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Fat: A matter of disturbance for the immune system

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Abstract

Obesity is increasingly being recognized as a risk factor for a number of benign and malignant gastrointestinal conditions. However, literature on the underlying pathophysiological mechanisms is sparse and ambiguous. There is compelling evidence that both overnutrition and undernutrition negatively interfere with the immune system. Overnutrition has been found to increase susceptibility to the development of inflammatory diseases, autoimmune diseases and cancer. In the regulation of immune and inflammatory processes, white adipose tissue plays a critical role, not only as an energy store but also as an important endocrine organ. The obese state is characterised by a low-grade systemic inflammation, mainly as a result of increased adipocytes as well as fat resident- and recruited-macrophage activity. In the past few years, various products of adipose tissue including adipokines and cytokines have been characterised and a number of path-

ways linking adipose tissue metabolism with the immune system have been identified. Activation of the innate immune system plays a major role in hepatic steatosis. Non-alcoholic fatty liver disease includes a wide spectrum of diseases, from pure steatosis to non-alcoholic steatohepatitis in the absence of significant alcohol consumption. Although steatosis is considered a non-progressive disease, non-alcoholic steatohepatitis may deteriorate in advanced chronic liver diseases, cirrhosis, and hepatocellular carcinoma. An important parallel between obesity-related pathology of adipose tissue and liver pertains to the emerging role of macrophages, and growing evidence suggests that Kupffer cells critically contribute to progression of non-alcoholic fatty liver disease. Moreover, a close link between specific immune activation and atherosclerosis has been well established, suggesting that fat can directly trigger immune responses. This review discusses the role of fat as "a matter of disturbance for the immune system" with a focus on hepatic steatosis.

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Key words: Adipocytokine; Adipose tissue; Fat; Immune system; Kupffer cell; Natural killer; Steatosis

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INTRODUCTION

Obesity predisposes individuals to an increased risk of

developing many diseases, including atherosclerosis, diabetes, non-alcoholic fatty liver disease (NAFLD), cancer and immune-mediated disorders, such as asthma^[1-3]. Obesity is typically assessed clinically with the surrogate measure of body mass index (BMI). Individuals with a BMI ≥ 30 kg/m² are considered obese. The incidence of obesity and its associated disorders is increasing markedly worldwide. Data from the most recent NHANES (National Health and Nutrition Examination Survey; 2005-2006) indicate that the prevalence of obesity was 33%-35% among US adults^[4]. In another recent NHANES survey based on the combined years of 2003-2006, 16% of children or adolescents aged 2-19 years were obese^[5]. In Europe, several surveys conducted since 2000 and using direct anthropometric measurements, showed that the prevalence of obesity ranges from 15% to 30% in men and from 11% to 34% in women, with considerable geographic variation (rates being higher in Central, Eastern, and Southern Europe)^[6]. Urbanization and unbalanced diet, associated with genetic susceptibility have allowed the emergence of the obese phenotype.

In mammals, adipose tissue (AT) occurs in two forms: white adipose tissue (WAT) and brown adipose tissue (BAT). Most AT in mammals is WAT and this is thought to be the site of energy storage. In contrast, BAT is found mainly in human neonates and is important for the regulation of body temperature through non-shivering thermogenesis. In addition to adipocytes, which are the most abundant cell type in WAT, adipose tissue also contains pre-adipocytes or stromal vascular cells (which are non-fat cells): endothelial cells, fibroblasts, leukocytes and, most importantly, macrophages. Body fat distribution, rather than adiposity *per se*, is an important risk factor for obesity-related disorders. An excess of intra-abdominal fat rather than subcutaneous fat (central *vs* peripheral obesity) is associated with metabolic syndrome (MS) and cardiovascular disease (CVD). The mechanisms responsible for this association are still unknown, but several hypotheses, which are not mutually exclusive, have been formulated^[7]. The first hypothesis proposed a direct effect of visceral AT depots on insulin resistance, lipoprotein metabolism, and blood pressure. Metabolic products of omental and mesenteric AT depots are released into the portal vein, which provides direct delivery to the liver. Lipolysis of omental and mesenteric AT depots releases free fatty acids (FFAs) that can induce hepatic insulin resistance and provide substrate for lipoprotein synthesis and neutral lipid storage in hepatocytes. In addition, specific proteins and hormones produced by omental and mesenteric AT, such as inflammatory molecules, angiotensinogen, and cortisol can also contribute to MS and CVD. Another hypothesis suggests that the limited capacity of subcutaneous fat to store excess energy results in overflow of fatty acids to intra-abdominal fat and "ectopic" sites such as liver, muscle, and islets. In this paradigm, excess intra-abdominal fat is merely a marker of fatty acid overflow from subcutaneous depots.

whose sole function was the storage of fat. However, it is now recognized that AT is an active endocrine organ that secretes numerous adipokines, cytokines and chemokines including leptin, adiponectin, resistin, retinol binding protein 4 (RBP4), tumor necrosis factor α (TNF- α), interleukin (IL)-1 β , IL-6, and monocyte chemoattractant protein 1 (MCP-1)^[8,9]. All of these play a central role in the regulation of energy and vascular as well as immune system homeostasis by acting both locally and at distant sites influencing various metabolic and immune processes. Moreover, organs other than AT may contribute to systemic levels of some adipokines.

Obesity is associated with a low-grade inflammation of WAT resulting from chronic activation of the innate immune system, which can subsequently lead to insulin resistance, impaired glucose tolerance and even diabetes^[10,11]. In addition to these associations between obesity and disease, research in the past few years has identified important pathways that link metabolism with the immune system and *vice versa*. Many of these interactions between metabolic and immune systems seem to be orchestrated by the complex network of soluble mediators derived from immune cells and adipocytes^[1].

The effects of obesity on the immune system are not restricted to local effects within AT. Elevated levels of pro-inflammatory cytokines have been noted in the serum of asymptomatic obese individuals, the cytokine levels being related to the degree of obesity^[12]. TNF- α is only present at very low levels in human blood suggesting that TNF- α released by adipose tissue has only autocrine/paracrine actions. IL-6, however, is present at much higher levels. Adipocyte-derived IL-6 has been estimated to comprise 30% of the circulating IL-6 suggesting an endocrine action^[13]. Furthermore, these elevated levels of IL-6 are associated with increased circulating levels of C-reactive protein suggesting that although the elevation in levels is modest compared with those seen in sepsis, they could be having real effects on innate immune function.

Obesity is also associated with altered functioning of circulating immune cells^[12,14]. Decreased T- and B-cell function, increased monocyte and granulocyte phagocytosis and oxidative burst, and an increase in leukocyte count have been described. More recently, circulating mononuclear cells from obese subjects have been shown to exhibit increased nuclear factor κ B (NF κ B) nuclear binding with decreased levels of NF κ B inhibitor, together with increased mRNA expression of IL-6, TNF- α and migration inhibition factor. Furthermore, there is a good correlation between the markers of macrophage activation and plasma levels of FFAs^[15]. It has previously been demonstrated that macronutrient challenges in normal subjects increase NF κ B nuclear binding in circulating mononuclear cells, raising the possibility that the activated state of mononuclear cells is due to increased circulating levels of FFAs found in the obese. Indeed, hyperlipidaemia in mice mediates an inflammatory response by the same signalling cascade through which lipopolysaccharide activates the innate immune system (this engages a receptor complex comprising Toll 4 CD14, CD14 and MD-2)^[16]. Table 1

FAT AND THE IMMUNE SYSTEM

Adipose tissue was once thought to be an inert mass

Table 1 Adipocytokines, pro-inflammatory cytokines and chemokines, and other factors synthesised by adipocytes and macrophages in white adipose tissue

Adipocytes	Macrophages
Adiponectin	TNF- α
Leptin	IL-1 β
Resistin	IL-6
RBP4	MCP-1
TNF- α	Resistin
IL-1 β	
IL-6	
MCP-1	
Visfatin	
MIP	

RBP4: Retinol binding protein 4; TNF- α : Tumor necrosis factor α ; IL: Interleukin; MCP: Monocyte chemotactic protein; MIP: Macrophage inflammatory protein.

summarises the secretion of adipokines, cytokines and other factors by adipocytes and macrophages in WAT. Finally, recent research has implicated the innate immune system in the pathophysiology of obesity-related liver damage^[17,18].

Obesity is a high risk factor for NAFLD. Studies in an animal model of obesity-related liver disease revealed the involvement of dysfunctional hepatic immune cells^[19]. In this review we analyse the relationship between hepatic steatosis and the immune system.

HEPATIC STEATOSIS AND THE IMMUNE SYSTEM

Hepatic steatosis is the histological hallmark of alcoholic liver disease (ALD) and NAFLD, which are among the commonest causes of cirrhosis and liver failure in the developed world^[20-24]. Steatosis may also alter the natural history of other liver diseases such as chronic viral hepatitis^[25]. Excessive consumption of alcohol in humans results in a spectrum of liver abnormalities, ranging from simple fatty liver to steatohepatitis and cirrhosis, which may be present independently or in combination. Infiltration of the liver by lymphocytes and neutrophils is an important feature of alcoholic hepatitis; it initiates a cascade of effector mechanisms that ultimately lead to hepatocyte death, fibrosis, and cirrhosis. Only a minority of consistently heavy drinkers with steatosis ever develop clinically important liver disease^[24,26] implying that host or environmental factors determine the evolution of alcohol-related liver damage. Ingestion of alcohol leads to increased production of reactive oxygen species (ROS), which are generated during the metabolism of alcohol by cytochrome P450 2E1 enzyme, and excessive alcohol consumption is associated with increases in lipid, protein, and DNA peroxidation. Consistent with this disease model, risk factors for the development of progressive liver damage in alcohol drinkers include both polymorphisms in alcohol-metabolizing enzymes and polymorphisms in genes associated with a more vigorous inflammatory response in

addition to exogenous factors including obesity, exposure to other hepatotoxins, and infection with hepatitis C and/or B virus^[27-29].

NAFLD is increasingly recognized as a leading cause of liver dysfunction and cirrhosis in the developed world and is part of a spectrum of metabolic diseases associated with central (intra-abdominal) obesity, hypertension, dyslipidaemia, insulin resistance, and type 2 diabetes mellitus^[22,30]. Similar to alcoholic liver disease, NAFLD is a spectrum of disorders, beginning as simple steatosis that is mostly considered an innocent condition. Being both the source and the result of insulin resistance, however, steatosis may be associated with an increased risk for cardiovascular morbidity^[31]. Most importantly, in about 15% of all patients with NAFLD, steatosis may evolve into steatohepatitis (NASH), a medley of inflammation, hepatocellular injury, and fibrosis, often resulting in cirrhosis and even hepatocellular carcinoma^[32]. Although this full sequence of progression is relatively rare, the overwhelming prevalence of NAFLD predicts a major healthcare burden. Epidemiology, pathogenesis, and approach to treatment of NAFLD follow the same trends as other metabolic disorders, and insulin resistance is the key event linking NAFLD to these diseases^[33-35].

ROLE OF ADIPOCYTOKINES IN ALCOHOLIC AND NON-ALCOHOLIC STEATOHEPATITIS

Many of the initial proinflammatory changes seen in NAFLD may be the consequence of altered metabolism rather than the underlying immune pathogenic event, and adipokines provide a link between fat, inflammation, and immunity (for more details see review by Tilg *et al*^[9]). More than 50 adipokines have been identified so far. Of these, leptin and adiponectin can influence the immune response, and their serum levels are increased and decreased, respectively, in NASH^[9]. While many adipokines are associated with adverse biological functions, adiponectin, the most abundant adipose-derived hormone, seems to have a protective effect in NAFLD. Adiponectin inhibits TNF- α induced endothelial cell adhesion molecule expression, induces production of anti-inflammatory cytokines such as IL-10, and reduces T and B lymphocyte responses. In particular, full-length adiponectin (Acrp30) and its cleavage derivative, globular adiponectin (gAcrp), have been credited with anti-diabetic, anti-inflammatory and anti-atherogenic properties^[36]. Adiponectin stimulates hepatic fatty acid oxidation and ketogenesis, while it inhibits cholesterol and triglyceride synthesis^[36]. While these metabolic activities primarily occur in hepatocytes, adiponectin has potent anti-inflammatory effects in macrophages. Thus, adiponectin is able to suppress the effects of lipopolysaccharides (LPS) in macrophages, including activation of NF- κ B and ERK1/2^[37-39]. Similarly, adiponectin prevents LPS-mediated inflammatory signalling in Kupffer cells^[40]. These anti-inflammatory effects of adiponectin may involve IL-10 signalling pathways^[41]. Interestingly, NADPH oxidase is a

major IL-10 target in various cell systems including macrophages^[42].

Decreased levels of adiponectin are definitely related to a variety of unfavourable effects, but the precise origin of adiponectin reduction has not been clarified. TNF- α has been demonstrated to suppress the transcription of adiponectin in an adipocyte cell line, which might explain the lower levels of serum adiponectin in obese individuals^[9]. Expression of adiponectin is also regulated by other pro-inflammatory mediators such as IL-6, which suppresses adiponectin transcription and translation in an adipocyte cell line^[9].

In a recent study, Kolak *et al.*^[43] evaluated subcutaneous AT biopsies obtained from healthy women both with and without increased liver fat (LFAT) ($2.3\% \pm 0.3\%$ *vs* $14.4\% \pm 2.9\%$, respectively), with similar BMIs and percentage body fat. Expression of cytokines and chemokines including CD68 (which correlates with the number of macrophages), MCP-1, macrophage-inflammatory protein (MIP-1 α), and PAI-1 were significantly increased, whereas peroxisome proliferator-activated receptors (PPAR)- γ and adiponectin were significantly decreased in women with high levels of LFAT compared with women with normal levels of LFAT, even though subcutaneous fat cell size, BMI, and percentage body fat were similar.

Leptin activates neutrophils, stimulates proliferation in human circulating monocytes, and appears to induce Th1-type cytokine production while inhibiting Th2-type cytokines. In addition, leptin has marked effects on the innate immune response by promoting activation and phagocytosis of macrophages, presumably through JAK/STAT signalling^[44]. Expansion of adipocytes in obesity leads to the recruitment of macrophages and the release of TNF- α , IL-6, and MCP-1 from macrophages and lymphocytes. TNF- α and IL-6 suppress the transcription of adiponectin, and TNF- α and IL-1 stimulate the production of leptin^[9,44]. Accordingly, hyperleptinaemia associated with obesity may contribute to progression of NAFLD, although this issue remains controversial^[45].

Resistin is another pro-inflammatory adipokine secreted by monocytes/macrophages and adipocytes in response to pro-inflammatory signals. Resistin induces NF κ B-dependent secretion of TNF- α and IL-6 by monocytes and increases ICAM-1 and VCAM-1 expression in endothelial cells, suggesting that it contributes to endothelial activation and leukocyte recruitment^[46]. In particular, in pure steatosis there is no significant increase in adhesion molecule expression but distinctive patterns are associated with both alcoholic hepatitis and cirrhosis, and in murine models of NASH elevated ICAM-1 expression is seen^[47]. Alcoholic hepatitis is characterized by increased expression of E-selectin and ICAM-1 on portal and hepatic venous endothelium and of ICAM-1, VCAM-1, and VAP-1 on sinusoidal endothelium as a consequence of local pro-inflammatory cytokines, particularly TNF- α ^[48-51]. In alcoholic cirrhosis, increased expression of endothelial adhesion molecules including ICAM-1, VCAM-1, and P-selectin is largely restricted to portal and septal vessels.

Endothelial ICAM-1 expression is increased in periseptal areas where LFA-1 is also increased in leukocytes, however, in contrast to alcoholic hepatitis, there is little increased ICAM-1 expression on hepatocytes^[48].

Visfatin, the characteristic adipokine of mesenteric AT, was previously identified as a protein involved in immune B-cell maturation (pre-B colony enhancing factor)^[52]. More recently, visfatin was described to be a highly expressed protein with insulin-like functions that was predominantly found in visceral AT, from which the name visfatin was derived^[53]. Thus, visfatin was identified as nicotinamide phosphoribosyltransferase, the rate-limiting enzyme that converts nicotinamide (a form of vitamin B3) to nicotinamide mononucleotide, a NAD precursor^[54]. Visfatin also has pro-inflammatory properties by inducing TNF- α and IL-6 in monocytes^[55]. Further studies are needed to fully understand the effect of this adipokine in Kupffer cells.

Figure 1 summarises the effects of adipocytokines on the regulation of the immune response.

HEPATIC STEATOSIS AND NATURAL KILLER CELLS

One experimental model which has generated a significant body of evidence regarding potential mechanisms of NAFLD pathogenesis and its relationship with the immune system is the *ob/ob* mouse. *Ob/ob* mice, which are leptin deficient as a result of a spontaneous mutation in the leptin gene, exhibit a number of metabolic and inflammatory features which mimic human NAFLD^[56] including insulin resistance, hyperlipidaemia, hepatic steatosis, and TNF- α elevation. One of the principal applications of the *ob/ob* mouse has been the identification of susceptibility of the steatotic liver to inflammatory insult (exemplified by the response to LPS) as a key factor in the development of NASH^[57]. A number of immuno-regulatory abnormalities have been identified in *ob/ob* mice which may contribute to their increased susceptibility to inflammatory damage. These include selective depletion in the liver (but not other organs) of Natural Killer (NK) T cells, a key population of immuno-regulatory/effecter lymphocytes which express phenotypic features of both "classical" T cells (CD3) and NK cells [NK1.1 (CD161 in humans)]^[58,59]. In their most characteristic form, NKT cells show specificity, through a semi-invariant surface T-cell receptor, for highly conserved glycolipid antigens presented by the MHC class I homolog CD1d. NKT cells, which are specifically enriched within the liver, have characteristic cytokine release patterns {Th-1 dominant [interferon (IFN)- γ], mixed, and Th-2 dominant (IL-4) depending on the mechanism of stimulation} which endow, in addition to their effector function, significant immuno-regulatory properties^[60]. The observation that liver NKT cells are depleted in steatosis in *ob/ob* mice has led to the suggestion that these cells play a key role in mediating and/or regulating inflammatory effects critical to the development of NAFLD. Although of potential value in the understanding of the pathogenesis of NAFLD, conceptual problems arise with regard to the *ob/ob* mouse

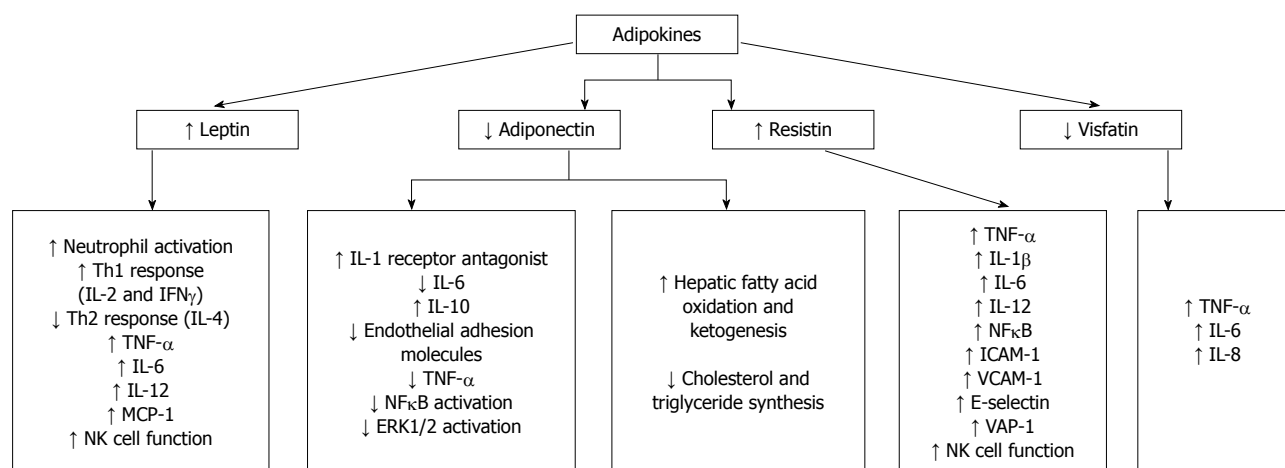


Figure 1 Effects of adipocytokines on regulation of the immune response. IL: Interleukin; IFN: Interferon; MCP: Monocyte chemotactic protein; NK: Natural Killer; TNF- α : Tumor necrosis factor α ; NF κ B: Nuclear factor κ B.

as a model for human disease due to its markedly different leptin phenotype (absent *vs* elevated), and to the fact that leptin is itself a key immunomodulatory cytokine^[61]. There are, therefore, potential mechanisms whereby leptin deficiency could modulate the immune response independent of its effects on hepatic fat accumulation. Li *et al.*^[62] used a natural obese/steatosis model to study the effects of hepatic steatosis on hepatic innate immune system function in leptin complete animals. C57Bl/6 mice fed a high-fat diet showed excess weight gain and the development of hepatic steatosis^[63]. Although total hepatic mononuclear cell levels were similar in the high- and low-fat diet groups, the percentage of hepatic (but not splenic) NKT cells was significantly reduced. Within both the hepatic T-cell and NKT fractions, the numbers of cells showing cytoplasmic staining for TNF- α and IFN- γ were, conversely, increased in the high-fat diet group (and serum IFN- γ levels were elevated), suggesting Th1 skewing of the response phenotype resulting from induced NKT cell effects. Finally, the livers of obese mice appeared to be sensitized to LPS injury, presumably reflecting the augmented Th1-type inflammatory cytokine response. These observations suggest that the development of hepatic steatosis *per se* can be associated with significant changes in liver NKT cell function. This finding would be compatible with the NKT cell changes seen in the *ob/ob* mice occurring as a result of hepatic steatosis that occurs in these animals, rather than the specific absence of leptin. The findings do, however, raise a number of issues which will determine whether this model is suitable for the study of human NAFLD. The first issue is the mechanism responsible for liver NKT cell “loss”, and Th-1 skewing of the residual cells, in obese C57Bl/6 mice. Theoretically, a reduction in liver NKT cells in obese C57Bl/6 mice could result from a decreased rate of NKT cell recruitment to, or development in, the liver, an increased rate of NKT cell death or migration from the liver, a loss of surface markers identifying the cells as NKT cells or any combination of these effects. The liver recruitment aspect of NKT cell homeostasis was not addressed

in the Li’s study^[60,64,65]. Instead, the authors argue that increased cell loss is the dominant effect, with evidence presented to suggest increased NKT cell apoptosis and increased hepatic expression of IL-12 (postulated to be a promoter of NKT cell apoptosis). There is an emerging consensus, however, that NKT cells are in fact relatively resistant to activation-induced cell death^[66]. An alternative (albeit non-mutually exclusive) explanation for the Li’s data would be that endogenous IL-12 released by Kupffer cells (KC) at elevated levels in the context of obesity^[62,67] acts as a cofactor for the stimulation of IFN- γ release (as opposed to IL-4 release which occurs in the absence of IL-12) by physiologically activated NKT cells, with the resulting “loss” of cells occurring as a consequence of post-activation surface phenotypic shift^[68]. If elevation of KC-released IL-12 in response to steatosis were to prove to be a factor in human fatty liver development^[69], its well-established ability to promote breakdown of self-tolerance may explain the increasingly recognised tendency towards autoantibody formation reported in NASH patients^[70,71]. The possibility that NKT cell activation is responsible, through activation-induced cell death and/or post-activation phenotypic change, for “reduction” in hepatic NKT cells in obese C57Bl/6 mice, and through cytokine release, for liver damage, raises the important question of the mechanism of this activation. Most previous work on NKT cell activation has used non-physiological ligands (anti-CD3 and anti-TCR). Although the recent identification of α -galactosylceramide has highlighted the potential importance of glycolipids as natural ligands for NKT, it is unlikely, given its marine sponge origin, that this agent is a physiological ligand in mice or humans. At present, the identity of the *in vivo* physiological ligand for NKT cells, the extent to which TCR-mediated as opposed to cytokine-driven mechanisms (such as *via* IL-12) are required for activation, and the extent to which different activation pathways result in different cytokine response phenotypes, remain areas of speculation. One potentially highly intriguing link between hepatic steatosis and NKT cell activation has emerged

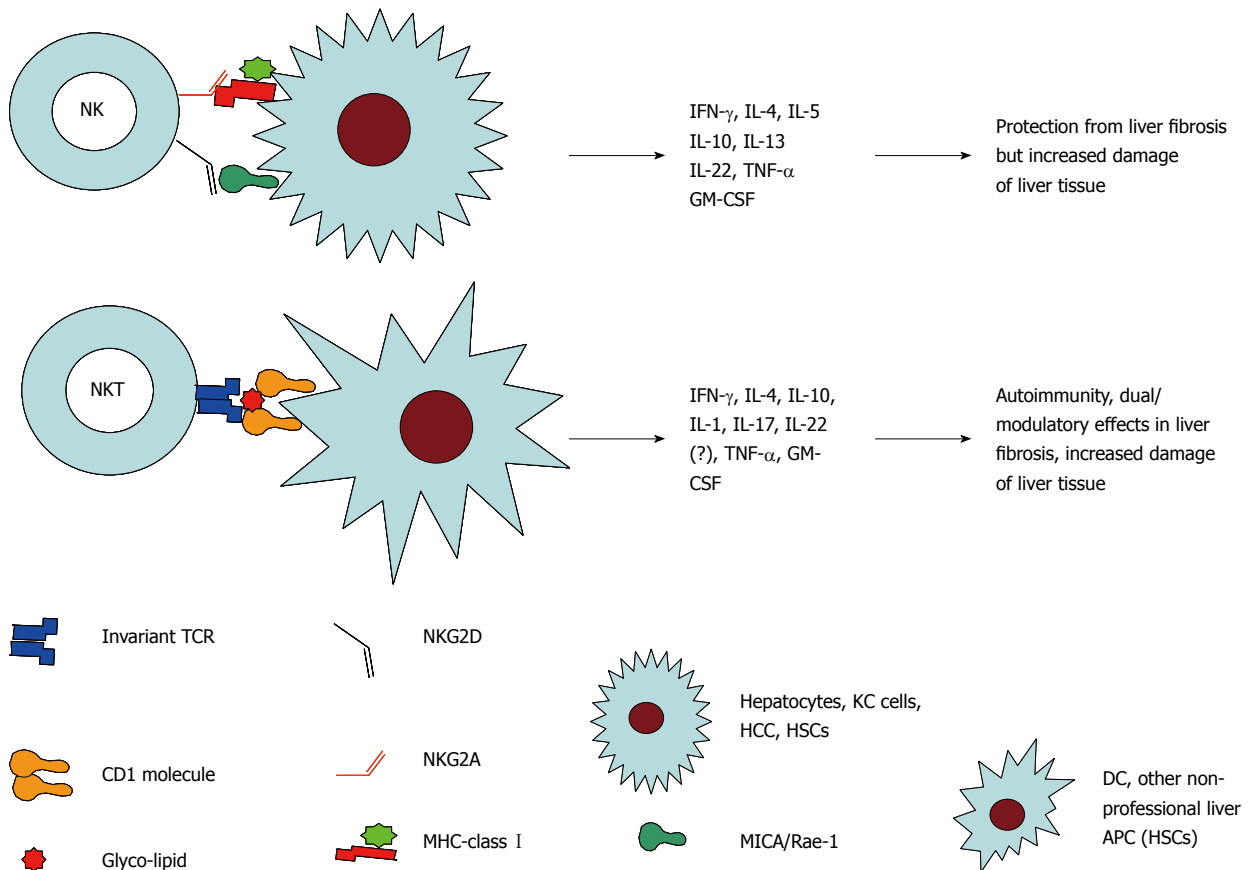


Figure 2 Simplified scheme of Natural Killer/Natural Killer T cell role in liver diseases. Natural Killer (NK) interacts with major and minor histocompatibility antigens expressed on several liver cells and kill and/or produce cytokines having several effects on the tissue. More complex is the role of NKT cells. These cells specifically recognize an antigen expressed in the context of a CD1 molecule and, upon recognition through an invariant TCR, secrete a large amount of cytokines having pleiotropic, sometimes controversial effects, whose overall results are due to the cytokine milieu and to the conditioning of the functions of other immune cells. This scenario is further complicated by the fact that many soluble factors (for instance cytokines) and hedgehog ligands may activate NK or NKT. IL: Interleukin; TNF- α : Tumor necrosis factor α ; HCC: Hepatocellular carcinoma; HSCs: Hepatic stellate cells; GM-CSF: Granulocyte-macrophage colony stimulating factor; APC: Antigen presenting cell.

with the observation that microsomal triglyceride transfer protein, plays a key role in the acquisition of glycolipid antigens by CD1d^[72]. In partial support of this concept, deficiency of microsomal triglyceride transfer protein in mice is associated with hepatic steatosis, and functional polymorphisms of its encoding gene have shown significant associations with NASH in humans^[73]. One approach to dissecting out the mechanisms of NKT cell activation and loss in obese C57Bl/6 mice would be to utilize NKT cell adoptive transfer and tracking methodologies in recombinant NKT cell-deficient mice in combination with NKT cell activation and appropriate cytokine blocking.

In a recent paper, Hua *et al.*^[74] examined the mechanism of dietary fatty acid induced hepatic NKT cell deficiency and its causal relationship to insulin resistance and NAFLD, and found that dietary saturated fatty acids (SFA) or monounsaturated fatty acids (MUFA), but not polyunsaturated fatty acids (PUFA), caused hepatic NKT cell depletion with increased apoptosis. Dietary SFA or MUFA also impair hepatocyte presentation of endogenous, but not exogenous, antigen to NKT cells, indicating alterations of the endogenous antigen processing or presenting pathway. *In vitro* treatment of normal hepatocytes with fatty acids also demonstrates impaired ability of CD1d to present

endogenous antigen by dietary fatty acids. Furthermore, dietary SFA and MUFA activate the NF κ B signaling pathway and lead to insulin resistance and hepatic steatosis.

Recently, a new subset of T helper cells, named Th17 due to the ability to produce IL-17 and other cytokines, has been correlated to processes underlying hepatic steatosis. In particular, Th17 largely express a NKT marker, CD161, and they have been described to be closely involved in the immune responses in several anatomical sites including skin, liver and gut^[75,76]. Th17 produce cytokines besides IL-17 such as IL-22 which is indicated to play a pivotal role in hepatic steatosis as recently shown^[77].

Figure 2 shows a simplified Scheme of NK/NKT cell role in liver diseases.

KUPFFER CELLS AND STEATOSIS

Hepatocellular accumulation of lipids is a key morphologic feature of NAFLD. Lipidomic analysis of human liver tissue is a promising novel approach to associate abnormal fat composition with various stages of NAFLD. Thus, total and damaged phospholipids are more abundant in simple steatosis at the expense of triglycerides^[78], while the increased ratio of stearic to arachidonic acid in NASH may

correlate with fibrosis^[79]. Altered abundance and composition of liver tissue lipids may modulate the biological activity of KC in NAFLD through a number of mechanisms. First, the space-occupying effect of fat-laden hepatocytes may lead to impaired sinusoidal perfusion^[80]. Leukocytes trapped in narrowed sinusoids may increasingly engage KC in the microvascular inflammatory response^[80]. Second, excessive exposure of KC to fatty acids may modulate pathways of inflammation and insulin resistance through interaction with cell surface receptors and intracellular mediators^[81]. Third, anomalous deposition of lipids in the plasma membrane may alter the structure of lipid raft domains and interfere with clustering and function of cell surface receptors^[82]. Altered lipid composition may also affect proper functioning of intracellular membranes as seen with free cholesterol loading of mitochondria^[83]. Finally, abundant or abnormal lipids may confound recognition of fatty hepatocytes as dangerous and promote adverse interactions with KC^[17]. Nevertheless, the existence of a lipid-derived quintessential alarm expressed or released by steatotic hepatocytes remains speculative.

Recent findings indicate that TLR-mediated recognition of fatty acid moieties is an important mechanism by which lipids regulate pathways of inflammation and innate immunity^[82]. Depending on fatty acid composition, the outcome of this effect may be highly variable. Saturated fatty acids, implicated in the development of chronic conditions such as atherosclerosis, have been shown to activate TLR4 signalling in adipocytes and macrophages through both Myd88-dependent and TR-IF-dependent pathways^[84,85]. In contrast, polyunsaturated fatty acids inhibit these events in several cell types including macrophages^[85]. Consequently, TLR4 is a sensor of endogenous fatty acid levels and composition, and KC most likely benefit from this ability.

Emerging evidence indicates that altered cholesterol metabolism may directly affect the function of KC. Thus, high-fat diet fed to LDL receptor deficient mice rapidly results in significant hepatic inflammation, but only if the diet contains cholesterol^[86]. The presence of “foamy” KC suggests that scavenging of modified lipoproteins may induce this early inflammatory response^[86]. While these findings need to be extrapolated to human NAFLD with caution, they point to the importance of altered cholesterol metabolism. In addition, some of these observations challenge the “second-hit” concept since steatosis is not necessarily a forerunner of hepatic inflammation as these events may develop simultaneously^[86,87].

There is evidence that steatosis promotes Th1 polarization of the cytokine balance favouring innate or classic activation of macrophages in NAFLD^[88]. PPAR- α , PPAR- γ , and PPAR- σ and liver X receptors LXR- α and LXR- β are members of the nuclear hormone receptor superfamily of transcription factors that coordinate complex genetic programs of metabolism^[89,90]. Therapeutic use of synthetic ligands to target these receptors and exploit their biological functions is increasing. The beneficial effects of PPAR- γ in hepatocellular lipid homeostasis have prompted

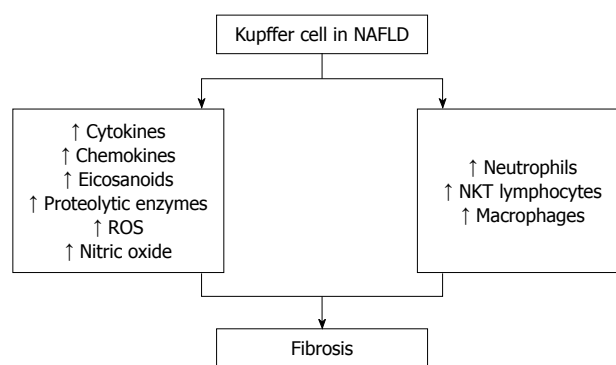


Figure 3 Effects of the activation of Kupffer cells in non-alcoholic fatty liver disease. NAFLD: Non-alcoholic fatty liver disease; NKT: Natural Killer T cells; ROS: Reactive oxygen species.

large clinical trials to assess impact on NAFLD and these efforts have been recently reviewed elsewhere^[91]. However, the recognition that nuclear hormone receptors link lipid metabolism to alternative activation of macrophages adds a new dimension to their potential use in the treatment of NAFLD^[88,92]. While PPAR- γ promotes alternative activation of macrophages that contribute to valuable metabolic changes such as improved insulin sensitivity^[93,94], recent research indicates that PPAR- σ is specifically required for a similar program in KC^[95,96]. Thus, signature gene expression of PPAR σ -deficient KC is greatly reduced in the livers of obese mice and in response to IL-4 stimulation^[95,96]. Moreover, PPAR σ ablation results in severe steatosis and insulin resistance^[95,96]. Notably, the effect of PPAR σ in KC is modulated by fatty acids^[95] and may fail due to altered lipid homeostasis and hepatic microenvironment in NAFLD. Thus, hepatocytes as a previously unsuspected source of Th2 cytokines stimulate M2 gene expression in KC and this important regulatory circuit may be altered in steatosis^[96]. These findings raise the intriguing possibility that specific targeting of PPAR- σ in KC to induce alternative activation may improve both inflammation and steatosis in NAFLD. One important caveat is that the M2 phenotype includes stimulation of the extracellular matrix that may contribute to hepatic fibrosis^[97]. Figure 3 shows the effects of activation of Kupffer cells in NAFLD.

In the last few years, there is increasing evidence that ligands of Hedgehog (Hh) may have a critical role in processes leading to liver fibrosis. The Hh mediated activity is quite low in healthy liver but increases during the course of several liver diseases, as recently reviewed^[98]. In particular, it has recently been shown that damaged/dying hepatocytes may produce Hh ligands that mediate proliferation of myofibroblasts in the liver, thus promoting fibrosis^[99]. Moreover, Hh seems to be critical due to its properties in regulating NKT growth and functions in liver fibrosis^[100,101].

IMMATURE MYELOID CELLS AND STEATOSIS

Immature myeloid cells (CD11b⁺Gr-1⁺) play a role in the

induction of inflammatory cytokines^[102] through activation of innate immune pathways. The role that immature myeloid cell populations play in obesity-related liver disease is unknown. In a recent study, Deng *et al.*^[103] hypothesize that accumulation of immature myeloid cells in the liver may be an important component in the development of inflammatory responses in liver tissue that are triggered by obesity, which in turn contributes to metabolic consequences, such as steatohepatitis. In this study, the liver of obese mice was demonstrated as the major organ where CD11b⁺Ly6C⁺-Ly6G⁻ immature myeloid cells accumulate. It is not clear why these cells are preferentially recruited into the liver. Chemotactic cytokines and chemokines could direct the migration of immune cells including myeloid cells and may be responsible for the cell accumulation. Several hepatic cell populations, including hepatocytes, KC, sinusoidal endothelial cells, and hepatic stellate cells, can secrete chemokines upon activation. High-fat diet-derived products could activate one of these cells in the liver, resulting in the recruitment of these circulating activated immature myeloid cells into the liver. IL-6 is overexpressed in the NAFLD patient^[104], and IL-6 has been shown to block immature myeloid cell differentiation^[105]. As a result, these activated cells are accumulated in the liver. The specific role of chemokines or other factors in the recruitment of these cells to the liver warrants further investigation.

CONCLUSION

It is now recognized that adipose tissue is an active endocrine organ that secretes numerous molecules that play a central role in the regulation of energy and vascular as well as immune system homeostasis by acting both locally and at distant sites influencing various metabolic and immune processes. Many of these interactions between metabolic and immune systems seem to be orchestrated by this complex network of soluble mediators derived from immune cells and adipocytes that are briefly summarized in Figure 4.

NAFLD is becoming an increasingly relevant clinical issue, especially in the developed world. One of the unmet challenges of NAFLD is to satisfactorily predict its progression from simple steatosis into steatohepatitis. This transition represents a milestone in the natural history with a considerable probability for developing end-stage liver disease. Elucidation of molecular and cellular events that may lead to this outcome is therefore critically important. Fortunately, the past few years have brought remarkable advances in our understanding of NAFLD pathogenesis, often by extension of research in adipose tissue biology, obesity, and insulin resistance. These efforts point to the intricate relationship of the innate immune system and lipid homeostasis in NAFLD with a prominent role for Kupffer, myeloid and NKT cells and a number of biochemical and cellular mechanisms involved.

However, a number of questions regarding the role of macrophage infiltration in human obesity remain to be an-

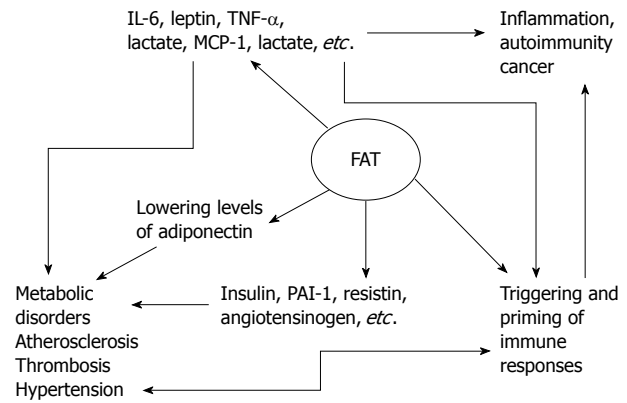


Figure 4 Complex network of soluble mediators derived from immune cells and adipocytes. MCP: Monocyte chemotactic protein; TNF- α : Tumor necrosis factor α ; IL: Interleukin.

swered. For example, what is the cause/s of macrophage infiltration? Does moderate fat gain alter macrophage number and/or macrophage phenotype in humans? Are some individuals predisposed to this? Is macrophage infiltration causal in the development of insulin resistance? The activation of NKT cells exacerbates macrophage infiltration in adipose tissue and glucose intolerance with obesity. Therefore, NKT cells enhance chronic inflammation in visceral adipose tissue and contribute to the development of metabolic disorders in obesity. The NKT cells may be the novel therapeutic targets in atherosclerosis, metabolic syndrome, and type 2 diabetes.

Further studies are needed to fully understand the interaction between fat, the immune system and steatosis.

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