specific molecular mechanisms of tumor inhibition by thiazolidinediones include pro-apoptotic effects via activation of p53 and PTEN, as well as inhibition of Bcl2 and the ERK pathway. Because most members of this class of anti-diabetic drugs have been implicated in major adverse events, the role of thiazolidinediones in NAFLD management and HCC prevention remains controversial.

Additional opportunities for chemoprevention of HCC in NAFLD

Statins are inhibitors of endogenous cholesterol synthesis that work by targeting 3-hydroxy-3-methylglutaryl coenzyme-A (HMG-CoA) reductase, and are widely used for primary and secondary prevention of cardiovascular diseases. Statins have been reported to decrease the risk of a number of cancer types through anti-proliferative, pro-apoptotic, anti-angiogenic, and immunomodulatory mechanisms. Specific molecular targets of statins include G proteins of the Ras/Rho superfamily, the RAF/MEK/ERK pathway, and the Myc oncoprotein. In a recent meta-analysis of 4,298 cases of HCC, statin users were less likely to develop HCC than non-users, with an adjusted odds ratio of 0.63 (95% CI, 0.52–0.76).

Farnesoid X receptor (FXR) is a nuclear hormone receptor with high levels of expression in the liver, and has been implicated primarily in bile acid sensing. Similar to other nuclear hormone receptors, FXR regulates a large number of genes involved in lipid and glucose metabolism, liver regeneration, inflammation, and cancer. FXR-deficient mice display sustained activation of the Wnt/β-catenin pathway, and reduced expression of the tumor suppressor N-myc downstream-regulated gene 2 (NDRG2) with spontaneous development of HCC and other malignancies.

FXR inhibits gankyrin, a proteasome factor that promotes degradation of multiple tumor suppressor proteins. Human studies indicate that FXR is downregulated in HCC, further supporting a protective role for FXR in tumor development. Overexpression of FXR in human hepatoma cells or stimulation by synthetic agonists such as GW40024 inhibited tumor growth in an orthotopic mouse xenograft model. These findings provide a basis for novel approaches in the prevention and treatment of HCC.

Conclusions

HCC is one of the most common malignancies in the world and has one of the worst survival rates of all the major cancers. HCC is among the few malignancies with an increasing incidence in the USA, a worrisome trend initially attributed to the aging population with chronic HCV, but increasingly linked to the rapidly spreading epidemic of obesity, diabetes, and NAFLD. Although cirrhosis provides the most conducive microenvironment for hepatocarcinogenesis, NAFLD-associated HCC may emerge in its absence, fittingly described as “a wolf in sheep’s clothing.” Because NAFLD could be a game changer for liver cancer, it is imperative to develop reliable risk assessment and prevention tools, explore the pathogenesis in non-cirrhotic HCC, and identify new molecular targets for effective therapy.

Conflict of interest

None

Author contributions

Conceiving and writing review (GB).

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