HYPERTENSION AND OBESITY (E REISIN, SECTION EDITOR)

Aldosterone Production and Signaling Dysregulation in Obesity

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Abstract In the past decades, we have extended the view of aldosterone effects beyond epithelial tissues. New evidence regarding the aldosterone/mineralocorticoid receptor (MR) pathway in active metabolic tissues, including adipose tissue, has confirmed its pathogenic role in systemic inflammation, endothelial dysfunction, insulin resistance, and dyslipidemia. Obesity, a current epidemic worldwide, increases aldosterone production by several adipocyte factors such as leptin but is also associated with local aldosterone production. In addition, obesity can modulate MR activation leading to signaling dysregulation and a pro-inflammatory profile of adipocytes. Current knowledge have deciphered that this phenotypical differences of obesity may be explained, at least in part, by novel non-genomic activation of MR, new inducers of aldosterone synthesis, and probably by several epigenetic modifications. In addition, with the understanding of the complex interplay of obesity, hormones, and receptors, targeted pharmacological therapy is expected and is currently under active research.

Keywords Aldosterone · Adipose tissue dysfunction · Mineralocorticoid receptor · Metabolic syndrome · GPR30 · Aldosterone synthase · Renin-angiotensin-aldosterone system · Adiponectin · Leptin · Obesity

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Introduction

Aldosterone is known as a mineralocorticoid hormone involved in the regulation of electrolyte balance and volume homeostasis in epithelial cells [1]. By acting on kidney distal nephron, aldosterone promotes sodium reabsorption, water retention, and potassium and magnesium loss, modulating the extracellular space volume and blood pressure [2]. The epithelial cells of the kidney express the mineralocorticoid receptor (MR) a nuclear receptor that after binding to aldosterone translocates to the cell nucleus and modulates expression of specific "aldosterone-induced" proteins that regulate electrolyte and fluid balance [3].

The main regulator of aldosterone secretion is the reninangiotensin system [4]. Renin is synthesized by the juxtaglomerular cells of the kidney in response to low circulating blood volume, decreased renal perfusion, or reduced tubular sodium chloride concentration. Renin catalyzes the conversion of angiotensinogen to angiotensin I (Ang I); this is, in turn, converted to angiotensin II (Ang II) by the angiotensin-converting enzyme. Ang II induces the synthesis of aldosterone by the zona glomerulosa in the adrenal cortex through to AT1 receptor. Potassium levels and the adrenocorticotropic hormone can also regulate aldosterone levels [5].

Once aldosterone is produced and secreted on epithelial cells from the renal tubule, vascular smooth muscle cells, or adipose tissue [6, 7], it exerts most of its biological activities through binding to the MR, a member of the steroid receptor family [8]. After binding to the MR, the complex dimerizes and translocates to the nucleus where it induces the expression of genes related to water and salt regulation, such as epithelial 3 sodium channel (ENaC), sodium-potassium ATPase, and serum/glucocorticoid regulated kinase 1 (SGK1). The main goal of these processes is to maintain the body blood pressure in a normal range by means of water and electrolyte homeostasis control [9, 10].



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During the past years, studies showing the expression of MR in tissues such as the heart [11], blood vessels [12], brain [13], and immune cells [14–17] have extended the view of aldosterone acting beyond epithelial tissues. Thus, aldosterone is not only linked to systemic inflammation, endothelial dysfunction, increased vascular stiffness, hypertension, and cardiac hypertrophy but also to impaired pancreatic β -cell function, skeletal muscle insulin sensitivity, liver deficiency, and increased release of pro-inflammatory cytokines from the adipose tissue leading to impaired glucose tolerance and dyslipidemia [18–20].

Aldosterone and Obesity

Aldosterone has been linked with obesity and metabolic syndrome (MetS) that is a combination of interrelated risk factors of metabolic origin, including arterial hypertension, dyslipidemias, glucose intolerance, abdominal obesity, and a pro-inflammatory and prothrombotic state [21]. Patients with central obesity have higher levels of aldosterone, which have been associated with the development of MetS. Recently, aldosterone dysregulation was associated with MetS and cardiometabolic risk factors [22•]. On the other hand, urinary aldosterone, as a proxy of aldosterone daily production, has been linked with central obesity and higher adipose tissue [23].

Primary aldosteronism (PA) showed a high prevalence of MetS proposing a negative effect of aldosterone excess on glucose metabolism, suggesting that higher rates of cardiovascular events in PA than in essential hypertensives might also be secondary to an increase of the MetS in the former condition [24]. Moreover, The Framingham Offspring Study demonstrated an association of aldosterone levels with the development of MetS and with the longitudinal change of its components, suggesting that aldosterone may play a key role in mediating metabolic risk [25]. Observations that aldosterone might have detrimental effects on glucose metabolism and/or insulin action were focused on the similarities between the action of aldosterone and glucocorticoids and in the role of hypokalemia as a link between aldosterone and insulin secretion [26-30]. Mosso et al. showed that PA patients have lower HOMA-bF index and C-peptide levels and a negative correlation between HOMAbF and ARR, suggesting a decrease in insulin secretion or production secondary to aldosterone excess. The significant increases of the C-peptide and insulin after spironolactone treatment support the hypothesis that this effect could be mediated by aldosterone [31].

Aldosterone plasma levels have been shown to be associated with insulin resistance (IR), characterized by a reduced ability of insulin to inhibit the production of glucose from the liver and to promote glucose uptake in adipose tissue and skeletal muscle in normotensive healthy subjects independent of traditional risk factors [32]. Although the pathophysiological determinants linking aldosterone excess and IR are still largely unknown, it is known that PA patients display an increased incidence of IR mediated by the decreased gene expression of lipid metabolism genes phosphoenolpyruvate carboxykinase, perilipin, adiponectin, and peroxisome proliferator-activated receptors (PPAR γ) in the visceral adipose tissue (VAT) compared to age-, sex-, and BMI-matched controls [33].

Excess adiposity has been associated with elevated urinary aldosterone levels [23], as well as higher plasma aldosterone has been associated with obese subjects' essential hypertension [34]. Engeli et al. demonstrated that 70 % of obese women with weight loss showed a reduction of at least one of the renin-angiotensin-aldosterone system (RAAS) component [35]. Further, animal studies have also shown that MR expression is increased in obesity, which suggests that in this disease there is a combination of higher ligand and receptor overexpression in its pathophysiology [36•].

Today it is known that components of the RAAS are produced by adipocytes [36•, 37]. In addition to angiotensin II (Ang II), adipocytes secrete mineralocorticoid-releasing factors that stimulate steroidogenesis in human adrenocortical cells which might explain the higher aldosterone levels often observed in obese subjects [36•, 38-40]. Fatty acid oxidation products, or endogenous ones from adipocytes, could stimulate aldosterone synthesis [41]. The peroxidation product of linoleic acid, 12,13-epoxy-9-oxo-10 (trans)-octadecanoic acid (EKODE), stimulates the production of aldosterone by isolated rat zona glomerulosa cells, suggesting that free fatty acids could stimulate synthesis and release of aldosterone in adipocytes independently of angiotensin II (Fig. 1). Interestingly, in humans, EKODE plasma levels correlated with aldosterone plasma levels [41], while a positive correlation between aldosterone and the adipocyte-derived hormone leptin has recently been established [42].

Recently, Huby et al. described that leptin is a direct regulator of aldosterone secretion, regulating aldosterone synthase (CYP11B2) expression and production and promoting endothelial dysfunction and cardiac fibrosis [43•]. On the other hand, aldosterone synthase-deficient mice (As (-/-)) fed on a high-fat diet display better profile in insulin secretion, adipocyte size, and fatty liver than wild type. These results provide further evidence for a novel role of aldosterone in obesity-induced beta cell dysfunction, hepatic steatosis, and adipose inflammation [44].

In addition to the well-known effect of sodium and potassium on aldosterone regulation, new mechanisms include inadequate aldosterone feedback of adipocyte-releasing factors and alternative MR modulators such as caveolin-1 or Rac1 [45, 46]. With this new information regarding the complexity



Fig. 1 Aldosterone signaling in the adipocyte. In the adipocyte, aldosterone income from both the adrenal gland and in situ generated by the adipocyte by its own RAAS axis machinery may signal through a variety of genomic and non-genomic mechanisms and molecular pathways affecting adipose physiology. Aldosterone may act through a non-genomic pathway by binding to the membrane-bound GPR30 receptor or by inducing oxidative stress through Rac1 within or independently of the NOX complex. Otherwise, aldosterone may act through a genomic pathway by ligand-independent Rac1-mediated MR activation or by classical aldosterone-mediated MR activation along with transcription factors induced by non-genomic pathways induction and promote gene transcription of cytokines and adipokines, which can be secreted and induce adipocyte dysfunction. Aldosterone-releasing factors derived from these adipocytes, such as leptin and free fatty acids, have been shown to modulate aldosterone synthesis in the adrenal gland

of the aldosterone/MR pathway, is not surprising that contemporary environmental factors such as high sodium intake and some prevalent non-communicable diseases such as obesity can disturb the established aldosterone homeostasis, resulting in secretory and signaling dysregulation leading to several cardiometabolic disorders.

Aldosterone in the Adipocyte Function and Differentiation

The action of MR on adipose tissue was reported by Zennaro et al. in 1998, using a transgenic mice model that overexpress MR promoter activity leading to liposarcoma development, thus demonstrating that MR was transcriptionally active in adipocytes [47]. Then, the effect of aldosterone/MR on adipose tissue identified an important role in T37i cells differentiation. This cells in the presence of aldosterone showed a white rather than brown phenotype with accumulation of intracytoplasmic lipid droplets, intracellular triglyceride induction, and inhibition of UCP1 expression and thermogenesis, effects that were reverted by MR antagonists but not by GR antagonist [48]. The treatment with aldosterone in this cell line also enhanced early adipogenic gene markers, confirming the activation of the differentiation process [49].

The role of MR and aldosterone in white fat differentiation was also explored by Caprio et al. [50], where they showed that aldosterone was able to induce white adipose differentiation in 3T3-L1, promoting intracellular accumulation of lipid droplets and induction of PPAR γ and adipokines such as adiponectin, leptin, and resistin; these effects were reversed by the MR blocker spironolactone. Moreover, individuals with hypertrophic adipocytes have elevated pro-inflammatory factors including IL-6, IL-8, and monocyte-chemotactic-protein-1 (MCP-1) [51] and reduced adiponectin and IL-10 [51–53].

Studies in animal models has demonstrated that the activation of MR contributes to the inflammatory state in obesity increasing pro-inflammatory cytokines gene expression such as tumor necrosis factor- α (TNF- α), MCP-1, and prothrombotic factor plasminogen activator inhibitor type 1 (PAI-1), compared with lean, wild-type mice effect, and this phenotype was prevented by eplerenone [54]. In addition, eplerenone significantly reduced macrophage infiltration and reactive oxygen species production in adipose tissue [55]. Taken together, these results suggest that MR activation contributes to the changes in adipose tissue that promote low-grade inflammation.

Recent findings have also emphasized the interplay existing between gluco- and mineralocorticoid hormones and their respective receptors in the modulation of adipose tissue biology, an active topic of research that is beyond the scope of this review [56, 57•]. To date, adipose tissue function had not been studied in MR knockout mice neither in the knockdown MR transgenic rodents, since these models have been focused on MR activation in epithelial tissues [58–62].

Non-genomic and Epigenetic Modulation of Aldosterone Signaling in Adipocytes

Steroid hormone receptors also induce rapid extranuclear signaling effects including the activation of kinase signaling cascades and increases in second messenger production without transcriptional activity. Aldosterone binds to the MR and also mediates rapid non-genomic effects through the transactivation of the epidermal growth factor receptor (EGFR), through phosphorylation signaling by the mitogenactivated protein kinase (MAPK) cascades [63], as demonstrated in renal, cardiovascular, colonic, and brain tissues through the sensitivity of these responses to MR antagonists such as spironolactone or eplerenone [64].

Thus aldosterone, through genomic and non-genomic actions, contributes to induce several abnormalities beyond classic renal and vascular effects such as impaired beta cell function, as well as reduced skeletal muscle and adipose tissue insulin sensitivity [64-66]. Many of the rapid non-genomic actions of aldosterone are also mediated by GPR30 (or GPER-1), a G-protein coupled receptor [67] (Fig. 1). In human adrenal cortex and aldosterone-producing adenoma cells, GPR30 is the second most expressed receptor [68]. Interestingly, GPR30 deficiency in male mice results in insulin resistance, dyslipidemia, and a pro-inflammatory state [69]. The activation of GPR30 by the natural ligand 17^β-estradiol (E2), and the specific agonist G1, was shown to inhibit lipid accumulation in 3T3-L1 cells, while such inhibition was reversed upon knockdown of GPR30 using specific siRNA or GPR30 antagonist [70].

Obesity has emerged in the last years as a potential contributor for the epigenetic regulation of gene expression. Epigenetic modifications included DNA methylation, posttranslational histone modifications, and non-coding RNA-mediated silencing processes [71, 72]. These mechanisms, together with other transcriptional regulatory events, ultimately regulate gene activity and expression during development and differentiation or in response to nutritional and environmental stimuli [72]. However, it is unknown whether obesity affects the epigenetic regulation of different genes involved in adipose tissue dysfunction through aldosterone and/or MR activation.

Recently, Demura et al. showed that angiotensinogen (AGT) gene is an epigenetic-regulated gene in adipocytes, where DNA methylation negatively regulates AGT expression, which dynamically changes in response to continuous AGT promoter stimulation [73•]. High-salt intake and excess circulating aldosterone cause DNA demethylation around the CCAAT enhancer-binding protein-binding sites, thereby converting the phenotype of AGT expression from an inactive to an active state in visceral adipose tissue [74]. Evidences of methylation were also found in key genes regulating aldosterone biosynthesis (i.e., CYP11B2, AVPR1a, and PRKCA). These genes that regulate steroidogenic signals and synthesis in adrenocortical cells were differentially methylated: AVPR1a and PRKCA were downregulated and hypermethylated, and CYP11B2 was upregulated and hypomethylated [75].

On the other hand, microRNA (miRNA) regulation has been found in genes regulating RAAS. In 2013, Robertson et al. showed that aldosterone synthase (CYP11B2) 3'UTR is regulated by miR-24, miR-125a-5p, and miR-125b in adrenal cells [76]. Similarly, post-transcriptional regulation of MR gene expression in kidney is modulated by miR-124 and miR-135a [77]. In a rodent and human vascular smooth muscle cells (VSMC), the expression of AT1R is downregulated by miR-483-3p, which is a potential negative regulator of steadystate levels of RAAS acting directly over 3'UTR region of four different RAAS genes [78]. All these recent evidences support the potential epigenetic regulation by methylation CpG or miRNA of RAAS genes in adipose tissue and encourage further studies in this field.

Targeting Aldosterone Production and Signaling Pathways in Adipocyte

Despite the fact that obesity is a worldwide public health problem, current antiobesity therapy has only modest effects in body weight and associated cardiometabolic disorders [79–81]. Thus, targeting the pathogenic role of aldosterone signaling in adipose tissue is an interesting strategy. Considering the integrated pathways between different organs as well as the autocrine/paracrine action of these compounds on adipocyte biology, the adipocyte RAAS system regulation, aldosterone secretion, MR blockade, and adipokine receptors modulation emerge as targets to unravel the molecular mechanism of aldosterone action in adipocytes and thus identify novel therapeutic approaches.

As explained above, fat cells appear to stimulate aldosterone secretion, but aldosterone also appears to mediate adipocyte differentiation, and the increase in adipocyte mass may generate an autocrine-positive feedback loop for RAAS components [82, 83]. Several angiotensin-converting enzyme inhibitors (ACEi) and angiotensin receptor blockers (ARBs) reduce the secretion of aldosterone or increase adiponectin levels [84-86]. However, despite treatment with ACEi or ARBs, the plasma aldosterone level of classes 2 and 3 obese patients was higher than non-obese and class 1 obese patients probably because of aldosterone "breakthrough" [87, 88]. Another RAAS system player, Ang 1–7, had an ameliorating effect in obesity, adiposity, and myogenic and adipogenic differentiation [89, 90]. On the other hand, aldosterone synthase (CYP11B2) inhibition reduces aldosterone levels. However, inhibition of the highly similar enzyme CYP11B1, responsible for the production of cortisol, must be avoided because of the resulting impairment of the cortisol-induced stress response [91•]. Nextgeneration CYP11B2 inhibitors are currently under development, showing improved selectivity index and reduction of aldosterone levels in humans [92]. The recently solved crystal structure of human CYP11B2 provides a structural basis for the design of novel aldosterone synthase inhibitors, and the development of suitable surrogates for the human enzymes will provide superior in vitro assay systems for bioactivity evaluation [93•, 94].

In relation to aldosterone/MR pathway in mice, interestingly MR antagonists have shown to control HF diet-induced impairment in glucose tolerance and prevent body weight gain and white fat expansion [95]. However, in a small doubleblind, placebo-controlled, randomized clinical study in rather healthy obese non-diabetic (NCT01406015), spironolactone 50 mg daily versus placebo for 6 weeks showed only decreased median systolic blood pressure with no effect on other metabolic disorders [96••]. Thus, future studies in selected high-risk subjects with confirmed aldosterone dysregulation are warranted.

Finally, modulating adipokines that have a pathogenic role in obesity is a novel pharmacologic strategy. Enhanced leptin signaling elevated CYP11B2 expression and plasma aldosterone levels. The advancement into the structural and mechanistic aspects of leptin receptor (LR) activation allows a rational design of leptin receptors antagonists that may block obesity-induced increases in CYP11B2 and aldosterone [97, 98]. On the other hand, adiponectin secreted from adipocytes binds to adiponectin receptors AdipoR1 and AdipoR2 and exerts beneficial metabolic effects via activation of AMPK and PPAR- α pathways, respectively [99, 100]. The recently reported crystal structure of human AdipoR1 and AdipoR2 receptors provides a basis for the identification of novel adiponectin receptor agonists, that will help understanding the roles and mechanisms of the AdipoR1/ AdipoR2 in lipid metabolism [101•]. Further studies will show the potential of these approaches to ameliorate obesityrelated disorders.

Conclusions

In the past decades, we have extended the view of aldosterone effects beyond epithelial tissues. New evidence regarding the aldosterone/MR pathway in active metabolic tissues, including adipose tissue, has confirmed its pathogenic role in systemic inflammation, endothelial dysfunction, insulin resistance, and dyslipidemia, which are clustered in obesity.

We have recently learned that adipocytes express MR and also secrete aldosterone. Thus, both in vitro and in vivo studies have demonstrated a role for the aldosterone and MR activation in adipose tissue differentiation and enhanced early adipogenic gene markers. Also, current knowledge have deciphered that obesity- associated metabolic disorders may be explained, at least in part, by novel non-genomic activation of MR, new inducers of aldosterone synthesis, such as leptin, and probably by several epigenetic modifications. With the understanding of this complex interplay of obesity, hormones, and receptors, novel targeted pharmacological therapy is under active research.

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Compliance with Ethical Standards

Conflict of Interest Dr. Carvajal reports grants and personal fees from Fondecyt No. 1130427, from Fondecyt No. 1150437, from IMII P09/016-F ICM, and from CORFO 13CTI-21526-P1.

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