Adipose Tissue Dysfunction and Alcoholic Liver Disease

Abstract
Alcoholic liver disease (ALD) remains an important health problem in the United States. The disease process is featured by early steatosis, steatohepatitis (steatosis with inflammatory cells infiltration and necrosis), with some individuals ultimately progressing to fibrosis/cirrhosis. Although the disease progression is well characterized, no FDA-approved therapies are currently available to halt or reverse this process in humans. The mechanisms underlying the initiation and progression of ALD are complex and multifactorial. Emerging evidence supports that adipose tissue dysfunction contributes to the pathogenesis of ALD. It has been long known that aberrant hepatic methionine metabolism is a major metabolic abnormality induced by chronic alcohol exposure and plays an etiological role in the pathogenesis of ALD. The recent studies in our group documented the similar effect of chronic alcohol drinking in adipose tissue. In this article, we briefly reviewed the recent research progress in the field with a focus on how abnormal methionine metabolism in adipose tissue contributes to adipose tissue dysfunction and liver damage.

Keywords: Alcohol, Liver, Adipose, Methionine, Lipolysis, Methylation

Alcoholic Liver Disease (ALD)
Although much progress has been made over last three decades of research on the mechanisms underlying alcoholic liver disease (ALD), it remains an important health problem in the United States. ALD ranks among the major causes of morbidity and mortality in the world, and affects millions of patients worldwide each year [1]. The disease process is characterized by early steatosis, steatohepatitis (steatosis with inflammatory cells infiltration and necrosis), with some individuals ultimately progressing to fibrosis/cirrhosis. Although the disease progression is well characterized, there is currently no FDA-approved therapy available to halt or reverse this process in humans.

Pathogenesis of Early-Stage ALD
Hepatic steatosis, characterized by excess fat accumulation in hepatocytes, is the most common and earliest response of the liver to chronic alcohol consumption. Although “pure” steatosis is clinically considered to be a benign condition, excessive fat accumulation makes hepatocytes vulnerable to the attack of “the second hit,” such as pro inflammatory cytokines and oxidative stress, leading to the progression to steatohepatitis [2,3]. The mechanisms involved in the development of alcohol-induced hepatic steatosis are multifactorial. Sterol regulatory element binding proteins (SREBP)-1c, a master transcription factor controlling de novo lipogenesis, is up regulated in the liver of mice chronically exposed to ethanol-containing diet [4]. Importantly, liver-specific knockout of SREBP-1c protect mice against alcohol-induced fatty liver and liver damage [5], supporting the notion that enhanced hepatic de novo lipogenic process plays a pivotal role in alcohol-triggered fat accumulation in the liver. Moreover, chronic alcohol exposure is associated with impaired fatty acid β-oxidation in hepatocytes, contributing to fat accumulation in hepatocytes. Suppressions of both adenosine monophosphate-activated protein kinase (AMPK) and peroxisome proliferator-activated receptor-alpha (PPAR-alpha), two regulatory proteins of fatty acids oxidation, are mechanistically involved in this process [6,7]. Furthermore, long-term alcohol consumption is reported to enhance uptakes of free fatty acids and triglyceride-rich lipoproteins by hepatocytes [8,9] and impair hepatic very-low-density lipoprotein (VLDL) secretion [10], thereby contributing to fatty liver after chronic alcohol exposure.

Adipose Tissue Dysfunction and ALD
Although it has been well-established that chronic alcohol consumption exerts a detrimental effect on hepatic fat synthesis and disposal, leading to the development of hepatic steatosis, emerging evidence supports that adipose tissue dysfunction also plays an important role in the pathogenesis of ALD. In the clinical setting, it has been reported that visceral fat accumulation is positively related to the onset of alcoholic liver damage and body mass index (BMI) represents an independent risk factor for fibrosis in alcoholic patients [11-14]. Moreover, adipose tissue inflammation is correlated with the severity of pathological features in the liver of patients with ALD [13]. Experimentally, long-term alcohol consumption is associated with adipose tissue oxidative stress, insulin resistance, inflammation, adipocyte cell death, and adiponectin decline [15-19]. Chronic alcohol feeding results in hyper-lipolysis (degradation of triglycerides) in adipose tissue, leading to elevated circulation free fatty acids concentrations and a significant loss of white adipose tissue [8]. A recent study demonstrated that moderate obesity and alcohol synergistically induced steatohepatitis [20], further supporting the critical role of adipose tissue dysfunction in the development of ALD. Importantly, both rosiglitazone [a
PPAR-gamma agonist mainly targeting adipocytes) [21] and recombinant adiponectin (an adipokine exclusively secreted by adipocytes) [22] improved ALD, suggesting that improving adipose tissue function represents a potential therapeutic approach for ALD.

**Pathogenesis of Early-Stage ALD**

**Methionine metabolism abnormality in ALD**

Aberrant hepatic methionine metabolism is a major metabolic abnormality induced by chronic alcohol exposure and plays an etiological role in the pathogenesis of ALD [23-26]. As illustrated in (Figure 1), intracellular methionine metabolism involves two major pathways, transmethylation reaction and transsulfuration reaction. The first step in methionine metabolism is the formation of S-adenosyl methionine (SAM) in a reaction catalyzed by methionine adenosyl-transferase (MAT). Under physiological conditions, most of the SAM generated per day is used in transmethylation reactions in which methyl groups are added to a vast number of molecules, including DNA, RNA, phospholipids, histones, and other proteins, via specific methyltransferases. In this process, SAM is converted to S-adenosylhomocysteine (SAH), followed by homocysteine (Hcy) and cysteine, a precursor for glutathione biosynthesis, via transsulfuration pathway. SAH is a potent competitive inhibitor of most methyltransferases studied and decreased SAM: SAH ratio has been widely employed as an indicator of suppressed transmethylation reactions [27,28]. While chronic alcohol exposure leads to hepatic SAM deficiency, both SAH and Hcy are increased in the liver in response to alcohol.

**Aberrant adipose methionine metabolism and hyper-lipolysis in adipose tissue**

Cellular/molecular mechanisms underlying alcohol-triggered adipocyte dysfunction are not completely known. Increased oxidative stress due to CYP2E1 over expression in adipose tissue represented one of the mechanisms [19]. Studies using both genetic and dietary animal models provided evidence supporting that hyper homocysteinemia (HHcy) is associated with adipose tissue dysfunction [29-32], suggesting that methionine metabolism regulates adipose tissue function. We are the first to report that, similar to its effect on the liver, chronic alcohol feeding induces methionine metabolism abnormality in adipose tissue, which is characterized by SAM deficiency, and Hcy and SAH accumulation, leading to significant decrease of SAM/SAH ratio, a strong indicator of inhibitory transmethylation reactions (hypomethylation). We subsequently showed that Hcy accumulation in adipose tissue contributes to adiponectin decline in ALD via a mechanism involving ER stress induction [16] and adipogenesis suppression via inhibiting PPAR-gamma transactivation [33]. Further, we provided evidence supporting that intracellular hypomethylation status in adipocytes in the setting of chronic alcohol feeding contributes to adipose tissue hyper-lipolytic response in ALD via suppressing protein phosphatase 2A (PP2A) activity, leading to hormone sensitive lipase over activation [34] (Figure 2). Our data support that rectification of methionine metabolism through dietary supplementation of betaine protects against alcohol-induced liver damage, at least partially via improving adipose tissue function. Taken together, the recent research in our group suggests that aberrant methionine metabolism in adipocytes contributes to alcohol-elicited adipose tissue dysfunction and liver damage.

**Figure 1:** Intracellular methionine metabolism: Chronic alcohol consumption causes SAM deficiency, but enhancement of homocysteine and SAH.


**Figure 2:** SAM-dependent methylation reactions are required for PP2A activation, which dephosphorylates (inhibits) HSL. Chronic alcohol consumption induces intracellular hypomethylation status in adipocytes, which suppresses PP2A activity, leading to uncontrolled HSL activation.

SAM: S-Adenosylmethionine; Me: Methyl Group; PP2A: Protein Phosphatase 2A; HSL: Hormone Sensitive Lipase.
Conclusion

Emerging evidence shows that adipose tissue plays an important role in both initiation and progression of liver damage induced by chronic alcohol consumption. The recent studies in our laboratory provide initial evidence suggesting that methionine metabolism abnormality contributes to alcohol-associated adipose tissue malfunction. Although much progress has been made, the exact underlying cellular/molecular mechanisms involved in adipose tissue dysfunction in ALD remain to be elucidated. Drugs targeting both adipose tissue and the liver represent ideal choices for the treatment of ALD.

References


