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Inflammation in Polycystic Ovary Syndrome: Underpinning of insulin resistance and ovarian dysfunction

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Abstract

Chronic low-grade inflammation has emerged as a key contributor to the pathogenesis of Polycystic Ovary Syndrome (PCOS). A dietary trigger such as glucose is capable of inciting oxidative stress and an inflammatory response from mononuclear cells (MNC) of women with PCOS, and this phenomenon is independent of obesity. This is important because MNC-derived macrophages are the primary source of cytokine production in excess adipose tissue, and also promote adipocyte cytokine production in a paracrine fashion.

The proinflammatory cytokine tumor necrosis factor- α (TNF α) is a known mediator of insulin resistance. Glucose-stimulated TNFα release from MNC along with molecular markers of inflammation are associated with insulin resistance in PCOS. Hyperandrogenism is capable of activating MNC in the fasting state, thereby increasing MNC sensitivity to glucose; and this may be a potential mechanism for promoting diet-induced inflammation in PCOS.

Increased abdominal adiposity is prevalent across all weight classes in PCOS, and this inflamed adipose tissue contributes to the inflammatory load in the disorder. Nevertheless, glucose ingestion incites oxidative stress in normal weight women with PCOS even in the absence of increased abdominal adiposity.

In PCOS, markers of oxidative stress and inflammation are highly correlated with circulating androgens. Chronic suppression of ovarian androgen production does not ameliorate inflammation in normal weight women with the disorder. Furthermore, *in vitro* studies have demonstrated the ability of pro-inflammatory stimuli to upregulate the ovarian theca cell steroidogenic enzyme responsible for androgen production. These findings support the contention that inflammation directly stimulates the polycystic ovary to produce androgens.

Keywords

Hyperandrogenism; inflammation; oxidative stress; insulin resistance; ovarian dysfunction; abdominal adiposity

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Introduction

Polycystic Ovary Syndrome (PCOS) is characterized by hyperandrogenism, chronic oligoor anovulation and polycystic ovaries [1]. Hyperandrogenism in particular, is a hallmark feature of PCOS because it is strongly implicated in the genesis of the disorder [2]; and is also associated with metabolic derangements that contribute to the underlying pathophysiology [3]. Consequently, the AE-PCOS Society maintains that the presence of hyperandrogenism is required to diagnose PCOS in consort with either chronic oligo- or anovulation or the presence of polycystic ovarios [4].

It is now clear that PCOS is a proinflammatory state, and emerging data suggests that chronic low-grade inflammation underpins the development of metabolic aberration and ovarian dysfunction in the disorder [5,6]. Most importantly, there is a strong association between hyperandrogenism and inflammation in PCOS that has been the focus of ongoing investigation [5,7-10]. Novel data presented herin suggests that in PCOS, diet-induced inflammation may directly invoke hyperandrogenism.

Insulin resistance and defective insulin signaling in PCOS

Insulin resistance is a common feature of PCOS affecting 50-70% of women with the disorder. The compensatory hyperinsulinemia is considered to be a promoter of the hyperandrogenism and chronic oligo- or anovulation. Android obesity is evident in \sim 52%-64% of women with PCOS, and is independently associated with metabolic abnormalities such as insulin resistance [1,11]. Abdominal adiposity in particular is present in ~30% of normal weight women with PCOS [12].

Circulating levels of the proinflammatory cytokine tumor necrosis factor- α (TNF α) are elevated in obesity, and are also elevated in PCOS independent of obesity [13,14]. In fact, the discovery of TNFα elevations in PCOS served as the initial clue that PCOS is a proinflammatory state. In obesity-related diabetic syndromes, TNFα is a known mediator of insulin resistance by causing increased serine phosphorylation of insulin receptor substrate-1 (IRS-1) in insulin sensitive tissues [15]. This leads to decreased expression of GLUT 4, the insulin sensitive glucose transport proteín [16].

The insulin receptor in PCOS is genetically and functionally normal. Insulin resistance in PCOS is also caused by a post-receptor defect in insulin signaling with increased serine phosphorylation implicated as the cause of decreased insulin-stimulated IRS-1 activation and decreased GLUT 4 expression [17,18]. Thus, the ability of TNF α to stimulate increased serine phosphorylation makes it an ideal candidate for initiating these molecular events in PCOS.

Although insulin resistance *per se* is considered to be the responsible entity for hyperandrogenism in PCOS, this mechanism does not explain the hyperandrogenism evident in women with the disorder without insulin resistance and/or excess adiposity. Inflammation may be the common thread in the induction of insulin resistance that is related to PCOS *per se*, or to superimposed excess adiposity. This concept raises the possibility that inflammation may be capable of directly inducing hyperandrogenism in PCOS.

Chronic low-grade inflammation in PCOS

There is a genetic basis for the chronic low-grade inflammation observed in PCOS. Several proinflammatory genotypes including those that encode TNF-α, and the type 2 TNF receptor as well as interleukin-6 (IL-6) and its signal transducer are associated with PCOS [19-21].

The majority of studies addressing the status of chronic low-grade inflammation in PCOS have focused on the measurement of circulating C-reactive protein (CRP) using highsensitivity assays. CRP is an acute phase reactant produced by the liver following stimulation by IL-6, the endocrine cytokine originating from adipose tissue in this instance [22]; CRP is also directly produced by adipose tissue [23]. CRP levels >3 mg/L are equally predictive of a cardiovascular event compared to the ATP III criteria for metabolic syndrome [24]. CRP also plays a functional role by promoting the uptake of lipids into foamy macrophages within atherosclerotic plaques [25].

A recent meta-analysis revealed that CRP is the most reliable circulating marker of chronic low-grade inflammation in PCOS [26]. However, the CRP elevation in normal weight women with PCOS (\leq 3.0 mg/L) is still much less compared to the obese ($>$ 3.0 mg/L) regardless of whether or not they have PCOS [27,28]. Thus, CRP elevations attributable to PCOS are obscured in the presence of obesity, and are below the range to predict metabolic or cardiovascular risk. This suggests that in PCOS, a single static circulating marker may not be reflective of inflammation at the molecular level.

The role of diet-induced inflammation in PCOS

Circulating mononuclear cells (MNC) and MNC-derived macrophages in tissue produce proinflammatory cytokines such as TNF α and IL-6 [29]. While TNF α is a known mediator of insulin resistance, the impact of IL-6 on insulin resistance is variable [15,30,31]. Moreover, IL-6 is clearly involved in the promotion of atherogenesis [32].

Circulating mononuclear cells utilize glucose during glycolysis for mitochondrial respiration. Some glucose is diverted to the hexose monophosphate shunt to generate nicotinamide adenine dinucleotide phosphate (NADPH) [33]. Membrane-bound NADPH oxidase is activated by translocation of a cytosol component known as p47^{phox} to the cell membrane [34,35]. Oxidation of NADPH by NADPH oxidase generates superoxide, a reactive oxygen species (ROS) that induces oxidative stress [36]. This in turn activates the transcription factor, nuclear factor κB (NFκB) by its dissociation from the inhibitory proteín, inhibitory κB (IκB). Activated NFκB translocates to the nucleus to promote TNFα and IL-6 gene transcription [15,37].

In PCOS, glucose ingestion induces an inflammatory response as evidenced by increased ROS-related oxidative stress, and increased NFκB activation that are independent of obesity (Fig. 1A-B) [5,9]. The release of TNF α and IL-6 from circulating MNC is also altered in PCOS by glucose ingestion *in vivo*, and by glucose exposure *in vitro* [7,8]. In addition, these markers of oxidative stress and inflammation are associated with glucose-challenged measures of insulin sensitivity and/or fasting measures of insulin resistance [5,8,9]. Thus, diet-induced inflammation in PCOS culminates in proinflammatory signaling known to be involved in the development of insulin resistance and atherogenesis.

The influence of adipose tissue on inflammation in PCOS

The proinflammatory state of obesity contributes to the promotion of insulin resistance and atherogenesis when present in PCOS. Hypoxia-related adipocyte death in response to adipose tissue expansion promotes an influx of MNC into the stromal-vascular compartment [38]. These MNC alter morphologically to become resident macrophages. MNC-derived macrophages are the prime source of $TNF\alpha$ and IL-6 production in adipose tissue, and also stimulate cytokine production in adipocytes through paracrine mechanisms [39].

The expression of molecular markers of inflammation is similar in adipose tissue of overweight women regardless of PCOS status, and there is lower expression in normal

weight women with PCOS compared to overweight individuals [40]. Thus, the inflammatory load derived from adipose tissue in PCOS is in proportion to body mass, but is not uniquely greater compared to that of individuals without PCOS. Even in the absence of frank obesity, increased abdominal adiposity is prevalent across all weight classes in PCOS [12].

Until recently, it remained unclear whether increased abdominal adiposity was the cause of the proinflammatory state in normal weight women with PCOS. However, it is now known that markers of oxidative stress such as MNC-derived ROS generation and p47^{phox} protein content increase in response to glucose ingestion in normal weight women with PCOS without increased abdominal adiposity [41]. This population is also insulin resistant, and exhibits higher testosterone levels and lower CRP levels compared to normal weight women with PCOS who have increased abdominal adiposity. Nevertheless, markers of oxidative stress are still greater in normal weight women with PCOS who have increased abdominal adiposity. There are also associations between CRP and abdominal adiposity, and between markers of oxidative stress and circulating androgens in normal weight women with PCOS. Thus, glucose-stimulated oxidative stress is independent of increased abdominal adiposity in normal weight women with PCOS, but increased abdominal adiposity contributes to the inflammatory load in the disorder. In addition, testosterone production in PCOS is greater in the absence of increased abdominal adiposity while CRP elevations are mostly an adiposityrelated phenomenon.

The relationship between inflammation and hyperandrogenism in PCOS

Circulating and molecular markers of oxidative stress and inflammation are highly correlated with circulating androgens [5,7-10]. These findings raise the possibility that in PCOS, either hyperandrogenemia pre-activate MNC to account for the hyperglycemiainduced inflammation, or conversely that glucose-stimulated inflammation promotes ovarian androgen production in PCOS. There is data to support that both mechanisms may occur [42-44].

Induction of hyperandrogenism in normal weight ovulatory women

In PCOS, MNC are pre-activated as evidenced by increased ROS generation and activated NFκB in the fasting state [45-46]. This accounts for the increased MNC sensitivity to glucose ingestion in the disorder. In contrast, MNC of normal weight ovulatory women are not sensitive to hyperglycemia, and do not exhibit an inflammatory response to glucose ingestion [5,7-9]. Acute oral androgen administration raises circulating androgen levels in normal weight ovulatory women to the range observed in PCOS. In the process, ROSrelated oxidative stress, activated NF κ B and TNF α RNA content from MNC increase in the fasting state, and in response to glucose ingestion (Fig. 2A-B) [42,43]. Thus, hyperandrogenemia to the degree present in PCOS, promotes MNC activation and increases MNC sensitivity to glucose ingestion. This suggests that hyperandrogenism, the hallmark feature of PCOS, is the progenitor of diet-induced inflammation in the disorder.

Suppression of androgens in women with PCOS and ovulatory women

Alteration in circulating CRP reflects exacerbation or amelioration of inflammation in clinical practice making it a useful measurement of inflammatory load [47-49]. CRP and body weight increase in obese women with PCOS, but remain unaltered in normal weight women with PCOS in response to chronic gonadotropin releasing hormone agonist (GnRH) agonist-induced androgen suppression. CRP is also unaltered in obese ovulatory women, but decreases in lean ovulatory women without significant weight change in either group during similar treatment (Fig. 3A-B) [44].

The ability of elevated circulating androgens to promote lipolysis may be responsible for the rise in CRP following chronic GnRH agonist administration in obese women with PCOS [50]. Testosterone in particular, is known to stimulate catecholamine-induced hormone sensitive lipase activity, which in turn, limits adipose tissue expansion [51]. A decrease in lipolysis following androgen suppression to castrate levels may explain the progressive weight gain in obese women with PCOS that most likely represents expansion of the adipose tissue compartment during chronic of GnRH agonist administration. In fact, circulating free fatty acids (FFA) decline in obese women with PCOS during chronic GnRH agonist administration [43]. Subsequent increases in IL-6 production from inflamed adipose tissue could result in the progressive rise in CRP observed in these individuals during this period.

Circulating androgens have a limited effect on lipolysis in normal weight women with PCOS and obese ovulatory women. Moreover, catecholamine resistance of subcutaneous adipose tissue precluding adequate induction of hormone-sensitive lipase activity is well documented in these individuals [52]. This phenomenon can limit expansion of inflamed adipose tissue to explain their unaltered levels of CRP and FFA and lack of significant change in weight during GnRH agonist administration. Thus, the factors responsible for limitation of fat mass in normal weight women with PCOS and obese ovulatory women are not dependent on the release of control by circulating androgens. In contrast, the decline in CRP in normal weight ovulatory women following GnRH agonist-induced androgen suppression corroborates the studies showing an increase in inflammatory load following androgen administration in this population [42].

These data demonstrate that hyperandrogenism in PCOS exerts an anti-inflammatory effect when obesity is present, but does not promote inflammation in the disorder. These unique observations support the contention that androgens have a pleiotropic effect on inflammation dependent on the combination of PCOS and weight status present in a given individual.

Induction of androgen production capacity by inflammation

Inflammation may be the promoter of hyperandrogenism in the disorder. Infiltration of the ovary by MNC-derived macrophages has been previously demonstrated [53]. *In vitro* studies show that CYP17, the ovarian steroidogenic enzyme responsible for androgen production is upregulated by proinflammatory stimuli, and inhibited by anti-inflammatory agents such as resveratrol and statins [6,54]. TNFα is a proinflammatory cytokine capable of stimulating *in vitro* proliferation of androgen producing theca cells [55]. It is possible that MNC recruited into the polycystic ovary may cause a local inflammatory response that stimulates ovarian androgen production in women with PCOS.

Conclusions

In PCOS, a dietary trigger such as glucose is capable of inducing oxidative stress to stimulate an inflammatory response even in the absence of excess adiposity. Hyperandrogenism may be the progenitor of chronic low-grade inflammation. Diet-induced inflammation in particular may be the underpinning of insulin resistance in the disorder. Inflammation directly stimulates excess ovarian androgen production. Increased abdominal adiposity contributes to the inflammatory load in PCOS, and its development may be controlled by the severity of hyperandrogenism.

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In Polycystic Ovary Syndrome

- **•** a prooxidant, proinflammatory state exists that is independent of excess adiposity,
- **•** inflammation triggered by glucose ingestion is associated with insulin resistance,
- **•** hyperandrogenism may be the progenitor of diet-induced inflammation,
- **•** oxidative stress and inflammation promotes hyperandrogenism, and
- **•** superimposed excess adiposity augments the inflammatory load.

Figure 1.

(A) The change from baseline (%) in ROS generation from mononuclear cells (MNC) when fasting samples (pre) were compared to the samples collected 2 hours after glucose ingestion (post). * the percent (%) change in ROS generation in normal weight women with PCOS was greater than that of normal weight ovulatory controls, $P < 0.009$. \dagger the % change in ROS generation in obese women with PCOS was greater than that of normal weight ovulatory controls, $P < 0.003$. (B) Representative EMSA bands from the 4 study groups showing the change in quantity of NFκB in nuclear extracts from MNC when fasting samples (pre) were compared to the samples collected 2 hours after glucose ingestion (post). Densitometric quantitative analysis of intranuclear NFκB protein content in MNC.

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Compared to normal weight ovulatory controls, the % change in NFκB activation was significantly greater in obese ovulatory controls (*, P<0.03), in normal weight women with PCOS (†, P<0.006), and in obese women with PCOS (†, P<0.002). Adapted from González et al. [5,9], with permission. Copyright The Endocrine Society, 2006.

Figure 2.

(A) Representative EMSA bands from the two study groups showing the quantity of NFκB in nuclear extracts from mononuclear cells (MNC) in samples collected in the fasting state (0) and 2 hours post-glucose ingestion (2), before and after treatment with DHEA or placebo. Densitometric quantitative analysis comparing the change from baseline (%) in MNC-derived activated NFκB between the two study groups for fasting samples before and after (before versus after, 0) DHEA or placebo administration (left panel); and for fasting and 2 hour post-glucose ingestion samples for each OGTT (before, 0 versus 2; after, 0 versus 2) as a measure of the response to glucose challenge before and after DHEA or placebo administration (right panel). After DHEA administration, the % change in activated

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NFKB was significantly greater compared to placebo in the fasting state (*, P<0.04), and in response to glucose ingestion (†, P<0.005). (B) Comparison between groups of the change from baseline (%) in TNF α mRNA content in MNC for fasting samples before and after (before versus after, 0) DHEA or placebo administration (left panel); and for fasting and 2 hour post-glucose ingestion samples for each OGTT (before, 0 versus 2; after, 0 versus 2) as a measure of the response to glucose challenge before and after DHEA or placebo administration (right panel). Values are normalized to 28S rRNA expression. After DHEA administration, the percent (%) change in TNF α mRNA transcripts significantly increased compared to placebo in the fasting state $(*, P<0.05)$, and in response to glucose ingestion $(*,$ P<0.05). Adapted from González et al. [43], with permission. Copyright The American Physiological Society, 2011.

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Figure 3.

The incremental change (Δ) from baseline in (A) serum C-reactive protein (CRP) levels and (B) body weight after 3 and 6 months of gonadotropin-releasing hormone (GnRH) agonist administration. The incremental Δ in CRP was significantly (*, P<0.009) higher in obese women with PCOS compared to normal weight ovulatory controls after 3 and 6 months of GnRH agonist treatment; and compared to obese ovulatory controls $(\dagger, P<0.005)$ and normal weight ovulatory controls (\ddagger , P<0.007) after 6 months of treatment. The incremental Δ in body weight was significantly higher in obese women with PCOS compared to obese ovulatory controls (*, P<0.02) and normal weight ovulatory controls (†, P<0.04) after 3 and 6 months of GnRH agonist treatment. González et al. [44].