

Chapter 1

Function and Dysfunction of Adipose Tissue

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Abstract Adipose tissue is an endocrine organ which is responsible for postprandial uptake of glucose and fatty acids, consequently producing a broad range of adipokines controlling several physiological functions like appetite, insulin sensitivity and secretion, immunity, coagulation, and vascular tone, among others. Many aspects of adipose tissue pathophysiology in metabolic diseases have been described in the last years. Recent data suggest two main factors for adipose tissue dysfunction: accumulation of nonesterified fatty acids and their secondary products and hypoxia. Both of these factors are thought to be on the basis of low-grade inflammatory activation, further increasing metabolic dysregulation in adipose tissue. In turn, inflammation is involved in the inhibition of substrate uptake, alteration of the secretory profile, stimulation of angiogenesis, and recruitment of further inflammatory cells, which creates an inflammatory feedback in the tissue and is responsible for long-term establishment of insulin resistance.

Keywords Nutrient storage • Adipokines • Lipid intermediates • Hypoxia • Inflammation • Angiogenesis

1.1 Adipose Tissue Structure

Adipose tissue is a complex and heterogeneous tissue composed by cells with lipid storage functions, called adipocytes, and a stromovascular function, composed by endothelial and mesenchymal stem cells, preadipocytes, fibroblasts, and resident cells from the immune system (Rajala and Scherer 2003; Juge-Aubry et al. 2005a;

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Guilherme et al. 2008b; Christiaens and Lijnen 2010). During the embryonic development, the vascular network develops before adipocytes and the extracellular matrix which supports blood vessels is the first to be deposited, showing the crucial role of the vascular system in adipose tissue development (Neels et al. 2004; Christiaens and Lijnen 2010). In fact, during the embryonic development, a close communication between the stromovascular fraction and the adipocytes results in a mutual control between angiogenesis and adipogenesis. Recent data show that adipocytes may develop from capillary networks as the progenitor cells respond to pro-angiogenic stimuli in association with the expanding capillaries (Min et al. 2016).

In adult life, a well-developed vascular network is observable at the microscope and each adipocyte is surrounded by at least one capillary (Neels et al. 2004; Rutkowski et al. 2009; Christiaens and Lijnen 2010). The capillaries of the adipose tissue are fenestrated and are rich in trans-endothelial channels, which allow a close communication with the adipocytes (Christiaens and Lijnen 2010). Moreover, even in the adult life, this vascular network is very dynamic and is continuously adapting to changing nutritional fluxes. However, the mechanisms governing such remodeling are still far from being understood. Interestingly, the dynamics of vascular remodeling apparently influence adipocyte behavior during expansion. Adipocyte hypertrophy is usually associated with the formation of aberrant capillaries, while adipocyte hyperplasia is usually associated with increased angiogenesis and development of new capillaries. Hyperplasia is considered a harmless form of adipose tissue expansion, given the formation of smaller well-irrigated adipocytes with lower inflammatory activity than hypertrophic ones (Christiaens and Lijnen 2010). When the tissue is forced to expand, the formation of local phenomena of hypoxia leads to the expression of angiogenic factors which stimulate angiogenesis, including several cytokines and adipokines. The balance between these factors determines vessel density and permeability, and thus the “good” physiological or the “bad” pathophysiological expansion of the adipose tissue. The mechanisms will be detailed in the following sections.

1.2 Metabolic Functions of the Adipose Tissue

Although its important endocrine functions, the primary function of the adipose tissue is to store energy in the form of lipids, mainly in intracellular triglycerides droplets, also regulating lipid catabolism in different tissues due to the actions of adipokines (Rajala and Scherer 2003; Juge-Aubry et al. 2005a; Guilherme et al. 2008b). Lipid droplets are coated by a group of proteins, which the main one is Perilipin A (Per A), which prevent the contact between the stored triglycerides and the cytoplasm. In times of energetic need, such triglycerides are quickly hydrolyzed into free fatty acids and glycerol, a process called lipolysis, and released to the blood in order to feed other organs demands (Arner 2005; Guilherme et al. 2008b; Galic et al. 2010). Thus, the adipose tissue is able to recognize the metabolic state of the organism, not only by local energetic sensors, but also through different inputs coming mainly from the gut after and between meals. Moreover, adipocytes also regulate cholesterol metabolism, as they are able to produce high-density lipoproteins (HDL) in order to send excessive cholesterol to the liver.

When adipocytes accumulate excessive amounts of nonesterified fatty acids, for reasons that are currently under investigation, their metabolism and endocrine function are shifted in order to produce a broad range of proinflammatory factors, while adiponectin secretion is decreased. Such factors include cytokines and chemokines, growth factors, tumour necrosis factor (TNF)- α , interleukin (IL)-6, monocyte chemoattractant factor (MCP)-1, vascular endothelial growth factor (VEGF), leptin, and resistin (Wellen and Hotamisligil 2005; Tilg and Moschen 2006; Guilherme et al. 2008b). In fact, a strong relationship between metabolism and innate immunity is present at the adipose tissue. Many of the intracellular pathways involved in metabolic signaling are recruited as well during an immune response and most of the adipokines and adipose tissue-derived factors, besides regulating metabolism, also have paracrine and endocrine functions in regulating the immune response. Many authors support the existence of a metabolism—immunity axis, as any metabolic change immediately induces alterations in the immune response and the regulation of metabolic fluxes usually involves the activation of intracellular inflammatory and stress pathways (Rajala and Scherer 2003; Juge-Aubry et al. 2005a; Goossens 2008; Guilherme et al. 2008b; Rutkowski et al. 2009).

1.2.1 Mechanisms of Nutrient Uptake and Storage in Adipocytes

Triglycerides synthesis, a process called esterification, occurs from one molecule of glucose-derived glycerol and three fatty acyl chains. Adipocytes have a limited ability to store glycogen and thus all the glucose that is not consumed in the adipocyte metabolism is transformed into glycerol and stored in the triglycerides pool (Tilg and Moschen 2006; Goossens 2008; Guilherme et al. 2008b). Esterification is mainly stimulated by insulin, which induces tyrosine kinase activity in its receptor (Fig. 1.1). This in turn leads to the tyrosine phosphorylation and activation of the insulin receptor substrate-1 (IRS-1) and initiates a signaling pathway which involves PI₃K and Akt/PKB activation. Among other actions, the activation of this signaling cascade leads to the translocation to the membrane of GLUT4-containing vesicles, allowing glucose uptake (Wellen and Hotamisligil 2005; Bugianesi et al. 2005). Besides inducing glucose uptake, insulin is also responsible for lipolysis inhibition, through the inhibition of adenylate cyclase, the main enzyme involved in AMPc synthesis. AMPc activates PKA, which in turn phosphorylates and activates hormone-sensitive lipase (HSL), the main enzyme involved in triglycerides hydrolyzation (Fig. 1.1). On the other hand, in times of nutrient demand, contra-regulatory hormones, like glucagon, cortisol, growth hormone, and adrenaline, increase AMPc levels, leading to the activation of HSL and thus the release of fatty acids into the circulation (Fig. 1.1) (Tilg and Moschen 2006; Goossens 2008; Guilherme et al. 2008b). Increased lipolysis is also observed in obese individuals due to the development of insulin resistance and the increased secretion of proinflammatory cytokines that promote lipolysis (Langin 2006; Guilherme et al. 2008b). The subsequent flux of free fatty acids from the adipose tissue to the circulation may in turn cause their

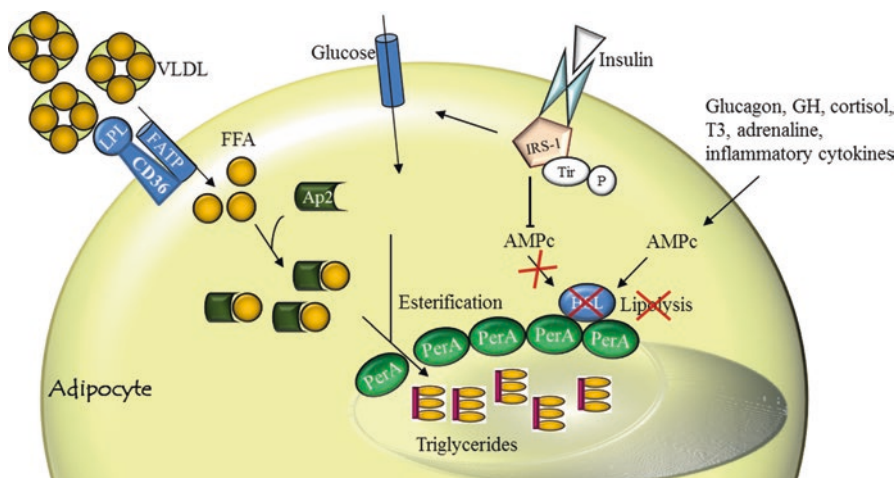


Fig. 1.1 Mechanisms of lipid storage in adipocytes and their mobilization from lipid droplets. Lipolysis is inhibited by insulin signaling and promoted by other hormones like glucagon, GH, cortisol, T3, or adrenaline, due to the stimulation of AMPc and HSL. *cAMP* cyclic adenosine monophosphate, *FATP* fatty acid binding protein, *FFA* free fatty acids, *HSL* hormone-sensitive lipase, *IRS-1* insulin receptor substrate-1, *LPL* lipoprotein lipase, *PerA* perilipin A, *VLDL* very-low density lipoprotein

ectopic accumulation in other tissues, such as the liver and the skeletal muscle, mechanism which will be described in the chapter dedicated to the pathophysiology of adipose tissue.

Circulating lipids are derived from hepatic incorporation into VLDL or from intestinal absorption and included in chylomicrons. Such lipoproteins bind to the CD36 receptor present at the adipocyte membrane. Lipoproteins are then hydrolyzed by the lipoprotein lipase (LPL) and the fatty acids are transported to the cytoplasm through the fatty acid transporter protein (FATP/CD36). Once at the cytoplasm the nonesterified fatty acids are captured by the protein ap2, which prevent their free circulation in the cell and the consecutive activation of inflammatory and stress pathways (Ram 2003; Wellen and Hotamisligil 2005). Fatty acids are finally esterified into triglycerides and stored in lipid droplets (Fig. 1.1).

Alternatively to esterification, fatty acids may be metabolized in several mediators of intracellular signaling pathways, namely eicosanoids, which are important activators of the peroxisome proliferation activated receptor-gamma (PPAR γ). This nuclear receptor controls the events involved in lipid esterification, being also activated by the pharmacological class of thiazolidinediones (TZD) (Fig. 1.2). The genes controlled by the activation of PPAR γ include proteins involved in fatty acid uptake (FATP, CD36 and LPL), metabolism (PEPCK and SCD-1), storage (perilipin A), and oxidation (Adiponectin and UCP-1). PPAR γ activation also inhibits cellular inflammatory pathways, including NF- κ B, which will be discussed in the chapter dedicated to adipose tissue pathophysiology (Wellen and Hotamisligil 2003; Ram 2003). Thus, PPAR γ promotes insulin sensitivity due to the reduction of cytoplasmic free fatty acids and due to the inhibition of inflammatory pathways (Ram 2003; Guilherme et al. 2008b).

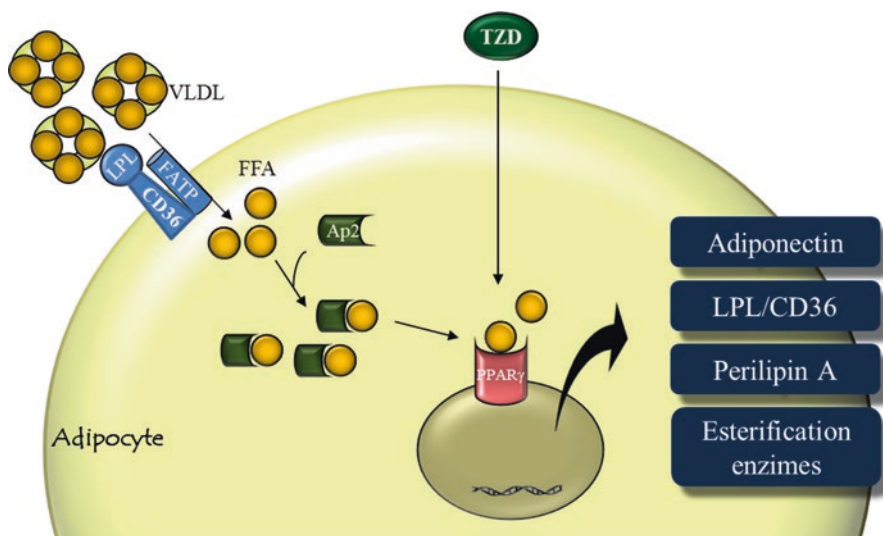


Fig. 1.2 Mechanisms of PPAR γ activation in response to fatty acids and their metabolites, stimulating the expression of adiponectin and proteins involved in fatty acid esterification and storage. The activity of PPAR γ can be improved by the pharmacological class of TZDs. *FATP* fatty acid binding protein, *FFA* free fatty acids, *LPL* lipoprotein lipase, *PPAR γ* peroxisome proliferating-activated receptor γ , *VLDL* very-low density lipoprotein

PPAR γ is expressed in adipocytes and macrophages, controlling lipid uptake, but also preadipocyte differentiation into mature adipocytes, a process called adipogenesis (Tamori et al. 2002; Lee et al. 2011). PPAR γ inhibition in obesity conducts to the inhibition of adipogenesis and to the hypertrophy of the existing adipocytes. Hypertrophic adipocytes have shown to be hypoxic and metabolically dysregulated (Trayhurn et al. 2008a; Trayhurn 2014). PPAR α controls the expression of genes involved in lipid oxidation and is expressed in tissues with catabolic activity like the liver and the skeletal muscle. It is regulated by adiponectin and by drugs like fibrates and metformin (indirectly), increasing the oxidation of fatty acids on mitochondria.

1.2.2 Cholesterol Fluxes

The adipocyte is also an important regulator of cholesterol storage and mobilization. Excessive intracellular cholesterol is incorporated into HDL/Apo-A1 particles, being transported to the liver where it used in the synthesis of biliary acids or excreted in bile (Yin et al. 2010). Two of the most important proteins in this transport are ABCA1 and ABCG1 (“ATP-Binding Membrane Cassete Transporter A1 e G1”), being involved in the transport of cholesterol, phospholipids, and other lipophilic substances to the HDL particles (Yin et al. 2010). The expression of ABC proteins is regulated by the PPAR and by the AMPc/PKA pathway, which is important in the mobilization of stored lipids (cholesterol). However, decreased PPAR activity in dysfunctional adipocytes

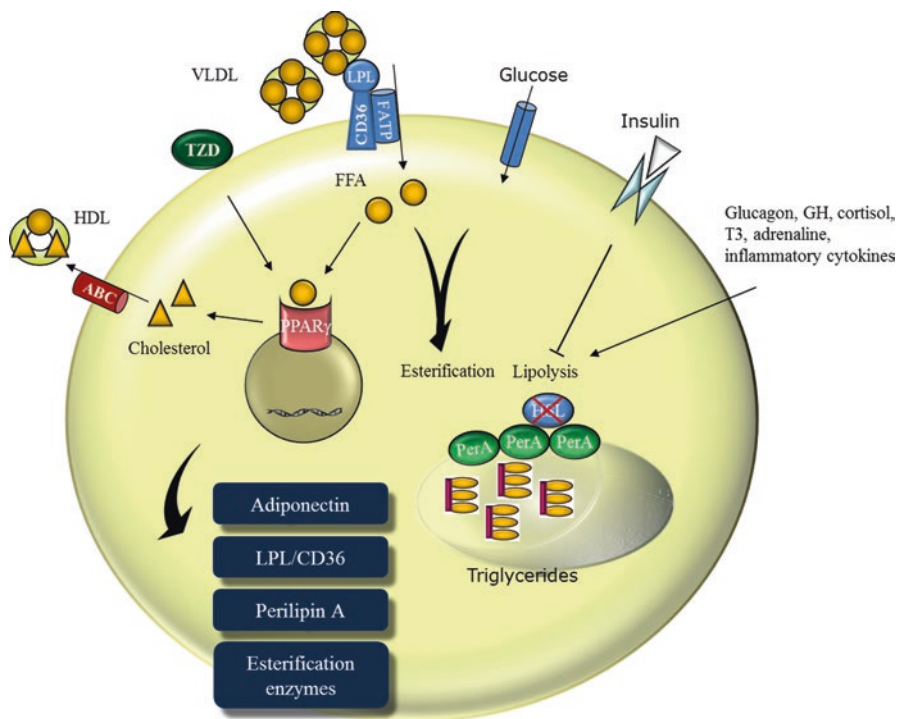


Fig. 1.3 Integration of the mechanisms involved in the uptake of lipids and glucose into the adipocyte, namely the role of insulin in inducing glucose uptake and in inhibiting lipolysis, as well as the role of PPAR γ in promoting fatty acid esterification and cholesterol efflux to HDL particles. *FATP* fatty acid binding protein, *FFA* free fatty acids, *HDL* high-density lipoprotein, *HSL* hormone-sensitive lipase, *LPL* lipoprotein lipase, *PerA* perilipin A, *PPAR γ* peroxisome proliferating-activated receptor γ , *VLDL* very-low density lipoprotein

may compromise cholesterol mobilization into the liver (Fig. 1.3) (Yin et al. 2010). Knockout models for ABCA1 have more infiltration of inflammatory cells in several tissues, due to decreased cholesterol transport to the liver and excessive deposition in the tissues. On the other hand, ABCA1 overexpression was shown to prevent cholesterol accumulation and atherosclerosis progression. Thus, ABCA1 prevents the accumulation of cholesterol in tissues, which is known to induce the formation of foam cells from recruited macrophages. The effect of ABCA1 and ABCG1 knockout is cumulative, suggesting that they have distinct roles in regulating cholesterol efflux. It is believed that ABCA1 and ABCG1 have distinct affinities for different HDL proteins (Yin et al. 2010).

1.3 Endocrine Function of the Adipose Tissue

The adipose tissue secretes a broad range of factors, including adipokines, cytokines, chemokines, angiogenic factors, coagulation factors, and vasoactive factors, among others. More than 600 different factors have been identified as produced and

secreted by the adipose tissue, affecting glucose and lipid metabolism, appetite, vascular function, inflammation, coagulation, or the cardiovascular function. However, the complete list of adipokines and their functions is not yet completely known (Wellen and Hotamisligil 2003; Trayhurn and Wood 2004; Guilherme et al. 2008b; Galic et al. 2010). Moreover, there are differences in the secretory profile between the different fat depots, with the visceral ones being more susceptible to nutritional signals due to their proximity to the intestine and liver.

1.3.1 Leptin

Leptin was the first factor originally identified as a product of adipose tissue and its discovery changed the view about this tissue, from a mere fat reserve to an endocrine organ able to control energy homeostasis. Leptin is almost exclusively produced by the adipocyte (95%) and its levels are proportional to the fat mass (Rajala and Scherer 2003; Meier and Gressner 2004; Golay and Ybarra 2005; Lorincz and Sukumar 2006; Vona-Davis and Rose 2007). Differences in leptin secretion from different fat depots were also shown, being the subcutaneous one more active (Nielsen et al. 2009). The main function of leptin is to inform the hypothalamus about nutrient availability. Leptin activates afferent nervous fibers and acts directly on the hypothalamus in order to suppress appetite and modulate energy expenditure (Fig. 1.4) (Kiess et al. 2008). Moreover, it acutely suppresses insulin secretion, but increases long-term β cell survival and function. Moreover, leptin increases fatty acid uptake and oxidation in the skeletal muscle and liver, also acting as a growth factor for endothelial cells (Rajala and Scherer 2003; Meier and Gressner 2004; Bugianesi et al. 2005; Lorincz and Sukumar 2006).

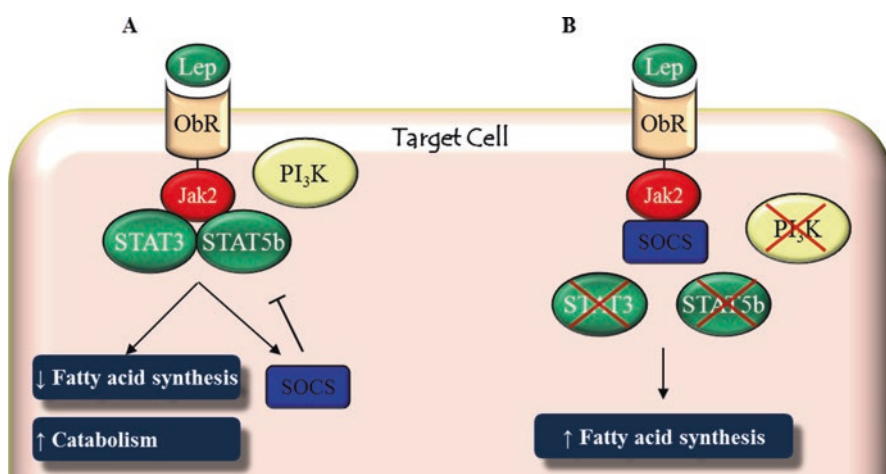


Fig. 1.4 Mechanisms of leptin signaling in target cells (a), as well the alterations occurring in leptin resistance (b). *Lep* Leptin, *ObR* leptin receptor, *SOCS* suppressor of cytokine signalling, *STAT* signal transducer and activator of transcription

Although leptin levels are proportional to fat mass, its secretion may be stimulated by insulin and inhibited by increased levels of AMPc, i.e., it increases after meals and decreases in times of energetic demand (Meier and Gressner 2004). When leptin levels are lower, appetite is stimulated and thyroid hormones, thermogenesis, and immune system function are inhibited (Münzberg et al. 2005). Mutations of the leptin or leptin receptor genes are known to induce obesity. On the other hand, leptin administration leads to an increase of fatty acid oxidation and a reduction of their circulating levels. Such effects are not observed in obese patients, which are known to have a resistance to the action of the hormone (Rajala and Scherer 2003; Meier and Gressner 2004). Other functions of leptin include the stimulation of angiogenesis, the modulation of the immune response, and the expression of the key enzyme involved in estrogen synthesis, aromatase (Catalano et al. 2003, Juge-Aubry et al. 2005a, b Lorincz and Sukumar 2006).

Leptin receptor is linked to a Jak/STAT pathway, which first protein is Jak2 and leads to the activation of STAT3 and STAT5. STAT proteins are transcription factors that stimulate catabolic processes and insulin secretion in β cells (Ueki et al. 2004a, b; Laubner et al. 2005; Kaneto et al. 2010; Blüher and Mantzoros 2015) (Fig. 1.4). STAT proteins also activate SOCS proteins, involved in the negative feedback. Thus, hyperleptinemia leads not only to increased signaling but also to increased negative feedback, contributing to the leptin resistance observed in obese patients mainly in the hypothalamus (Laubner et al. 2005). SOCS proteins also inhibit insulin signaling, contributing to insulin resistance. In fact, leptin and insulin signaling have a common pathway, namely the activation of IRS-1, PI3K, and Akt (Ahima and Flier 2000; Ueki et al. 2004a; Münzberg et al. 2005; Ahima 2005; Imrie et al. 2010). Leptin also activates AMPK inducing lipid oxidation and promoting insulin sensitivity (Rajala and Scherer 2003; Meier and Gressner 2004; Ahima 2005; Juge-Aubry et al. 2005a). Moreover, leptin inhibits fatty acid synthesis, due to the suppression of the transcription factor SREBP-1c and the key proteins of the biosynthetic pathway ACC and FAS. On the other hand, such mechanisms were shown to be activated by SOCS proteins (Fig. 1.4) (Ueki et al. 2004a, b).

1.3.2 *Adiponectin*

Adiponectin is an important regulator of lipid metabolism and stimulator of its oxidation produced in the adipocyte as a consequence of the activation of PPAR γ . Pharmacological activators of PPAR γ , glitazones, are known to increase adiponectinemia. On the other hand, adiponectin secretion is inhibited by the activation of inflammatory pathways and cAMP (Fig. 1.5). Thus, insulin also induced adiponectin secretion by reducing cAMP levels, while its elevation in times of energy demand prevents fatty acid consume (Xu et al. 2003).

Adiponectin was shown to have cardioprotective effects by promoting cell viability and inhibiting apoptosis during ischemia (Ding et al. 2012; Park and Sweeney 2013; Smekal and Vaclavik 2017). As well, similar effects were demonstrated in β

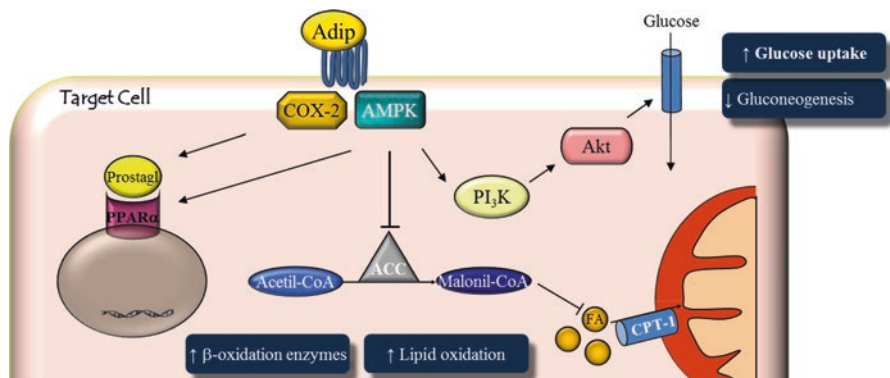


Fig. 1.5 Mechanisms of adiponectin signaling in target cell, including AMPK-mediated glucose uptake and inhibition of ACC. Inhibition of ACC prevents fatty acids synthesis and prevents inhibition of fatty acid uptake by the ACC product malonyl-CoA. Activation of PPAR α increases the expression of fatty acids oxidation enzymes. ACC acetyl-CoA carboxylase, Adip adiponectin, AMK AMP-activated protein kinase, COX-2 cyclooxygenas-2, PPAR γ peroxisome proliferating-activated receptor γ

cells, improving insulin secretion. Adiponectin was shown to exert anti-inflammatory effects on the vessel wall preventing atherosclerosis through the inhibition of adhesion molecules expression and smooth muscle cell proliferation (Fig. 1.5) (Juge-Aubry et al. 2005a). Protective effects of adiponectin have also been reported in different pathologies like cancer and neurodegenerative diseases (Vona-Davis and Rose 2007; Pais et al. 2009; Matafome 2013; Letra et al. 2014).

Adiponectin circulates in three molecular forms (trimeric, hexameric, and high-molecular weight isoform (HMW) composed of several hexamers). Reduced levels of the HMW isoform have been specifically associated with metabolic disorders (Rajala and Scherer 2003; Vona-Davis and Rose 2007, 2009; Pais et al. 2009). The globular region of the adiponectin chain binds to two membrane receptors, AdipoR1 and AdipoR2. While AdipoR1 is found mainly in the skeletal muscle, AdipoR2 is more abundant in the liver (Meier and Gressner 2004). A third receptor, T-cadherin, was recently identified as an adiponectin receptor, but without intracellular signaling and mainly involved in the anchorage of the HMW isoform to the membrane (Takeuchi et al. 2007). Adiponectin binding to AdipoR1 and AdipoR2 leads to AMPK and PPAR α activation, which in turn leads to increased uptake and oxidation of glucose and lipids (Fig. 1.5) (Kamon et al. 2003; Yamauchi et al. 2003; Lorincz and Sukumar 2006; Pais et al. 2009). Moreover, AMPK is involved in promoting glucose uptake through GLUT4 translocation to the membrane and in inhibiting gluconeogenesis (Juge-Aubry et al. 2005a; Nawrocki et al. 2006). Similar to leptin, adiponectin directly inhibits the enzymes involved in the fatty acids synthesis pathway, through the inhibition of acetyl-coA carboxylase (ACC), the key enzyme of such pathway (Fig. 1.5) (Ouchi et al. 2001; Xu et al. 2003; Nawrocki et al. 2006). On the other hand, the activation of PPAR α increases the expression of the enzymes involved in lipid oxidation and uptake to the mitochondria (Fig. 1.5) (Kamon et al. 2003; Yamauchi et al. 2003; Gealekman et al. 2008, 2012).

1.3.3 Resistin

Resistin is an adipokine associated with the establishment of insulin resistance as its levels are increased in models of obesity and type 2 diabetes. Studying the role of human resistin is not easy as significant differences were found to murine resistin. Resistin gene was found in different chromosomes and human resistin is mainly produced in macrophages while murine resistin is produced in adipocytes (Lazar 2007). The activation of the immune system after metabolic dysregulation induces resistin secretion, in order to activate pathways which block nutrient uptake by hypertrophic adipocytes and promote the release of those already stored. Resistin also induces angiogenesis aiming to increase blood flow and thus adipocyte oxygenation. Importantly, resistin increases lipid uptake by macrophages, also inhibiting cholesterol efflux from these cells, in order to store adipocyte-derived lipids. However, the chronic activation of such mechanisms leads to the formation of foam cells, common in atherosclerotic lesions (Fig. 1.6) (Lazar 2007; Robertson et al. 2009).

Resistin is directly involved in blocking insulin signaling in adipocytes, but also in the liver and skeletal muscle. Resistin knockout mice were shown to be protected of obesity-related insulin resistance, suggesting that it may be an important link between the activation of the immune system and glucose metabolism (Lazar 2007; Qatanani and Szwegold 2009; Robertson et al. 2009). Moreover, resistin neutralization increases insulin sensitivity and decreases hepatic glucose production, while

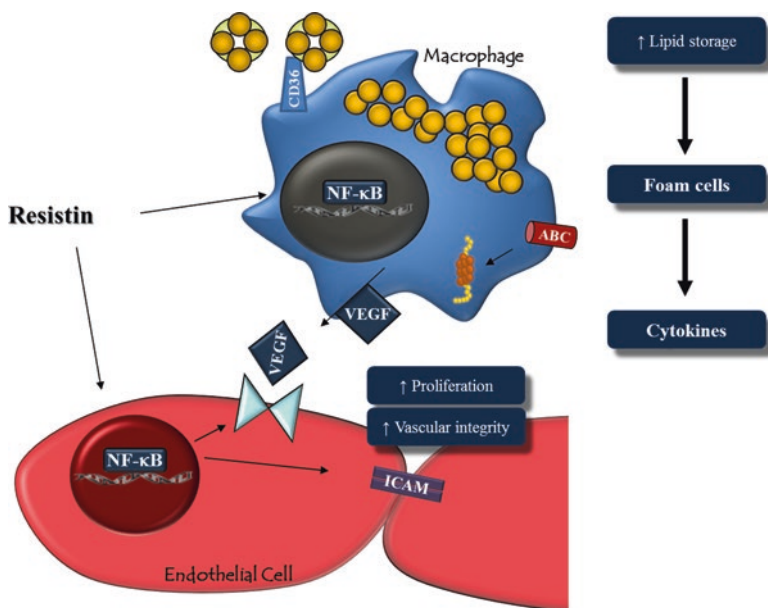


Fig. 1.6 Mechanisms of resistin action in macrophages and endothelial cells, promoting the formation of foam cells and endothelial proliferation. *ICAM* intercellular adhesion molecule, *NF-κB* nuclear factor-kappa B, *VEGF* vascular endothelial growth factor

resistin administration induces severe insulin resistance (Bugianesi et al. 2005; Juge-Aubry et al. 2005a; Lazar 2007). The expression of human resistin specifically in the macrophages of resistin knockout mice was able to increase adipose tissue lipolysis and fatty acid ectopic accumulation in the skeletal muscle, which led to insulin resistance (Qatanani and Szwegold 2009).

Resistin is a cysteine-rich protein which forms a dimer responsible for its biological activity (Bugianesi et al. 2005; Juge-Aubry et al. 2005a; Lazar 2007; Robertson et al. 2009; Smith and Yellon 2011). Human resistin was shown to induce the proinflammatory activity of macrophages and endothelial cells, by activating the Nuclear Factor- κ B (NF- κ B) and inhibiting the insulin signaling (Fig. 1.6). Resistin promotes the proliferation and adhesion of endothelial cells through NF- κ B activation and increased expression of the VEGF receptors (VEGFR1 and VEGFR2), matrix metalloproteinases (MMP-1 and MMP-2), and adhesion proteins (VCAM and ICAM) (Fig. 1.6) (Juge-Aubry et al. 2005a).

In macrophages, resistin promotes adipocyte-like mechanisms of lipid uptake resulting in the formation of foam cells. Such mechanisms include the upregulation of scavenger receptors (SR-A and CD36), also promoting the degradation of lipid efflux proteins in the proteasome (ABCA1, ABCG1). Foam cells are known for their proinflammatory and proangiogenic potential which is thus promoted by resistin actions (Fig. 1.6) (Lazar 2007; Lee et al. 2009; Qatanani and Szwegold 2009).

1.3.4 Other Adipose Tissue Products

Recently, a broad range of other adipokines has been identified, including apelin, visfatin, vaspin, omentin, and others involved in canonical mechanisms like coagulation, vascular tone, and immunity.

Visfatin was initially discovered as a factor controlling B lymphocytes development. On the other hand, an enzyme called Nampt (nicotinamide 5-fosforibosyl-1-pyrophosphate transferase) catalyzing the formation of nicotinamide adenine dinucleotide (NAD) was also described as product from the same gene (Kiess et al. 2008; Gallí et al. 2010; Saddi-Rosa et al. 2010). Thus, visfatin appears to be a factor with a simultaneous enzymatic and endocrine activity. Its circulating concentrations are correlated with the amount of visceral adipose tissue and increases during weight gain. Moreover, its synthesis was identified mostly in macrophages residing and infiltrating the tissue (Kiess et al. 2008; Gallí et al. 2010; Saddi-Rosa et al. 2010). Increased circulating visfatin levels were found in a number of pathological situations like portal inflammatory infiltration, steatohepatitis, coronary accumulation of oxidized LDL and atherosclerotic plaque instability, higher incidence of ischemic events and decreased kidney function (Saddi-Rosa et al. 2010). On the other hand, visfatin was observed to produce a rapid insulin-independent hypoglycemic effect after its administration, while animal models lacking visfatin have shown hypoinsulinemia and lower insulin-stimulated glucose uptake (Kiess et al. 2008; Gallí et al. 2010; Saddi-Rosa et al. 2010). Visfatin was also

observed to increase adipocyte and myocyte glucose uptake and to suppress hepatic glucose production (Saddi-Rosa et al. 2010). In vitro and in vivo studies also demonstrated protective effects of visfatin in ischemic events and its ability to increase endothelial cell activation, promoting angiogenesis (Mocan Hognogi and Simiti 2016). Thus, contradictory results about visfatin effects have been observed. Apparently, it is a compensatory mechanism for hyperglycemia with protective effects at multiple levels (Saddi-Rosa et al. 2010). Such effects were recently attributed to its ability to synthesize NAD, which improves the activity of several enzymes involved in glucose metabolism and mitochondrial function (Gallf et al. 2010; Mocan Hognogi and Simiti 2016).

Apelin was only recently identified, after the identification of its receptor APJ. Its functions are not very clear but apparently it regulates distinct biological processes like angiogenesis, vascular tone, heart function, and insulin secretion. Apelin was shown to antagonize Angiotensin (Ang)-2 effects of endothelial cells, increasing NO-dependent vasodilation, but it was also shown to increase smooth muscle contraction. Although these are contradictory results, apelin also has the ability to bind to antidiuretic hormone (ADH) producing neurons in the hypothalamus, decreasing its production. Thus, together with its ionotropic actions in the heart, apelin is thought to be involved in the control of blood pressure (Glassford et al. 2007; Pitkin et al. 2010; Fasshauer and Blüher 2015). Moreover, insulin was observed to increase Apelin secretion, but apelin apparently has inhibitory functions on insulin secretion, suggesting that it might be involved in a negative feedback to insulin secretion (Glassford et al. 2007; Pitkin et al. 2010; Fasshauer and Blüher 2015). Nevertheless, little is yet known about apelin and more research is needed in this field.

Vaspin is another adipose-derived hormone, initially identified as inhibitor of serine proteases, but with benefic effects on insulin sensitivity. Although the mechanisms of vaspin stimulation and secretion are unknown, it was shown to be secreted by visceral fat mass and nutritionally regulated, due to its circadian release after meals (Jeong et al. 2010; Fasshauer and Blüher 2015). Vaspin levels are decreased in type 2 diabetes and its administration was shown to improve glucose tolerance and insulin sensitivity (Fig. 1.7) (Klötting et al. 2006; Jeong et al. 2010; Fasshauer and Blüher 2015). Moreover, diet or surgery-induced weight loss leads to a decrease of vaspin levels, which may result from decreased fat mass (Klötting et al. 2006; Handisurya et al. 2010).

Omentin is mainly produced in the omental adipose tissue, but it is expressed in vascular cells and not in adipocytes (Schäffler et al. 2005; Yang et al. 2006; Yamawaki et al. 2010). Omentin was recently suggested to increase insulin-stimulated glucose uptake, being correlated with insulin sensitivity. Its concentrations were observed to be decreased in obese patients and to increase after weight loss (Fig. 1.7) (Yang et al. 2006; De Souza Batista et al. 2007; Moreno-Navarrete et al. 2010). Importantly, omentin was shown to induce eNOS-dependent vessel relaxation, although the mechanisms are still unknown (Yamawaki et al. 2010).

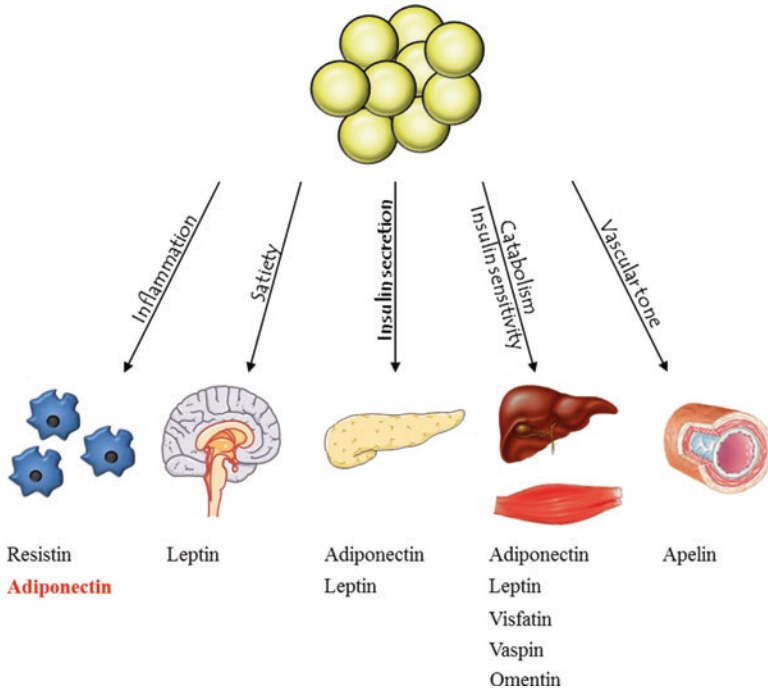


Fig. 1.7 Effects of adipose tissue-derived endocrine factors on inflammation, appetite, sensitivity and secretion of insulin and vascular tone

1.4 Dysregulation of Adipose Tissue Function in Obesity

Adipocyte hypertrophy is a process occurring after excessive nutrient uptake, especially in visceral adipose tissue, which is more susceptible to nutrient fluxes. Hypertrophy has been recently associated with the development of hypoxic regions, with a strong influence on the metabolic and secretory functions of the adipose tissue.

Several studies demonstrated inflammation as the link between metabolic dysregulation and hypoxia, changing the secretory profile to create a proangiogenic environment (Fig. 1.8). Inflammation blocks nutrient uptake (insulin resistance and PPAR γ inhibition) in order to prevent uncontrolled cell and tissue expansion and promotes lipolysis. Dysfunctional adipose tissue secretes a broad range of cytokines, but also chemokines, which recruit circulating monocytes to the tissue. Infiltrating macrophages are then important in maintaining the inflammatory environment and in uptake of excessive lipids, leading to the formation of adipose tissue foam cells. Many authors claim the existence of a metabolism—immunity axis, given that all metabolic changes lead to an immune response and many of the inflammatory factors simultaneously regulate metabolism and the immune system (Fig. 1.8) (Rajala and Scherer 2003; Golay and Ybarra 2005; Juge-Aubry et al.

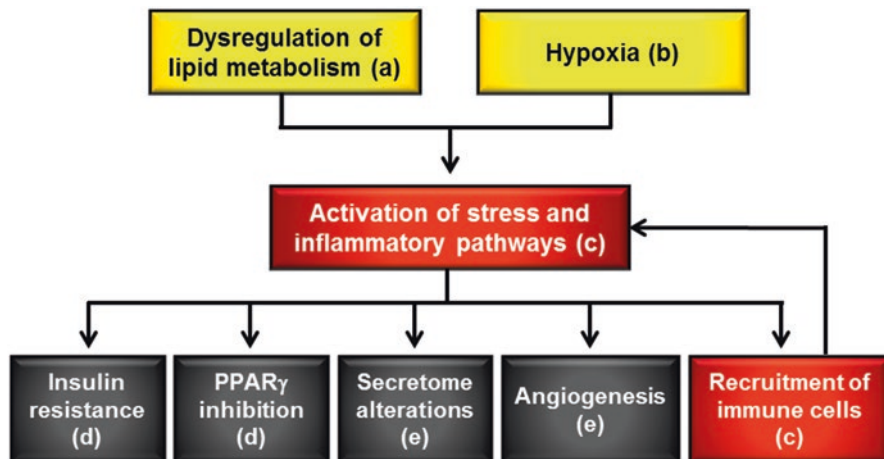


Fig. 1.8 The role of hypoxia (b) and dysregulation of the lipid metabolism (a) in activating inflammatory mechanisms (c), aiming to restore homeostasis. Inflammation inhibits nutrient uptake and promotes lipolysis (d), also increasing angiogenesis (e) and the secretion of inflammatory adipokines (e). The mechanisms involved will be discussed in the following sections according to the letters in each figure's box

2005a; Ye et al. 2007; Goossens 2008; Guilherme et al. 2008b; Rutkowski et al. 2009; Maury and Brichard 2010).

1.4.1 *Dysregulation of Lipid Metabolism*

During the process of fat accumulation, adipocytes continuously accumulate triglycerides, increasing the expression of enzymes involved in fatty acid esterification. When adipose tissue is not able anymore to store all the lipids from the diet, adipocytes release fatty acids to the circulation, which then accumulate in tissues like skeletal muscle and liver. Type 2 diabetic patients usually have hepatic and muscle steatosis, leading to insulin-resistance and morphological alterations in such tissues (Rajala and Scherer 2003; Guilherme et al. 2008b; Kawano and Cohen 2013; Lonardo et al. 2015). The increasing availability of fatty acids also increases their esterification into triglycerides and oxidation in the mitochondria (Bugianesi et al. 2005; Golay and Ybarra 2005). However, lipid oxidation is a limited process and in such conditions intermediates of fatty acid metabolism like ceramides and diacylglycerols accumulate. Such intermediates inhibit insulin signaling and glucose uptake (Fig. 1.9). In physiological condition, such mechanisms prevent excessive nutrient uptake, which would cause cell death by oxidative stress. However when such mechanisms are chronically activated they conduce to insulin resistance and contribute to the onset of type 2 diabetes (Golay and Ybarra 2005; Guilherme et al. 2008a).

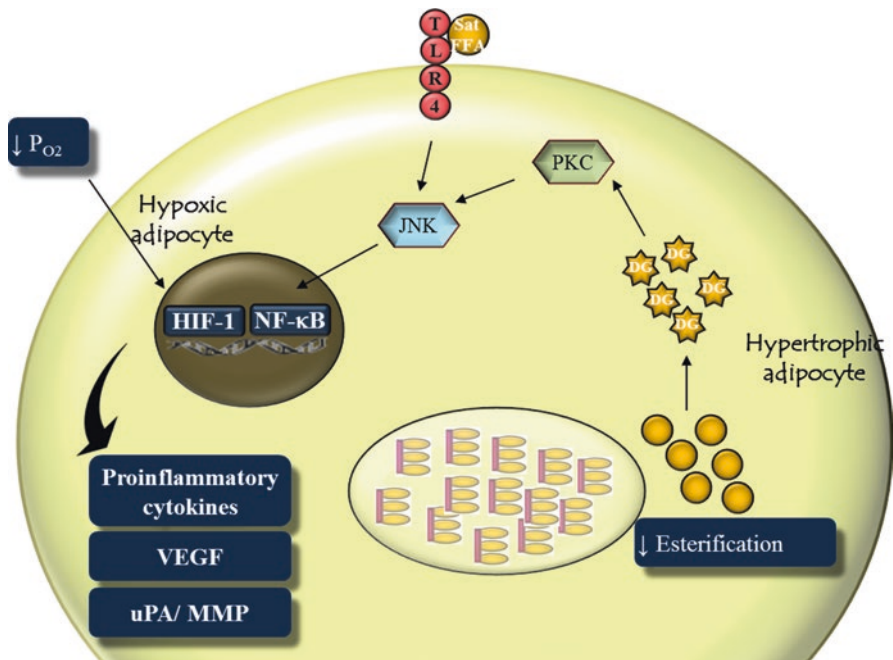


Fig. 1.9 Main mechanisms conducting to the activation of inflammatory pathways in adipocytes, namely the cytoplasmatic accumulation of lipid mediators and activation of membrane receptors by saturated fatty acids. *FFA-Sat* saturated free fatty acids, *HIF-1* hypoxia inducible factor-1, *JNK c-jun n-terminal kinase*, *MMP* matrix metalloproteinase, *P_{O2}* oxygen pressure, *TLR4* toll-like receptor 4, *TNF- α* tumor necrosis factor alpha, *uPA* plasminogen activator urokinase-type, *VEGF* vascular endothelial growth factor, *VEGFR* VEGF receptor

The accumulation of diacylglycerols directly activates several PKC isoforms which besides inhibiting insulin receptor activation through serine phosphorylation, also promotes the activation of the IKK β (Fig. 1.9) (Moeschel et al. 2004; Boden et al. 2005; Guilherme et al. 2008b). IKK β is also a serine kinase which directly inhibits the insulin receptor and leads to the activation of the NF- κ B. Moreover, JNK is also activated by the PKC and by fatty acid-activated toll-like receptor 4 (TLR4), contributing to the same final objective of NF- κ B activation (Qatanani and Lazar 2007; Zhang et al. 2010). In turn, NF- κ B activated the expression of proinflammatory and proangiogenic molecules (MCP-1, TNF- α , VEGF, adhesion molecules, etc.), which further contribute to the inhibition of substrate uptake (Fig. 1.9).

1.4.2 Adipose Tissue Hypoxia

It has been demonstrated that angiogenesis is a vital process in adipose tissue expansion and proper nutrient storage. Tissue expansion is believed to lead to the formation of physiological hypoxic regions, which induce the secretion of angiogenic factors and the reestablishment of tissue homeostasis. However, chronic hypoxia

has been suggested to dramatically change adipokine secretion, with a decrease of adiponectin and an increase of proinflammatory adipokines (Hosogai et al. 2007; Wang et al. 2007; Goossens 2008; Trayhurn et al. 2008b; Wood et al. 2009). Despite the reasons to the development of chronic hypoxia are not known, it was described that obese individuals lack the postprandial increase of blood supply to the adipose tissue observed in lean individuals (Trayhurn et al. 2008b; Wood et al. 2009). Hypoxic regions have been shown in several diet-induced and genetic animal models of obesity, showing deficient angiogenesis, oxygen pressure, with decreased levels of endothelial cell markers and increased accumulation of the hypoxia probe pimonidazole and lactate formation (Hosogai et al. 2007; Ye et al. 2007; Goossens 2008; Pang et al. 2008; Trayhurn et al. 2008b; Wood et al. 2009). Despite being currently controversial, a current view supports that the oxygen diffusion distance (100 μm) is lower than adipocyte diameter in obesity (150–20 μm). This would lead to hypoxia-inducible factor-1 (HIF-1) stabilization and consequent expression of GLUT1 and secretion of angiogenic factors like the VEGF (Wood et al. 2007, 2009; Ye et al. 2007; Wang et al. 2007; Rausch et al. 2008; Goossens 2008; Trayhurn et al. 2008a; He et al. 2011). However, this view is recently being abandoned and Goossens and colleagues have reported that the adipose tissue of obese individuals may even be hyperoxic under certain circumstances due to decreased metabolic activity (Goossens et al. 2011). Thus, the real causes for impaired vascular function and hypoxia (if confirmed) are currently unknown.

1.4.3 Activation of Stress and Inflammatory Pathways by Lipid Mediators and Hypoxia: Consequences for Inflammatory Cell Recruitment

There is no consensus about the role of adipocytes and preadipocytes in tissue response to hypoxia as current models does not discriminate the secretome change occurring in each cell type. On the other hand, cell lines do not mimic all the events observed in vivo. Nevertheless, several authors have suggested NF- κ B and HIF-1 activation leading to the expression of proinflammatory cytokines like IL-6, MCP-1, plasminogen activator inhibitor-1 (PAI-1), transforming growth factor (TGF)- β , and matrix metalloproteinases (MMP), but also leptin (Ye et al. 2007; Trayhurn et al. 2008b; Halberg et al. 2009; Wood et al. 2009). NF- κ B activation in hypoxia is well documented in different models of hypoxia like tumors, interacting with HIF-1 to increase the expression of inflammatory cells (Fig. 1.9) (van Uden et al. 2008). It has been described that cultured hypoxic adipocytes secrete higher levels of MCP-1, TNF- α , and VEGF and increase GLUT1 expression. Preadipocytes apparently are more susceptible to hypoxic stimuli, acquiring the ability to secrete leptin. However, hypoxia inhibits preadipocytes differentiation to adipocytes and promotes their transformation to macrophages. The release of chemoattractant factors like MCP-1 further increases the number of macrophages in the tissue (Trayhurn et al. 2008b; Wood et al. 2009).

1.4.3.1 Adipose Tissue Immune System: Recruitment of Inflammatory Cells and Secretion of Inflammatory Cytokines

Changes in the metabolic status are detected by the immune system, which acts in order to establish homeostasis. The activation of the immune system is an important mechanism in times of excessive nutritional fluxes, avoiding excessive nutrient uptake by the cells and maintaining cell viability and nutrient distribution by the different cells of the body. However, the low-grade inflammation observed in obesity resulting from chronic nutrient excess leads to insulin resistance in adipocytes (Galic et al. 2010). Such mechanisms include MCP-1 expression, which is involved in recruiting cells from the immune system to the tissue. Such cells, mainly monocytes-derived macrophages, further support the inflammatory feedback in the tissue, with the release of inflammatory mediators like TNF- α . In turn, inflammatory mediators further inhibit nutrient accumulation and tissue growth, showing the existence of a metabolism—immunity continuous (Schäffler et al. 2007; Galic et al. 2010).

A large number of circulating monocytes can be recruited to the adipose tissue through the action of proinflammatory cytokines and chemokines and develop into macrophages (Schäffler et al. 2007; Guilherme et al. 2008a; Surmi and Hasty 2010; Olefsky and Glass 2010). Macrophages may be derived from recruited monocytes and from differentiation from preadipocytes or mesenchymal cells and can become 50% of the total number of cells in the adipose tissue in obesity (Schäffler et al. 2007; Guilherme et al. 2008a; Surmi and Hasty 2010; Olefsky and Glass 2010). MCP-1, a major chemokine, was shown to be increased in obese patients and animal models. Its inhibition in diabetic and obese animal models improves insulin resistance and glucose tolerance (Guilherme et al. 2008a; Olefsky and Glass 2010).

The reason why preadipocytes can differentiate into macrophage is because they share the same cell lineage, sharing several intracellular mechanisms of lipid uptake, storage, and metabolism (Schäffler et al. 2007). Macrophages are known to surround dead adipocytes forming crown-like structures, which are typically found in the adipose tissue of obese patients and animal models (Trayhurn et al. 2008a). Such macrophages uptake and store large amounts of fat released from their lipid droplets, leading to the formation of foam cells. In turn, foam cells have an increased inflammatory activity with increased activation of the NF- κ B and secretion of proinflammatory factors (Ye et al. 2007; Rutkowski et al. 2009; Maury and Brichard 2010; Galic et al. 2010). Adipose tissue hypoxic regions were shown to be strongly populated by macrophages, being strongly positive for their membrane marker F4/80, and by CD4+ and CD8+ T cells, although their role in metabolic syndrome is currently unknown (Ye et al. 2007; Rausch et al. 2008; Goossens 2008; Trayhurn et al. 2008a; Wood et al. 2009).

Two populations of macrophages may be found at the adipose tissue: M1 macrophages recruited to the tissue by the gradient of chemokines like the MCP-1 typically have a more pronounced proinflammatory secretome, originating most of the foam cells. Tissue resident M2 macrophages secreting a broad range of angiogenic and tissue remodeling factors like metalloproteinases are involved in tissue adapta-

tion to hypoxia and have a more modest inflammatory activity (Maury and Brichard 2010; Olefsky and Glass 2010; Galic et al. 2010).

Activation of endothelial cells by inflammatory cytokines leads to the overexpression of adhesion molecules, which are important for macrophage binding and migration to the tissue matrix (Rutkowski et al. 2009; Maury and Brichard 2010). Moreover, the increased HIF-1-dependent expression of the migration inhibition factor (MIF-1) in hypoxia inhibits macrophage exit from the tissue, contributing to their accumulation (Ye et al. 2007). Most of the cytokines released by the adipose tissue in obesity, namely TNF- α , IL-6, and resistin, are derived from the infiltrating macrophages. TNF- α acts on adipocytes in order to inhibit insulin receptor and the PPAR γ and to activate NF- κ B, completing the inflammatory feedback in the tissue aiming to block further nutrient uptake and to increase angiogenesis and tissue remodeling (Wellen and Hotamisligil 2003; Moeschel et al. 2004; Sandu et al. 2005; Qatanani and Lazar 2007; Guilherme et al. 2008a; Olefsky and Glass 2010; Min et al. 2016). On the other hand, TNF- α and MCP-1 inhibition in models of diet-induced insulin resistance was shown to improve insulin sensitivity (Wellen and Hotamisligil 2003; Bugianesi et al. 2005; Guilherme et al. 2008b).

1.4.4 Inhibition of Nutrient Uptake by Inflammatory Stimuli: Insulin Resistance and PPAR γ Inhibition

Insulin resistance is characterized by a deficient cell response to physiological insulin concentrations, leading to increased β -cell insulin secretion and compensatory hyperinsulinemia (Yki-Järvinen 2005, 2015; Yki-Järvinen and Westerbacka 2005). Insulin resistance is not associated with the body mass index (BMI), but depends on impaired adipose tissue lipid storage and activation of inflammatory pathways (Yki-Järvinen 2005, 2015; Yki-Järvinen and Westerbacka 2005). Anti-inflammatory molecules like PPAR γ agonists (thiazolidinediones and polyunsaturated fatty acids) and salicylates are known to improve insulin sensitivity. As well, physical exercise is known to decrease inflammation and to improve insulin sensitivity (Bugianesi et al. 2005; Guilherme et al. 2008b; Oliveira et al. 2011).

Fatty acids are known to directly cause insulin resistance. Even nondiabetic individuals were shown to decrease insulin sensitivity after an acute exposure to fatty acids, which is a mechanism important to prevent excessive accumulation of non-esterified fatty acids in the cytoplasm (Golay and Ybarra 2005; Einstein et al. 2010). Circulating free fatty acids are also strong inducers of insulin resistance acting through the membrane receptor TLR4 (Qatanani and Lazar 2007; Guilherme et al. 2008a; Zhang et al. 2010; Olefsky and Glass 2010).

Hypoxia is a strong inhibitor of nutrient uptake. In cultured adipocytes hypoxia-dependent HIF-1 activation leads to the upregulation of stress pathways that blunts insulin signaling (Regazzetti et al. 2009; Ye 2009; Wood et al. 2009; Zhang et al. 2010). As well, the same mechanisms also inhibit PPAR γ activity, leading to decreased secretion of adiponectin (Gentil et al. 2006; Chen et al. 2006; Goossens

2008; Ye 2009). However, it was shown that the overexpression of an inactive form of HIF-1 during a high-fat diet challenge aggravates insulin resistance, suggesting distinct acute and chronic effects of hypoxia in adipose tissue. Such evidences support the idea that acute activation of hypoxia response mechanisms are important regulators of nutrient uptake be adipocytes, but long-term activation of such mechanisms is important to the regulation of mitochondrial biogenesis and angiogenesis (Zhang et al. 2010; He et al. 2011). In fact, adipose tissue vascular density was shown to correlate with insulin ability to suppress lipolysis, i.e., insulin sensitivity (Pasarica et al. 2010).

A main contributor to the long-term activation of such mechanisms is macrophage-derived inflammatory signals. Activation of inflammatory signals results not only in the inhibition of nutrient uptake but also the release of those already accumulated, namely through the stimulation of lipolysis. Cytokines like the TNF- α and IL-6 are known to be mostly produced in adipose tissue infiltrating macrophages. These cytokines were shown to activate NF- κ B through serine kinases like JNK and IKK, leading to the transcription of inflammatory factors and proteins involved in stress and inflammatory pathways (Fig. 1.10) (Tilg and Moschen 2006; Kiess et al. 2008; Olefsky and Glass 2010). Moreover, serine kinases phosphorylate

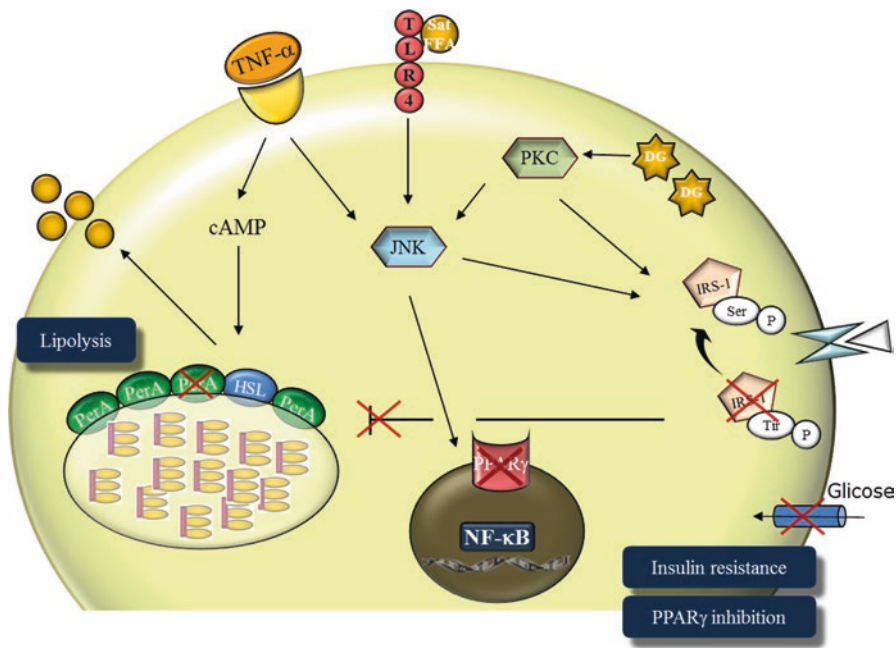


Fig. 1.10 Consequences of adipocyte inflammatory mechanisms for inhibition of insulin signaling and PPAR γ activity, inhibiting nutrient uptake and promoting lipolysis. *FFA-Sat* saturated free fatty acids, *IRS-1* insulin receptor substrate-1, *JNK* c-jun n-terminal kinase, *NF- κ B* nuclear factor kappa B, *PerA* perilipin A, *PKC* protein kinase C, *PPAR γ* peroxisome proliferating-activated receptor γ , *Ser* serine, *TLR4* toll-like receptor 4, *TNF- α* tumor necrosis factor alpha

the insulin receptor and its substrate (IRS-1) in serine residues instead of the stimulatory tyrosine phosphorylation, leading to the inactivation of the pathway (Fig. 1.10) (Moeschel et al. 2004; Boden et al. 2005; Wellen and Hotamisligil 2005; Qatanani and Lazar 2007; Guilherme et al. 2008a; Wang et al. 2009; Kaneto et al. 2010; Maury and Brichard 2010; Olefsky and Glass 2010; Galic et al. 2010). JNK was also shown to increase the expression of SOCS proteins, which are known to bind to and signalize IRS-1 to proteasome degradation (Ueki et al. 2004a; Howard and Flier 2006). Such mechanisms perpetuate the inhibition of insulin signaling in response to inflammatory cytokines.

Regarding lipid storage, TNF- α was shown to inhibit PPAR γ activity, causing the downregulation of key enzymes in fatty acid uptake (LPL and FATP) and esterification, as well as adiponectin (Boden et al. 2005; Juge-Aubry et al. 2005a; Qatanani and Lazar 2007; Guilherme et al. 2008b). TNF- α was also shown to increase intracellular levels of cAMP, which is known to activate hormone-sensitive lipase (HSL) and thus to induce lipolysis (Fig. 1.10) (Wellen and Hotamisligil 2003; Juge-Aubry et al. 2005a; Guilherme et al. 2008a; Maury and Brichard 2010; Galic et al. 2010). Interestingly, hypoxia is also associated with decreased levels of ATP and increased cAMP levels, also contributing to lipolysis (Ye 2009).

1.4.5 Activation of Angiogenesis

The formation of new blood vessels is a highly regulated process involving several cell types, which produce a broad range of factors controlling vessel permeability and stability. The plasticity of the vascular network is also modulated by a series of hormones involved in the metabolism, showing that the nutritional status has a major influence in the angiogenic stimulus. The activation of the immune system in metabolic syndrome dramatically changes the stimulus to endothelial cells, promoting vascularization as an attempt to restore homeostasis.

Obese patients with adipose tissue hypoxia were shown to have normal oxygen pressure and hemoglobin concentration and saturation, suggesting that hypoxia derives from local changes in blood supply (Goossens 2008; Rutkowski et al. 2009; Corvera and Gealekman 2014). Obese individuals may have a 30–40% reduction of blood supply than lean ones and a similar reduction was observed in obese Zucker rats (Ye 2009). Moreover, obese and type 2 diabetic animal models have been shown to have a decrease of the endothelial cells marker CD31, which may possibly be associated with the development of hypoxia (Goossens 2008; Ye 2009). Moreover, recent studies also demonstrated a reduction of endothelial progenitor cells, which may contribute to impaired angiogenic function (Neels et al. 2004; Gealekman et al. 2008; Tam et al. 2009; Corvera and Gealekman 2014).

Inhibition of angiogenesis was initially suggested as an effective strategy to prevent adipose tissue growth (Lijnen 2008). However, this hypothesis has been abandoned given that the vascular network was shown to be essential for proper tissue oxygenation and storage and metabolic functions and blood supply was shown to correlate with insulin sensitivity (Mannerås-Holm and Krook 2012). Moreover, it

was shown that adipose tissue expansion is necessary for nutrient storage and to prevent ectopic lipid deposition in liver and skeletal muscle.

The formation of hypoxic regions activates the secretion of angiogenic factors which ultimately restore homeostasis (Cao 2007; Goossens 2008; Rutkowski et al. 2009). However, if angiogenesis is not enough, hypoxia is maintained and HIF-1 levels increase in order to stimulate the expression of VEGF, angiopoietins, leptin, and matrix-remodeling factors (Minet et al. 2001; Yamakawa et al. 2003; Cao 2007; Hosogai et al. 2007; Goossens 2008). However, HIF-1 chronic activation also leads to the secretion of fibrotic factors (Rutkowski et al. 2009; Wood et al. 2009; Suga et al. 2010).

1.4.5.1 Pro and Antiangiogenic Factors

Adipose tissue angiogenesis is regulated by the balance between pro and antiangiogenic factors derived from all the cells of the tissue (Neels et al. 2004; Cao 2007; Rutkowski et al. 2009; Tinahones et al. 2012; Corvera and Gealekman 2014).

VEGF is the main endothelial cell growth factor, being associated with the development of obesity (Cao 2007; Rutkowski et al. 2009; Tinahones et al. 2012; Corvera and Gealekman 2014). Animal models of obesity and obese individuals have increased VEGF levels, which were shown to decrease after weight loss (Gómez-Ambrosi et al. 2010). VEGF is mostly produced by stroma cells and by mature adipocytes, acting in adjacent capillaries (Fig. 1.11) (Neels et al. 2004; Christiaens and Lijnen 2010). During adipose tissue expansion, a significant part of VEGF is produced by infiltrating inflammatory cells. VEGFR2 inhibition prevents diet-induced tissue expansion and preadipocyte differentiation, showing that angiogenesis is crucial to the development of new adipocytes (Tam et al. 2009; Tran et al. 2012; Min et al. 2016). The major regulators of VEGF expression are hypoxia and insulin, while inflammatory cytokines can also induce its expression. VEGF inhibition as a strategy to prevent obesity is still controversial. VEGF was shown to increase in genetic models of obesity, but not in models of metabolic syndrome, suggesting that hypoxia-induced VEGF overexpression may be lost in hyperglycemia (Trayhurn et al. 2008b; Wood et al. 2009). Such observations derive from impaired HIF-1 stabilization in hyperglycemic models, resulting in impaired VEGF expression and dysregulation of the angiogenic process (Bento et al. 2010b). Moreover, insulin resistance may also contribute to impaired VEGF expression in type 2 diabetes. Thus, the initial idea that inhibiting VEGF could be an effective strategy to prevent obesity is currently being changed and is now believed that VEGF is necessary for proper adipose tissue angiogenesis and function (Treins et al. 2002; Hausman and Richardson 2004; Ye 2009; Vona-Davis and Rose 2009).

VEGF effects on the vascular network depend on its interaction with angiopoietins (Ang). Ang-1 and Ang-2 have opposite effects on the vasculature, although both were shown to be necessary for embryonic development. While Ang-1 is involved in vessel stabilization, increasing cell-cell adhesion, Ang-2 induces destabilization of cell-cell and cell-matrix interactions in order to allow cell proliferation and migration (Cao 2007; Christiaens and Lijnen 2010; Corvera and Gealekman 2014).

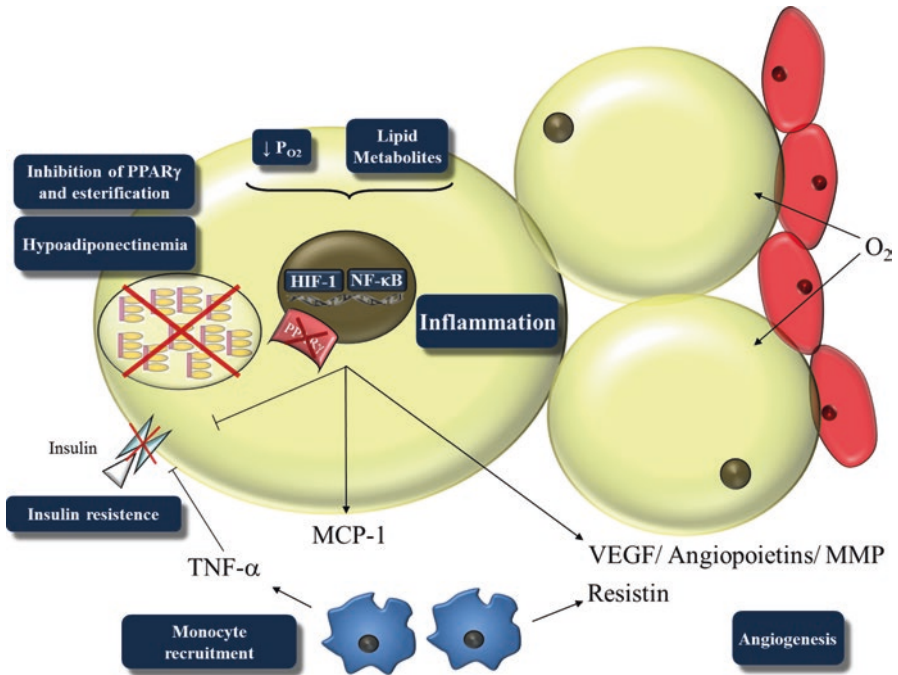


Fig. 1.11 Causes and consequences of inflammatory processes in adipose tissue. Inflammation directly inhibits excessive nutrient uptake by adipocytes and their continuous hypertrophy. Moreover, such mechanisms also stimulate angiogenic function in order to reestablish homeostasis. *HIF-1* hypoxia inducible factor-1, *MCP-1* monocyte chemoattractant protein-1, *MMP* matrix metalloproteinases, *NF- κ B* nuclear factor-kappa B, *P_{O₂}* oxygen pressure, *PPAR γ* peroxisome proliferating-activated receptor γ , *TNF- α* tumor necrosis factor alpha, *VEGF* vascular endothelial growth factor

Ang-1 promotes vessels stiffness through modulation of TGF β activity and deposition of extracellular matrix proteins (Cao 2007; Rutkowski et al. 2009; Christiaens and Lijnen 2010). On the other hand, Ang-2 expression is induced by hypoxia like VEGF and its receptors (Tie-2) are often co-localized with VEGFR2 (Christiaens and Lijnen 2010; Bento et al. 2010a). Ang-2 action in order to induce destabilization of cell interactions is necessary for VEGF-induced proliferation (Christiaens and Lijnen 2010; Bento et al. 2010a). The absence of VEGF after Ang-2 induced destabilization of endothelial cells leads to the formation of aberrant blood vessels. Such mechanisms have been suggested to be involved in the development of diabetic vascular complications and to be present in adipose tissue during the process of metabolic dysregulation (Bento et al. 2010a; Matafome et al. 2012).

Leptin was also shown to be proangiogenic, sharing many mechanisms with insulin, namely MAPK activation (Cao 2007; Vona-Davis and Rose 2007; Lijnen 2008; Christiaens and Lijnen 2010; Bento et al. 2010a). Leptin and insulin promote endothelial cells migration, proliferation, and survival, causing the increased VEGF secretion and VEGFR2 expression. Moreover, leptin was shown to increase MMP-

dependent matrix remodeling (Cao 2007; Vona-Davis and Rose 2007; Lijnen 2008; Christiaens and Lijnen 2010; Bento et al. 2010a). On the other hand, adiponectin was shown to counteract all these effects, by inhibiting VEGF actions in endothelial cells (Vona-Davis and Rose 2009). Adiponectin is mostly produced by small and non-inflamed adipocytes. Thus, when adipose tissue blood supply is enough, adiponectin prevents excessive blood vessel growth. On the other hand, leptin is produced proportionally to fat mass and thus its angiogenic functions are necessary to sustain tissue angiogenesis.

Other factors that have been suggested to be involved in the regulation of adipose tissue angiogenesis are fibroblast growth factor (FGF) (Lijnen 2008; Christiaens and Lijnen 2010), hepatic growth factor (HGF), and platelet-derived growth factor (PDGF) (Lijnen 2008; Pang et al. 2008; Christiaens and Lijnen 2010). As well, fibrotic and inflammatory factors like TGF- β and TNF- α have been shown to exert direct effects on adipose tissue endothelial cells. While TNF- α is involved in endothelial cell proliferation, TGF- β is important for the deposition of the extracellular matrix and vessel stabilization (Lijnen 2008; Niu et al. 2008; Christiaens and Lijnen 2010).

Angiogenesis is not only regulated by endothelial cell proliferation and migration, but also by matrix remodeling. Extracellular matrix degradation is a highly regulated process by the plasminogen system and by MMPs. The plasminogen system is known for its role on blood clotting but it is also shown to be involved in the regulation of extracellular matrix degradation (Lijnen 2008; Carter and Church 2009; Christiaens and Lijnen 2010). Animal models of obesity and metabolic syndrome have been shown to have increased levels of the inhibitor of plasminogen activator (PAI-1). Transgenic models of adipocyte PAI-1 expression have shown lack of visceral adipose tissue, showing that PAI-1 inhibits matrix degradation and vessel growth. However, although PAI-1 is apparently involved in preventing matrix degradation, its expression is increased by hypoxia and its levels are correlated with the body mass index (BMI) and insulin resistance, suggesting that its function in adipose tissue may be far more complex. Regarding MMPs, they are involved in matrix degradation, allowing endothelial cell migration. Their levels are increased in obesity and their overexpression results in decreased capillary density in adipose tissue. However, the mechanisms of their regulation and their real impact in adipose tissue pathophysiology are currently unknown (Lijnen 2008; Christiaens and Lijnen 2010; Corvera and Gealekman 2014).

1.4.6 Inflammatory Feedback in Adipose Tissue: Unifying Mechanism

The dysregulation of adipocyte metabolism occurs with the accumulation of fatty acids and their intermediates and with the development of hypoxia. All these conditions lead to the activation of stress pathways resulting in inhibition of nutrient uptake and increased expression of chemokines involved in the recruitment of inflammatory cells to the tissue (Galic et al. 2010). PPAR γ inhibition also leads to

decreased adiponectin secretion, which contributes to the inflammatory feedback in the tissue. Monocyte recruitment and their differentiation into macrophages further increase the release of inflammatory chemokines, cytokines like TNF- α , and angiogenic factors. Such events further chronically inhibit insulin signaling and fatty acid uptake, also increasing lipolysis (Fig. 1.11) (Qatanani and Lazar 2007; Olefsky and Glass 2010).

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