

Adipose tissue dysfunction and the pathogenesis of metabolic syndrome

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stress conditions in adipose tissue that not only correlates with insulin resistance but is also causative in its development. Oxidative stress may be a mechanistic link between several components of metabolic syndrome and cardiovascular diseases, through its role in inflammation and its ability to disrupt insulin-signaling. The study around adipose tissue dysfunction will help to understand the pathogenesis of metabolic syndrome and may bring effective therapy in treatment of metabolic syndrome related diseases. Therefore, this review mainly focuses on the roles of adipose tissue dysfunction in inflammation, insulin resistance, and oxidative stress in the pathogenesis of metabolic syndrome.

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Key words: Metabolic syndrome; Adipose tissue dysfunction; Insulin resistance; Inflammation; Oxidative stress

Abstract

Metabolic syndrome is a growing research area. The underlying mechanisms of metabolic syndrome are still not very clear. Insulin resistance, obesity, inflammation and oxidative stress may play an important role in the pathogenesis of metabolic syndrome. The role of adipose tissue dysfunction is emphasized during the development of obesity. Adipose tissue is identified as a complex endocrine organ and its metabolic functions extend well beyond the classical actions of thermoregulation and of storage and release of fatty acids. Chronic low-grade inflammation activated by the immune system in adipose tissue is a key contributing factor to type 2 diabetes mellitus and cardiovascular diseases. Visceral obesity results in cell autonomous impairment in insulin signaling that leads to insulin resistance. Chronic inflammation in adipose tissue has gained acceptance as a lead promoter of insulin resistance in obesity. Furthermore, obesity creates oxidative

Core tip Metabolic syndrome is a growing research area. Insulin resistance, obesity, inflammation and oxidative stress may play an important role in the pathogenesis of metabolic syndrome. The role of adipose tissue dysfunction is emphasized during the development of obesity in recent years. The study around adipose tissue dysfunction will help to understand the pathogenesis of metabolic syndrome and may bring effective therapy in treatment of metabolic syndrome related diseases. Therefore, this review mainly focuses on the roles of adipose tissue dysfunction in inflammation, insulin resistance, and oxidative stress in the pathogenesis of metabolic syndrome.

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INTRODUCTION

Metabolic syndrome (MetS) is associated with the mortality and morbidity in cardiovascular diseases (CVD). The underlying mechanisms of MetS are still not very clear. Insulin resistance, obesity, inflammation and oxidative stress may play an important role in the pathogenesis of MetS. The roles of adipose tissue dysfunction in obesity have been addressed in recent years. Several mechanisms including insulin resistance, sub-inflammatory state, over activity of renin-angiotensin-aldosterone system (RAAS), oxidative stress, autonomic dysregulation as well as mechanical compression on the kidneys are all activated by obesity^[1]. Therefore, this review mainly focuses on the roles of adipose tissue in inflammation, insulin resistance, and oxidative stress in the pathogenesis of MetS.

ADIPOSE TISSUE AND ENDOCRINE FUNCTION

The prevalence of obesity has increased throughout the last three decades due to genetic, metabolic, behavioral, environmental^[2], and epigenetics^[3,4] factors. Excess fat is no longer associated with wealth, but is instead recognized as a risk factor for many diseases, such as type 2 diabetic (T2DM), CVD, fatty liver disease and some forms of cancer. Adipose tissue is increasingly being identified as a vital, complex endocrine organ, and not simply as a fat store^[5]. It has become increasingly clear that adipose tissue is a much more complex organ than was initially considered and that its metabolic functions extend well beyond the classical actions of thermoregulation and of storage and release of fatty acids^[6]. Furthermore, obesity is associated with an increased mortality and morbidity for CVD and adipose tissue is recognized as an important player in obesity-mediated CVD^[7].

The discovery of leptin in 1994 sparked dramatic new interest in the study of white adipose tissue (WAT)^[8]. As a key endocrine organ, adipose tissue releases multiple bioactive substances, known as adipose-derived secreted factors or adipokines, which have proinflammatory or anti-inflammatory activities^[9]. These adipokines include leptin, free fatty acids (FFA), tumour necrosis factor- α (TNF- α), interleukin-6 (IL-6), C-reactive protein, resistin, angiotensinogen and adiponectin (ApN)^[10,11]. Retinol binding protein 4 (RBP4) is a recently identified adipokine suggested to link obesity with its comorbidities, especially insulin resistance, T2DM, and certain components of the MetS^[12]. It is also abundantly clear that the dysregulation of adipokine secretion and action that occurs in obesity plays a fundamental role in the development of a variety of cardiometabolic disorders, including the MetS, T2DM, inflammatory disorders, and vascular disorders, that ultimately lead to coronary heart disease^[6]. The link between obesity, inflammation and insulin resistance indicates the important secretory role of adipose tissue^[13,14]. In addition, adipocytokine have been proposed as additional molecules able to modulate

kidney function^[15]. Epidemiological studies have also repeatedly reported the association between insulin resistance and kidney dysfunction in both non diabetic and diabetic subjects^[15].

An increase in adipose tissue mass is associated with the augmented secretion of certain adipokines, such as IL-6, TNF- α and resistin, which cause endothelial dysfunction and hemostasis alterations that also favor a prothrombotic state^[11]. Obesity and the accompanying MetS are associated with endothelial dysfunction, such as arterial hypertension and atherosclerosis^[16]. This detrimental effect of obesity is mediated, in part, by excessive production of the adipose tissue hormone leptin. Under pathological conditions, such as obesity and MetS, the NO-mediated vasodilatory effect of leptin is impaired^[16]. ApN is an important white and brown adipose tissue hormone, which is an insulin sensitizing hormone^[17] and has anti-inflammatory properties^[18]. ApN exerts its action through its receptors AdipoR1, AdipoR2, and T-cadherin^[17] and exerts a pivotal role in vascular protection through activation of multiple intracellular signaling cascades^[18]. ApN is highly abundant in human serum but its levels are reduced in obesity and are even lower in patients with hepatic steatosis or non-alcoholic steatohepatitis^[19,20]. Decreased plasma ApN levels are implicated in the pathogenesis of the MetS and atherosclerosis and may serve as a diagnostic and prognostic biomarker as well as a rational pharmaco-therapeutic target to treat these disorders^[18]. The level of leptin increases with the increase in weight gain, while ApN decreases. Measuring circulating levels of leptin and ApN as a screening tool may help recognize those individuals who do not only have obesity as a major risk factor toward developing cardiometabolic disease but also may have an unfavorable "biomarker profile", putting them at highest risk^[21].

ADIPOSE TISSUE AND INFLAMMATION

Obesity and its comorbidities, including T2DM and CVD, are considered to be a state of chronic low-grade inflammation that can be detected both systemically and within specific tissues^[22,23]. This obesity-associated chronic tissue inflammation is a key contributing factor to T2DM and CVD^[24]. Furthermore, chronic low-grade inflammation occurring in the adipose tissue of obese individuals is causally linked to the pathogenesis of insulin resistance and the MetS^[25]. Pickup *et al*^[26] found that abnormalities of the innate immune system may be a contributor to the hypertriglyceridaemia, low HDL cholesterol, hypertension, glucose intolerance, insulin resistance and accelerated atherosclerosis of T2DM. Their initial studies supported the hypothesis that type 2 diabetes is caused by activated innate immunity and led to research that has uncovered links between insulin resistance, obesity, circulating immune markers, immunogenetic susceptibility, macrophage function and chronic infection^[27]. ApN, leptin and other inflammatory proteins have been shown to correlate with insulin resistance

and the MetS in adults^[22]. A higher inflammation status was significantly correlated with decreases in the levels of antioxidant enzymes, ApN and an increase in the risk of MetS^[28]. A number of studies have clearly demonstrated that the immune system and metabolism are highly integrated^[24]. This link allows mammals to adapt to changes in their internal and external environments and affects organism-wide function^[29]. Obesity-induced inflammation is mainly mediated by tissue resident immune cells, with particular attention being focused on adipose tissue macrophages (ATMs)^[30], as accumulating evidence has revealed a critical involvement of inflammatory responses triggered by lesional macrophages in the pathogenesis of MetS^[31]. Moreover, Gene silencing of inflammatory cytokines TNF- α or osteopontin in epididymal ATMs of obese mice caused significant improvement in glucose tolerance^[32]. These data were consistent with the hypothesis that cytokines produced by ATMs can exacerbate whole-body glucose intolerance^[32]. Based on *in vitro* studies, macrophages can be divided into M1 and M2 classifications^[33]. M1 macrophages, also termed “classically activated macrophages,” are highly proinflammatory, secreting the bulk of the cytokines that cause insulin resistance. M2 macrophages, also termed “alternatively activated macrophages,” are not inflammatory and give rise to cytokines that exert anti-inflammatory effects, such as IL-10 and IL-4^[33]. The overall macrophage-induced inflammatory state of the tissue is determined by the balance between these different macrophage subpopulations. In the obese state, the balance is clearly tilted toward the proinflammatory macrophage phenotype^[34]. Recently, more leukocyte subpopulations have been implicated in obesity, including neutrophils, eosinophils, and mast cells^[35]. Neutrophils, which participate in inflammation-induced metabolic disease^[36], and mast cells^[37] are increased in obese adipose tissue, and studies in mice have indicated that these two cell types can promote insulin resistance. The involvement of multiple leukocyte subpopulations underlines the complexity of obesity-associated AT inflammation^[35]. The role of innate immune cells, such as macrophages in AT inflammation has been well demonstrated. In contrast, less is known about the role of lymphocytes^[25]. However, more recently, cells of the adaptive immune system, specifically B and T lymphocytes, have emerged as unexpected promoters and controllers of insulin resistance^[31], and participate in modulating adipose tissue inflammation during the development of obesity^[38]. Furthermore, fluctuations in weight have been associated with worsened metabolic and cardiovascular outcomes^[39]. Weight cycling did increase the number of CD4⁺ and CD8⁺ T cells in AT, indicating that an exaggerated adaptive immune response in adipose tissue may contribute to metabolic dysfunction during weight cycling, although adipose tissue macrophage number and polarization were not modulated by weight cycling^[39]. Molecular mechanisms are complicated in VAT inflammation. Several studies during

the past two decades have highlighted the key role of the IKK kinase (IKK)/nuclear factor- κ B (NF- κ B) pathway in the induction and maintenance of the state of inflammation that underlies metabolic diseases such as obesity and T2DM^[40]. Excess adipose tissue is hypothesized to contribute to a state of chronic inflammation which promotes development of insulin resistance as well as other metabolic complications by stimulating NF- κ B and Jun N-terminal kinase (JNK) pathways in adipocytes and the liver^[2]. JNK in macrophages is required for the establishment of obesity-induced insulin resistance and inflammation^[41,42]. Furthermore, bone marrow mesenchymal stem cells from high-fat diet animals showed increased production of IL-1, IL-6, and TNF- α and increased NF- κ B and reduced peroxisome proliferator-activated receptor gamma (PPAR- γ) expression^[43], suggesting the inflammatory responses during weight gain. In addition, mice with a null mutation for transient receptor potential vanilloid (TRPV4) or wild-type mice treated with a TRPV4 antagonist showed elevated thermogenesis in adipose tissues and were protected from diet-induced obesity, adipose inflammation, and insulin resistance^[44]. A causal role for iron in adipocytes as a risk factor for MetS and a role for adipocytes in modulating metabolism through ApN in response to iron stores have also been reported^[45], suggesting that adipocyte iron regulates ApN and insulin sensitivity^[45].

Excess visceral fat causes local chronic low-grade inflammation and dysregulation of adipocytokines, which contribute to the pathogenesis of the MetS^[46]. The amount of visceral adipose tissue (VAT) and the liver fat content are important factors responsible for the link between abdominal obesity and features of the MetS^[47]. In addition, visceral fat adiposity also correlates with inflammation in peripheral blood cells^[46]. Individuals with MetS have a higher degree of endothelial dysfunction and inflammation compared with individuals with multiple CV risk factors and may therefore have an increased CV risk beyond the contributions of multiple traditional risk factors^[48]. The inflammatory profile often observed among sedentary overweight/obese individuals with an excess of VAT/liver fat may be a consequence of a more primary defect in subcutaneous adipose tissue^[47]. To address the hypothesis that lowering inflammation will lower vascular event rates, two large-scale placebo controlled trials using targeted anti-inflammatory agents for the secondary prevention of myocardial infarction have been initiated^[49]. These inflammatory pathways are potential novel pharmacological targets for the management of obesity-associated insulin resistance^[50]. Areas of active investigation focus on the molecular bases of metabolic inflammation and potential pathogenic roles in insulin resistance, diabetes, and CVD. Translating the information gathered from experimental models of insulin resistance and diabetes into meaningful therapeutic interventions is a tantalizing goal with long-term global health implications^[23].

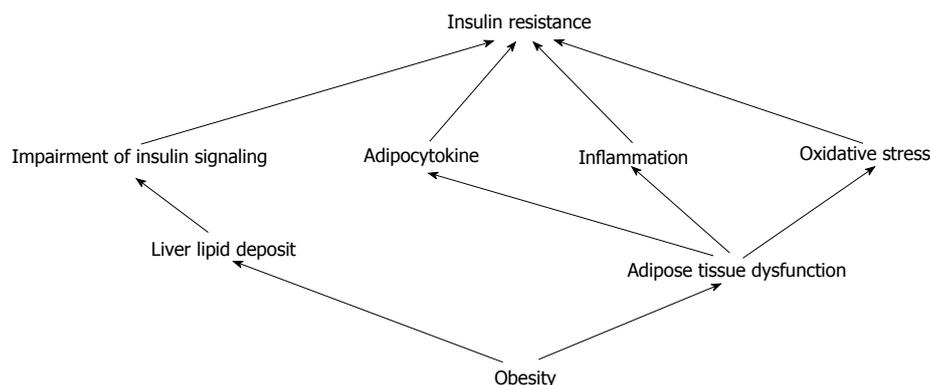


Figure 1 Adipose tissue dysfunction and insulin resistance. Obesity leads to adipose tissue dysfunction and liver lipid deposit. Adipose tissue dysfunction contributes to adipocytokine secretion, inflammation, and oxidative stress, etc. These factors finally lead to insulin resistance. Liver lipid deposit is closely associated with (hepatic) insulin resistance.

ADIPOSE TISSUE AND INSULIN RESISTANCE

Insulin resistance is a major characteristic of visceral obesity^[51]. The association between obesity and insulin resistance is an area of much interest and enormous public health impact^[52] (Figure 1). Visceral obesity, but not subcutaneous, results in cell autonomous impairment in insulin signaling that leads to insulin resistance. The mechanisms by which visceral obesity results in insulin resistance may be related to excess lipid accumulation in liver^[52]. Furthermore, it is not merely an increased mass of adipose tissue that directly leads to attenuation of insulin action, but rather adipose tissue inflammation activated by the immune system in obese individuals that leads to insulin resistance^[53]. VAT is prone to inflammation and inflammatory cytokine production, which also contribute to impairment in insulin signaling^[52]. Chronic inflammation in VAT has gained acceptance as a lead promoter of insulin resistance in obesity^[54-57]. The chronic state of insulin resistance in established obesity may be largely mediated by macrophage-induced proinflammatory actions, whereas the early-onset insulin resistance during high-fat diet feeding may be more likely related to acute tissue lipid overload^[58].

During obesity, many immune cells infiltrate or populate in adipose tissue and promote a low-grade chronic inflammation. Perturbation of inflammation is critically linked to nutrient metabolic pathways and to obesity-associated complications such as insulin resistance and T2DM^[59]. A great deal of evidence has pointed to the role of innate immune cells, in particular, adipose tissue macrophages, in the regulation of fat inflammation and glucose homeostasis^[56]. An increased accumulation of macrophages occurring in WAT has emerged as a key process in metabolic inflammation^[23,55] and insulin resistance in obesity^[60]. An association between adipose macrophage content and systemic insulin resistance was reported in diabetic humans^[61], suggesting the important role of inflammation in insulin resistance. Furthermore, IFN- γ is a central regulator of macrophage function and

play an important role in the regulation of inflammation and glucose homeostasis in obesity through multiple potential mechanisms, including effects on adipogenesis, cytokine expression, and macrophage phenotype^[60].

Recently, cells of the adaptive immune system, specifically B and T lymphocytes, have emerged as unexpected promoters and controllers of insulin resistance during the development of obesity^[56]. Adipose tissue contains a population of invariant natural killer T (iNKT) cells, whose abundance decreases with increased adiposity and insulin resistance^[62]. Adipose tissue-resident iNKT cells maintain healthy adipose tissue through direct interplay with adipocytes and prevent insulin resistance^[63]. Activation of NKT cells promotes M2 Macrophage polarization in adipose tissue and improves systemic glucose tolerance *via* IL-4/STAT6 protein signaling axis in obese adipose tissue^[62]. However, activation of iNKT by lipid excess promotes tissue inflammation and insulin resistance in obese mice, suggesting the role of iNKT cells in the complex network linking lipid excess to inflammation in obesity^[64]. In addition, iNKT cells do not affect glucose clearance but rather modulate lipid metabolism in both liver and adipose tissue^[65]. These effects on lipid metabolism are mainly mediated in the liver^[65]. A unique population of VAT-resident regulatory T (Treg) cells was recently implicated in control of the inflammatory state of adipose tissue and, thereby, insulin sensitivity. Unexpectedly, PPAR- γ expression by VAT Treg cells was necessary for complete restoration of insulin sensitivity in obese mice by the thiazolidinedione drug pioglitazone^[66]. Obese patients with insulin resistance displayed significantly decreased natural Tregs but an increase in adaptive Tregs in their VAT as compared with lean control subjects, suggesting a potential therapeutic value of Tregs to improve insulin resistance and end organ damage in T2DM by limiting the proinflammatory milieu^[67]. Obesity is characterized by circulating immune cells that are activated and insulin resistant, with the T-cell balance polarized towards a pro-inflammatory Th1 phenotype. The loss of insulin-induced suppression of inflammatory phenotypes in circulating immune cells could contrib-

ute to the systemic and adipose tissue inflammation^[68].

Visceral obesity results in insulin resistance appear to be related to excess lipid accumulation in liver^[52]. Visceral obesity is a main risk factor for non-alcoholic fatty liver disease (NAFLD)^[19], which is a low-grade chronic inflammatory state^[69]. Visceral obesity lead to inappropriate storage of triglycerides in adipocytes and higher concentrations of FFA may add to increased hepatic lipid storage and insulin resistance^[19]. The liver is directly exposed to increasing amounts of FFA and pro-inflammatory factors released from visceral fat into the portal vein of obese patients, promoting the development of hepatic insulin resistance, liver steatosis^[70] and progressive liver damage^[19]. Although liver lipid is closely associated with, and likely to be an important contributor to, (hepatic) insulin resistance^[71,72], it may also be in part the consequence of the lipogenic pathway of insulin action being up-regulated by hyperinsulinemia and unimpaired signaling^[71]. The 5-lipoxygenase pathway plays a major role in mounting inflammation in hepatic tissue and has emerged as a pathogenic factor in obesity-induced NAFLD. Therefore, modulation of lipoxygenases represents a novel target in the prevention of adipose tissue and hepatic dysfunction related to the MetS^[73]. Moreover, insulin resistance and lipotoxicity represent the missing links (beyond the classical cardiovascular risk factors) that help explain the accelerated rate of CVD in type 2 diabetic patients^[74]. AMP-activated protein kinase (AMPK) is considered as a master switch in regulating glucose and lipid metabolism. In the liver, activation of AMPK results in decreased production of plasma glucose, cholesterol, triglyceride and enhanced fatty acid oxidation^[75]. Interestingly, genetic deletion of the AMPK $\beta 1$ subunit in mice (referred to herein as $\beta 1(-/-)$ mice) reduced macrophage AMPK activity, acetyl-CoA carboxylase phosphorylation, and mitochondrial content, resulting in reduced rates of fatty acid oxidation. Thus, activation of AMPK $\beta 1$ and increasing fatty acid oxidation in macrophages may represent a new therapeutic approach for the treatment of insulin resistance^[76].

Endoplasmic reticulum (ER) stress in various cells plays an important role in the pathogenesis of several diseases^[77]. During the last decade, ER stress has emerged as a new player in the field of obesity, T2DM and insulin resistance, and a considerable number of recent studies have pointed out its role in the onset of insulin resistance^[51]. Furthermore, ER stress plays important pathophysiological roles in obesity-induced adipose tissue dysfunction^[78]. When the adipocyte endoplasmic reticulum is no longer capable of processing the excess nutrients, the so-called “endoplasmic reticulum stress” develops. This triggers efflux of FFA from adipocytes into the circulation and causes triglyceride overload in skeletal muscle, liver and pancreas^[79]. FFA is an important factor that has been implicated in the pathogenesis of insulin resistance^[80]. With a positive caloric balance, more FFA is released into the portal system. Excess of

circulating FFA, TNF- α and other factors induces insulin resistance^[81]. The mechanisms are related to inhibiting insulin signaling through the activation of serin-kinases, which promote a mechanism of serine phosphorylation of insulin receptor substrates, leading to interruption of the downstream insulin receptor signaling^[81]. Reducing plasma FFA concentration in obese and T2DM subjects improves insulin sensitivity. Moreover, pharmacologic FFA reduction improves insulin signaling in muscle from insulin resistant subjects. This beneficial effect on insulin action could be related to a decrease in local inflammation^[80]. Adipose tissue expansion not only involves enlargement of fat cells, but also the accumulation of inflammatory cells and a shift in the production of adipokines and cytokines^[82]. Inflammatory cytokines, ROS and ectopic lipid deposition are the main mediators of insulin resistance and vascular impairment^[79]. Impaired insulin signaling on vascular endothelium, atherosclerotic plaque macrophages can alter progression of CVD in the MetS and affect development of microvascular complications of diabetes mellitus^[83]. The specific cellular underpinnings or mechanisms of insulin resistance are not clear^[84]. The precise causes of insulin resistance are varied, and the relative importance of each is a matter of ongoing research^[85]. Advances in understanding of the complex pathophysiology of insulin's effects on vascular tissues will offer new opportunities for preventing these cardiovascular disorders^[83].

ADIPOSE TISSUE AND OXIDATIVE STRESS

The excess activation and the imbalance in the metabolism of oxygen and production of excess free radicals contribute to “oxidative stress” in the heart, vascular and kidney tissue^[86]. NADPH oxidase is the enzyme responsible for much of the generation of $\cdot O_2^-$ in cardiovascular tissue^[86,87], which is comprised of several membrane and cytosolic subunits that mobilize and activate under various agonists such as Ang II, aldosterone as well as fatty acids^[86,87]. MetS is associated with high oxidative stress, which is caused by an increased expression of NADPH oxidase and a decreased expression of antioxidant enzymes in the adipose tissue^[88]. Obesity creates oxidant conditions that favor the development of comorbid diseases^[89]. Oxidative stress in adipose tissue not only correlates with insulin resistance but is also causative in its development^[90]. Adipose tissue plays a central role in maintaining metabolic homeostasis under normal conditions^[78]. Energy imbalances lead to the storage of excess energy in adipocytes, resulting in both hypertrophy and hyperplasia. These processes are associated with abnormalities of adipocyte function, particularly mitochondrial stress and disrupted endoplasmic reticulum function^[89]. Oxidative stress plays a pivotal role in the pathogenesis of the MetS and in the progression of its complications^[91]. Oxidative stress may be a mechanistic

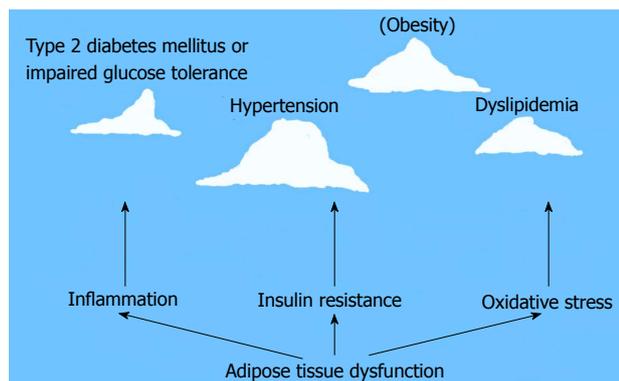


Figure 2 The role of adipose tissue dysfunction in the pathogenesis of metabolic syndrome. The mechanisms of metabolic syndrome are just as the base of iceberg under the sea level, such as insulin resistance, adipose tissue dysfunction, inflammation, and oxidative stress, etc. The relative clinical manifestations are as the top part of iceberg above the sea level, including hypertension, obesity, type 2 diabetes mellitus or impaired glucose tolerance, dyslipidemia, etc.

link between several components of MetS and CVD, through its role in inflammation and its ability to disrupt insulin-signaling^[7]. The cross-talk between impaired insulin-signaling and inflammatory pathways enhances both metabolic insulin resistance and endothelial dysfunction, which synergize to predispose to CVD^[7].

All components of the RAAS are expressed in and have independent regulation of adipose tissue. This local adipose RAAS exerts important auto/paracrine functions in modulating lipogenesis, lipolysis, adipogenesis as well as systemic and adipose tissue inflammation^[92]. The role of the RAAS on the development of insulin resistance and T2DM is an area of growing interest^[93]. Excess visceral adiposity contributes to inappropriate activation of the RAAS despite a state of volume expansion and of salt retention that contributes to subclinical elevations of pro-oxidant mechanisms. These adverse effects are mediated by excess generation of ROS and diminished antioxidant defense mechanisms^[86]. Extending beyond Ang II as the classical effector peptide, aldosterone has been shown to promote vascular production of oxidative stress through the enzyme complex NADPH oxidase independent of Ang II^[86,94,95]. In addition, aldosterone has been shown to potentiate the impact of Ang II impairments in endothelium-dependent relaxation both directly and indirectly through increased vascular oxidative stress resulting in reductions in the bioavailable nitric oxide^[86,94-96]. Inappropriate mineralocorticoid receptor activation has been demonstrated to be a causal factor in several pathologic conditions such as vascular inflammation, endothelial dysfunction, insulin resistance and obesity^[97].

Oxidative stress is positively associated with VAT as well as diffuse and focal carotid atherosclerosis in apparently healthy men and women^[98]. Increased adipose tissue oxygen tension in obese compared with lean men is accompanied by insulin resistance, and inflammation^[99]. Moreover, there is a synergistic effect of redox-

inflammatory processes to each of the components of the MetS^[100]. Using the available plasma oxidative stress biomarkers, many clinical studies have shown the presence of systemic oxidative stress in obese insulin resistant subjects, and its decrease after the successful treatment of obesity^[90]. Therefore, the evaluation of oxidative status may allow for the identification of patients at an increased risk of complications^[89].

CONCLUSION

MetS is a growing research area. The roles of VAT dysfunction in pathogenesis of MetS are emphasized in recent years. Adipose tissue dysfunction may lead to insulin resistance, inflammation and oxidative stress, even over activation of RAAS. These pathological mechanisms of MetS are just as the base of iceberg under the sea level. Hypertension, obesity, T2DM or impaired glucose tolerance, and dyslipidemia etc. are observed as the top part of floating iceberg which is above the sea level (Figure 2). The study around VAT dysfunction will help to understand the pathogenesis of MetS and may bring effective therapy in treatment of MetS related diseases.

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We apologize for the inability to cite numerous examples of important work in the field.

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