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# Obese Reproductive-Age Women Exhibit a Proatherogenic Inflammatory Response During Hyperglycemia

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# Abstract

**Objective**—The objective was to determine if physiological hyperglycemia induces a proatherogenic inflammatory response in mononuclear cells (MNCs) in obese reproductiveage women.

**Research Methods and Procedures**—Seven obese and 6 age-matched lean women (20 to 39 years of age) underwent a 2-hour 75-g oral glucose tolerance test. The release of interleukin-6 (IL-6) and interleukin-1 $\beta$  (IL-1 $\beta$ ) from MNCs cultured in the presence of lipopolysaccharide (LPS) was measured after isolation from blood samples drawn fasting and 2 hours after glucose ingestion. Reactive oxygen species (ROS) generation and intra-nuclear nuclear factor  $\kappa$ B (NF $\kappa$ B) from MNCs were quantified from the same blood samples. Insulin resistance was estimated by homeostasis model assessment of insulin resistance (HOMA-IR). Total body fat and truncal fat were determined by DXA.

**Results**—Obese women had a higher (p < 0.03) total body fat ( $42.2 \pm 1.1 \text{ vs. } 27.7 \pm 2.0\%$ ), truncal fat ( $42.1 \pm 1.2 \text{ vs. } 22.3 \pm 2.4\%$ ), and HOMA-IR ( $3.3 \pm 0.5 \text{ vs. } 1.8 \pm 0.2$ ). LPS-stimulated IL-6 release from MNCs was suppressed during hyperglycemia in lean subjects ( $1884 \pm 495 \text{ vs.} 638 \pm 435 \text{ pg/mL}$ , p < 0.05) but not in obese women ( $1184 \pm 387 \text{ vs. } 1403 \pm 498 \text{ pg/mL}$ ). There was a difference (p < 0.05) between groups in the hyperglycemia-induced MNC-mediated release of IL-6 ( $-1196 \pm 475 \text{ vs. } 219 \pm 175 \text{ pg/mL}$ ) and IL-1 $\beta$  ( $-79 \pm 43 \text{ vs. } 17 \pm 12 \text{ pg/mL}$ ). In addition, the obese group exhibited increased (p < 0.05) MNC-derived ROS generation ( $39.3 \pm 9.9 \text{ vs. } -1.0 \pm 12.8\%$ ) and intra-nuclear NF $\kappa$ B ( $9.4 \pm 7.3 \text{ vs. } -23.5 \pm 13.5\%$ ). Truncal fat was positively correlated with the MNC-derived IL-6 response ( $\rho = 0.58$ , p < 0.05) and intra-nuclear NF $\kappa$ B ( $\rho = 0.64$ , p < 0.05).

**Discussion**—These data suggest that obese reproductive-age women are unable to suppress proatherogenic inflammation during physiological hyperglycemia. Increased adiposity may be a significant contributor to this pro-inflammatory susceptibility.

# Keywords

inflammation; atherosclerosis; hyperglycemia; abdominal adiposity

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# Introduction

Obesity is a major risk factor for developing atherosclerosis and hyperglycemia (1,2). Obesity is also a pro-inflammatory state as evidenced by elevated plasma concentrations of interleukin-6 (IL-6),<sup>1</sup> interleukin-1 $\beta$  (IL-1 $\beta$ ), and C-reactive protein (CRP) (3-6). Circulating IL-6 and IL-1 $\beta$  promote atherosclerosis by inducing endothelial expression of adhesion molecules and chemokines that cause peripheral mononuclear cells (MNCs) to attach to the endothelial layer of the blood vessel wall, and to subsequently migrate into the vascular interstitium (7-10). Several studies have documented overexpression of IL-6 and IL-1 $\beta$  in adipose tissue when obesity or type 2 diabetes mellitus is present, with roughly 30% of circulating IL-6 originating from fat depots (3,6,11,12). IL-6 and, to a lesser extent, IL-1 $\beta$ also stimulate CRP synthesis (13-15). CRP is a major predictor of atherosclerosis in asymptomatic individuals and may also play a functional role by promoting the uptake of lipids into MNC-derived foamy macrophages within atherosclerotic plaques (16-18). Although the majority of circulating CRP is produced by the liver, recent studies have shown that adipose tissue is an additional source (19).

It has recently been shown that MNCs of the obese are activated in a pro-inflammatory state (20). This is important because MNCs are known to migrate into adipose tissue to activate adipocyte cytokine production (21,22). However, it is now clear that roughly one half of IL-6 produced in adipose tissue of the obese comes from MNC-derived macrophages present in the stromal-vascular compartment (22,23). MNCs exhibit increased reactive oxygen species (ROS) generation in response to hyperglycemia. This induces oxidative stress, which, in turn, activates nuclear factor  $\kappa$ B (NF $\kappa$ B), a pro-inflammatory transcription factor that promotes both IL-6 and IL-1 $\beta$  gene transcription (24-26).

In our previous studies of obesity-related insulin resistance, we evaluated tumor necrosis factor-alpha (TNF- $\alpha$ ), another pro-inflammatory cytokine that is a known mediator of insulin resistance. We found that in older men and in obese young women, lipopolysaccharide (LPS)-stimulated TNF- $\alpha$  release from MNC was altered in response to hyperglycemia, and that the increased abdominal adiposity of these individuals was related to the altered TNF- $\alpha$  response (27,28). However, the hyperglycemic response of MNC-derived proatherogenic cytokines has never been explored in obese young women.

Thus, we embarked on a study to determine the status of IL-6 and IL-1 $\beta$  release from MNC in response to hyperglycemia in asymptomatic obese women of reproductive age. We also examined plasma IL-6 and CRP levels, as well as MNC-derived ROS generation and

<sup>1</sup> Nonstandard abbrevia	ations:
IL-6	interleukin-6
IL-1β	interleukin-1β
CRP	C-reactive protein
MNC	mononuclear cell
ROS	reactive oxygen species
NFκB	nuclear factor <i>k</i> B
TNF-α	tumor necrosis factor-alpha
LPS	lipopolysaccharide
OGTT	oral glucose tolerance test
HOMA-IR	homeostasis model assessment of insulin resistance

activated intra-nuclear NF $\kappa$ B. We hypothesized that LPS-stimulated release of IL-6 and IL-1 $\beta$ , ROS generation, and intra-nuclear NF $\kappa$ B from MNC of obese reproductive-age women are altered in response to an oral glucose challenge, as compared with age-matched lean controls.

# **Research Methods and Procedures**

#### **Subjects**

Thirteen women (7 obese and 6 lean) between 20 and 39 years of age participated in the study. Obesity was defined as a BMI between 30 and 40 kg/m<sup>2</sup>. Lean subjects had a BMI between 18 and 25 kg/m<sup>2</sup>. All subjects were ovulatory, as evidenced by regular menses, and had a luteal phase serum progesterone level >5 ng/mL. All subjects were screened for diabetes, inflammatory illnesses, or endocrinopathies, and none was taking medications that would affect carbohydrate metabolics or immune function. Based on Adult Treatment Panel III guidelines to diagnose the metabolic syndrome, none of the subjects exhibited 3 or more of the following features: waist circumference >88 cm, plasma triglyceride ≥150 mg/dL, plasma high-density lipoprotein cholesterol <50 mg/dL, blood pressure ≥130 mm Hg systolic/≥ 85 mm Hg diastolic, and fasting glucose ≥110 mg/dL (29). None of the subjects was involved in any regular exercise program for at least 6 months before the time of testing. All of the subjects provided written informed consent in accordance with Institutional Review Board guidelines for the protection of human subjects.

# Study Design

All study subjects underwent an oral glucose tolerance test (OGTT) between days 5 and 8 after the onset of menses. Before the OGTT, they were provided with a healthy diet consisting of 50% carbohydrate, 35% fat, and 15% protein for 3 consecutive days (Days 1 to 3). The test was performed on the morning of Day 4 after an overnight fast of ~12 hours. All subjects also underwent body composition assessment on the same day the OGTT was performed.

#### OGTT

Baseline blood samples were drawn for glucose and insulin determination. A 75-g glucose beverage was subsequently ingested over 10 minutes. Blood samples were again drawn for glucose and insulin determination 2 hours after glucose ingestion. On completion of the test, subjects were fed a high carbohydrate snack. Plasma glucose concentrations were assayed immediately from the blood samples collected. Additional plasma was isolated from the same blood samples and stored at -80 °C until assayed for CRP, IL-6, and lipids. Glucose tolerance was assessed by the World Health Organization criteria with normal glucose tolerance defined as a 2-hour glucose-stimulated value <140 mg/dL (30). Insulin resistance was estimated by homeostasis model assessment of insulin resistance (HOMA-IR) using the following formula: fasting glucose (mM) × fasting insulin /22.5 (31).

#### **Body Composition Assessment**

Height without shoes was measured to the nearest 1.0 cm. Body weight was measured to the nearest 0.1 kg. Waist circumference was measured at the level of the umbilicus and used to estimate abdominal adiposity (32). In addition, all subjects underwent DXA to determine percentage of total body fat and percentage of truncal fat using the QDR 4500 Elite model scanner (Hologic, Inc., Waltham, MA). Truncal fat content was defined as the area between the dome of the diaphragm (cephalad limit) and the top of the greater trochanter (caudal limit) (33).

# MNC Culture

MNCs were isolated by Histopaque-1077 density gradient centrifugation from blood samples drawn while fasting and 2 hours after glucose ingestion (34). The cells were washed twice in pyrogen-free saline, resuspended in RPMI (0.3 mg/mL <sub>L</sub>-glutamine, 100 U/mL penicillin, 100  $\mu$ g/mL streptomycin) with TCH Serum Replacement (MD Biomedicals, Inc., Irvine, CA), and seeded in coated culture plates (VWR International, West Chester, PA) at a concentration of  $2.5 \times 10^6$  cells/mL. The cells were then incubated (95% humidity, 5% CO<sub>2</sub>, 37 °C) for 24 hours with LPS endotoxin (1 ng/mL). Cell supernatants were subsequently collected by centrifugation (10,000g for 2 minutes) and stored at -80 °C until analysis.

# **ROS Generation**

MNCs were isolated by density gradient centrifugation using polymorphonuclear cell isolation medium (Robbins Scientific Corp., Sunnyvale, CA) from blood samples drawn while fasting and 2 hours after glucose ingestion. The cells were washed twice with Hank's buffered saline solution, and reconstituted to a concentration of  $4 \times 10^5$  cells/mL in Hank's buffered saline solution. Respiratory burst activity of MNCs was measured by detection of superoxide radical via chemiluminescence. Duplicate cuvettes containing 500  $\mu$ L of MNC (400 cells/µL) were placed into a two-channel lumi-aggregometer (Chronolog Corporation, Haverton, PA). Fifteen microliters of 10 mM luminol followed by 1  $\mu$ L of 10 mM formylmethionyl leucine phenylalanine were added to each cuvette. Chemilumunescence was recorded in mV by computer software (Chronolog Aggrolink). This method was developed by Thusu and Dandona (35) and is similar to that published by Tosi and Hamedani (36). In this assay system, measurement of superoxide radical release has been shown to be linearly correlated with the ferricytochrome C method (37). In addition, superoxide dismutase, catalase, and diphenylene, a specific inhibitor of nicotinamide adenine dinucleotide phosphate oxidase, have been shown to inhibit chemiluminescence in a dose-dependent fashion. The specific inhibitory effect of diphenylene iodonium on nicotinamide adenine dinucleotide phosphate oxidase has been established by Hancock and Jones (38). As previously validated, the variation of ROS generation by MNCs in humans using this method varies by <8% over a 2-week period (39).

#### NFkB Electrophoretic Mobility Shift Assay

Nuclear extracts of DNA-binding protein were prepared from MNCs using the method described by Andrews and Faller (40). Total protein concentrations were determined using the bicinchoninic acid protein assay (Pierce Chemical Co., Rockville, IL). An NF $\kappa$ B gel retardation assay was performed using the NFKB-Binding Protein Detection Kit (Life Technologies, Inc., Long Island, NY). A double-stranded oligonucleotide containing a tandem repeat of the consensus sequence for the NF $\kappa$ B-binding site (Santa Cruz Biotechnology, Inc., Santa Cruz, CA) was radiolabeled with  $\gamma$ -P<sup>32</sup> (GE Healthcare Bio-Sciences Corporation, Piscataway, NJ) using T4 kinase (Invitrogen, Carlsbad, CA). Nuclear extract (7.5  $\mu$ g) was mixed with the incubation buffer, and the mixture was pre-incubated at 4 °C for 15 minutes. Labeled oligonucleotide (60,000 cpm) was added, and the mixture was incubated at room temperature for 20 minutes. The samples were electrophoresed on 6% non-denaturing polyacrylamide gels. The gels were dried under vacuum and exposed to X-ray film. Densitometry was performed using Kodak 1-Day Image Analysis software version 3.6 (Rochester, NY).

#### **Plasma Measurements**

Plasma glucose concentrations were measured by the glucose oxidase method (YSI, Yellow Springs, OH) while plasma insulin concentrations were measured by a double antibody radioimmunoassay (Linco Research, Inc., St. Charles, MO). Plasma CRP concentrations

were measured by a high sensitivity enzyme-linked immunosorbent assay (Alpha Diagnostics International, San Antonio, TX). The concentrations of IL-6 and IL-1 $\beta$  were also measured by enzyme-linked immunosorbent assay (eBioscience, San Diego, CA). Levels of total cholesterol, triglyceride, and high-density lipoprotein cholesterol were measured by enzymatic methods (SYNCHRON LX20 PRO automatic analyzer; Beckman Coulter, Inc., Fullerton, CA). Low-density lipoprotein cholesterol was calculated using Friedewald's formula (41). All samples from each subject were measured in duplicate in the same assay at the end of the study. The inter-assay and intra-assay coefficients of variation for all assays were 7% and 12%, respectively.

## Statistics

The StatView statistical package (SAS Institute, Inc., Cary, NC) was used for data analysis. The difference between the pre- and post-glucose challenge values for primary dependent variables, such as the release of IL-6 and IL-1 $\beta$  from MNCs, was calculated to represent the incremental change. The percentage change between the pre- and post-glucose challenge values was used to express alterations in ROS generation and nuclear-bound NF $\kappa$ B from MNCs due to the inter-individual variability. Descriptive data and the change of variables were compared between-groups using the unpaired Student's *t* test. Differences between pre- and post-glucose challenge variables within-groups were analyzed using the paired Student's *t* test. Regression analyses were performed using Pearson (r) correlation for parametric data and Spearman rank order ( $\rho$ ) correlation for non-parametric data. All values are expressed as means ± standard error. An  $\alpha$  level of 0.05 was used to determine statistical significance.

## Results

#### Body Composition, Glycemic Status, and Insulin Resistance

Age, height, and lipid levels were similar between groups, and all subjects were normotensive. The obese group had significantly higher (p < 0.003) weight, BMI, percentage of total body fat, percentage of truncal fat, and waist circumference (Table 1).

Levels of glucose while fasting and 2 hours post-glucose ingestion were similar in both groups (Table 2). All subjects had a normal glucose response during the OGTT with 2-hour glucose levels ranging between 62 and 138 mg/dL. However, the 2-hour glucose level in the obese group and the 2-hour insulin level in both groups were significantly increased (p < 0.02) compared with fasting levels. HOMA-IR was significantly greater (p < 0.05) in the obese group compared with the lean control group. HOMA-IR was positively correlated with percentage of total body fat (r = 0.67, p < 0.02), percentage of truncal fat (r = 0.67, p < 0.02), and waist circumference (r = 0.82, p < 0.004) for the combined groups (data not shown).

## IL-6 and IL-1β Release

LPS-stimulated IL-6 release from MNC in the fasting state was similar in both groups. However, fasting LPS-stimulated IL-1 $\beta$  release was significantly (p < 0.02) lower in the obese group. Hyperglycemia resulted in significant (p < 0.05) suppression of LPS-stimulated IL-6 release and modest suppression of IL-1 $\beta$  release from MNC of lean controls, but no change in either of these cytokine responses in obese women (Figure 1). In addition, the incremental change in the MNC-derived IL-6 and IL-1 $\beta$  response between the two groups was significantly different (p < 0.05).

There was a direct correlation between the incremental change in IL-6 release from MNC and that of IL-1 $\beta$  ( $\rho = 0.64$ , p < 0.04). There was also a direct relationship between the

MNC-derived incremental change in IL-6 release and percentage of total body fat, percentage of truncal fat, and waist circumference for the combined groups (Table 3). There was no correlation between the incremental change in IL-1 $\beta$  release from MNC and any of these body composition parameters.

#### Plasma CRP and IL-6, ROS Generation, and Intra-nuclear NFkB

Fasting plasma concentrations of CRP and IL-6 were significantly (p < 0.05) higher in the obese group but remained unchanged after glucose ingestion in both groups (Table 2). The percentage change in MNC-derived ROS generation and intra-nuclear NF $\kappa$ B in response to the oral glucose load was significantly (p < 0.05) higher in the obese group compared with that of lean controls (Figure 2). For the combined groups, there was a positive correlation between fasting plasma IL-6 and CRP. Each of these levels was positively correlated with BMI, percentage of total body fat, percentage of truncal fat and waist circumference. Plasma CRP was also positively correlated with ROS generation and intra-nuclear NF $\kappa$ B. HOMA-IR was positively correlated with fasting plasma IL-6 and CRP, and with intra-nuclear NF $\kappa$ B. There was also a positive correlation between ROS generation and BMI, and between intra-nuclear NF $\kappa$ B and percentage of truncal fat (Table 3).

# Discussion

Our data clearly show that obese reproductive-age women have impaired suppression of IL-6 and IL-1 $\beta$  release from MNC in response to physiological hyperglycemia. In contrast, the release of these cytokines from MNC was suppressed in lean controls of similar age under postprandial-like conditions. ROS generation, a promoter of oxidative stress; intranuclear NF $\kappa$ B, the cardinal signal of inflammation; and plasma IL-6 and CRP levels were also elevated in the obese group compared with lean controls. These findings in our study population provide further support for the role of inflammation in the development of atherosclerosis, and that in obese women, MNCs are likely to be involved in atherogenesis at an early age. Furthermore, the independent associations between multiple biomarkers of inflammation and abdominal adiposity suggest that the accumulation of abdominal fat is an important contributing factor in promoting atherogenesis in obese reproductive-age women.

Suppression of MNC-derived inflammation may be the normal in vivo response to physiological hyperglycemia. Lean controls in the present study showed decreases in LPSstimulated release of IL-6 and IL-1 $\beta$  from MNC in response to the oral glucose load. These results are similar to the hyperglycemia-induced TNF- $\alpha$  response that we have previously reported in lean young healthy men and women (27,28). Modest declines in MNC-derived ROS generation and intra-nuclear NF $\kappa$ B were also observed in lean controls. Our findings differ from those of previous studies that reported pro-inflammatory effects after an oral glucose challenge in normal subjects (42-44). The majority of subjects in these studies were men, which may account for the different results. Because insulin is an anti-inflammatory hormone (39,45), the postprandial insulin release in the more insulin sensitive lean controls may reverse the hyperglycemia-induced pro-inflammatory response. In this scenario, the suppressed inflammatory response may serve as a protective mechanism to preserve blood vessel integrity in the presence of hyperglycemia. Increased release of IL-6 and IL-1 $\beta$  after NFkB activation by ROS-induced oxidative stress may trigger MNCs to adhere to the endothelium and migrate into the vascular interstitium (10,24-26). Thus, young lean individuals may be capable of retarding atherogenesis by controlling the release of IL-6 and IL-1 $\beta$  in the postprandial state.

In contrast, the MNCs of obese reproductive-age women may have an impaired ability to down-regulate the inflammatory response to physiological hyperglycemia. Indeed, the elevations in plasma IL-6 and CRP observed in this group confirm previous reports

demonstrating that obesity is a pro-inflammatory state (5,6,46). The increases in ROS generation and intra-nuclear NFkB, and the failed suppression of MNC-derived IL-6 and IL-1 $\beta$  release in the obese group provide further corroboration, and is consistent with our previous report of failed suppression of TNF- $\alpha$  from MNCs in obese young women (28). It is paradoxical that IL-1 $\beta$  release in the fasting state is low in the obese group, raising the possibility that the study of MNC in vitro may not reflect physiology in vivo. The direct relationship of plasma CRP with ROS generation, intra-nuclear NFkB, and plasma IL-6 are consistent with previous investigations demonstrating that CRP production in the obese is the culmination of oxidative stress-induced ROS generation that activates NF $\kappa$ B to alter the transcription of IL-6 and IL-1 $\beta$  (24-26). Furthermore, obese reproductive-age women also exhibited evidence of insulin resistance based on an increase in HOMA-IR, a feature highly associated with atherosclerosis (47). This is confirmed in the present study by the direct relationship of HOMA-IR with intranuclear NFkB and plasma IL-6 and CRP levels. Further corroboration is provided by recent evidence that, in the obese, tyrosine phosphorylation of the insulin receptor is inversely related to pro-inflammatory mediators and HOMA-IR (48). Thus, our data suggest that chronically increased macronutrient intake leading to obesity at an early age results in an increase in oxidative stress and inflammation. A subsequent failure to suppress MNC-derived inflammation in the postprandial state may promote atherosclerosis. This concept is further supported by previous reports of a reduction in oxidative stress and inflammatory mediators after caloric restriction in the obese, and after a 2-day fast in normal subjects (49-51).

Our data suggest that there may be a link between obesity and MNC-derived inflammation that promotes atherosclerosis. There was a direct relationship between MNC-derived ROS generation, intra-nuclear NF $\kappa$ B and IL-6 release, as well as plasma IL-6 and CRP in response to physiological hyperglycemia, and measures of adiposity, particularly abdominal adiposity. Obese reproductive-age women with insulin resistance exhibited increased BMI, percentage of total body fat, percentage of truncal fat, and waist circumference. Because activated MNC-derived macrophages produce roughly one half of the IL-6 in the expanded adipose mass of obese individuals, it is possible that the inflamed adipose tissue of the obese, especially in the abdominal region, perpetuates the inability to suppress inflammation after hyperglycemia (22,25). It remains to be established whether failed suppression of inflammation is also evident in MNC-derived macrophages within the adipose tissue stromal-vascular compartment of obese individuals. Nevertheless, these data are striking because they suggest that obesity-related glucose or macronutrient-induced inflammation may initiate a proatherogenic milieu in women at an early age.

In summary, obese reproductive-age women are unable to suppress inflammation during physiological hyperglycemia. The resultant increases in MNC-derived ROS generation and intra-nuclear NF $\kappa$ B, failed suppression of LPS-stimulated IL-6 and IL-1 $\beta$  release from MNC, and increased plasma IL-6 and CRP levels compared with a lean control group demonstrates that inflammation can be a factor that places these individuals at an increased risk of developing atherosclerosis. The association between mediators of inflammation and both total and abdominal fat suggests that increased adiposity is a significant contributor to the proinflammatory state in obese young women. Thus, it is intriguing to consider the possibility that when glucose increases during the postprandial period in lean individuals, inflammation is suppressed to protect against atherogenesis. Conversely, the loss of this postprandial response may be one of the factors that contribute to the promotion of atherogenesis in obese women at an early age.

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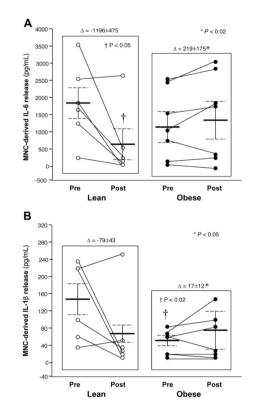
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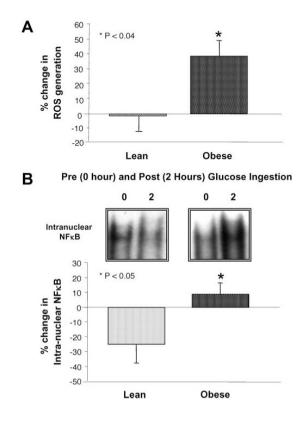
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#### Figure 1.

(A) Interleukin-6 (IL-6) release (pg/mL) from mononuclear cells cultured with lipopolysaccharide for 24 hours when fasting samples (pre) were compared with the samples collected 2 hours after glucose ingestion (post). \* Incremental change ( $\Delta$ ) in IL-6 release during oral glucose challenge in the obese group was significantly different from that of lean controls (p < 0.02). † Two hours post-glucose was significantly lower than fasting in the lean group (p < 0.05). (B) Interleukin-1 $\beta$  (IL-1 $\beta$ ) release (pg/mL) from mononuclear cells cultured with lipopolysaccharide for 24 hours when fasting samples (pre) were compared with the samples collected 2 hours after glucose ingestion (post). \* Incremental change in IL-1 $\beta$  release during oral glucose challenge in the obese group was significantly different from that of lean controls (p < 0.05). † Fasting was significantly lower than in the lean group (p < 0.02).

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### Figure 2.

(A) The percentage change in ROS generation from mononuclear cells (MNC) when fasting samples (pre) were compared with the samples collected 2 hours after glucose ingestion (post). \* ROS generation in the obese group was significantly greater compared with lean controls (p < 0.04). (B) Representative bands from the electrophoretic mobility shift assay showing the change in quantity of NF $\kappa$ B in nuclear extracts, and densitometric analysis of intra-nuclear NF $\kappa$ B protein content from MNC when fasting samples (pre) were compared with the samples collected 2 hours after glucose ingestion (post). \* The percentage change in intra-nuclear NF $\kappa$ B was significantly greater in the obese group compared with lean controls (p < 0.05).

### Table 1

Body composition, fasting lipid levels, and insulin resistance of subjects

	Lean ( <i>n</i> = 6)	<b>Obese</b> ( <i>n</i> = 7)
Age (yrs)	$33 \pm 2$	$29\pm3$
Systolic blood pressure (mm Hg)	$104 \pm 3$	$117\pm 6$
Diastolic blood pressure (mm Hg)	$56\pm3$	$76\pm4^*$
Height (cm)	$166 \pm 1.0$	$163\pm2.8$
Body weight (kg)	$59.1\pm2.2$	$92.1\pm 4.0^{\ast}$
BMI (kg/m2)	$21.6\pm0.8$	$34.8 \pm 1.2^{\ast}$
Total body fat (%)	$27.7\pm2.0$	$42.2\pm{1.1}^*$
Truncal fat (%)	$22.3\pm2.4$	$42.1\pm 1.0^{*}$
Waist circumference (cm)	$71.3\pm2.0$	$101.6\pm3.5^{*}$
Total cholesterol (mg/dL)	$164\pm7$	$201\pm23$
Triglycerides (mg/dL)	$55\pm 6$	$121\pm44$
HDL-cholesterol (mg/dL)	$52\pm3$	$50\pm3$
LDL-cholesterol (mg/dL)	$105\pm 6$	$121\pm19$

HDL, high-density lipoprotein; LDL, low-density lipoprotein. Values are mean  $\pm$  standard error.

\*Significantly different from lean group (p < 0.003).

# Table 2

Plasma glucose, insulin, IL-6, and C-reactive protein levels while fasting and in response to the oral glucose challenge, and HOMA-IR in lean and obese women

	Fasting	2 hours post-glucose
Glucose (1	mg/dL)	
Lean	$85.3\pm2.0$	$99.0 \pm 11.8$
Obese	$82.9\pm4.4$	$119.4 \pm 5.5^{*}$
Insulin (µ	U/mL)	
Lean	$8.5\pm1.2$	$33.1\pm8.8^\ast$
Obese	$14.0\pm2.3$	$88.2\pm24.2^*$
HOMA-II fastir (µU/r	ng insulin	
Lean	$1.8\pm0.02$	
Obese	$3.3\pm0.05^{\dagger}$	
Plasma IL (pg/n		
Lean	$0.68\pm0.06$	$0.72\pm0.11$
Obese	$2.26\pm0.61^{\dagger}$	$2.99\pm0.61^{\dagger}$
CRP (ng/n	nL)	
Lean	$219\pm92$	$221\pm99$
Obese	$7215\pm941^{\dagger}$	$6701 \pm 1001 ^{\dagger}$

 $IL-6, interleukin-6; CRP, C-reactive \ protein; HOMA-IR, homeostasis \ model \ assessment \ of \ insulin \ resistance. \ Values \ are \ mean \ \pm \ standard \ error.$ 

\*Two hours post-glucose significantly higher than fasting (p < 0.02).

<sup> $\dagger$ </sup>Significantly higher than the lean group (p < 0.05).

Table 3

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Spearman rank correlations for the combined groups

		Plasma IL-6 (pg/mL)	Plasma CRP (ng/mL)	ROS (% change)	NFkB (% change)	MNC IL-6 (pg/mL)	MNC IL-1/ (pg/mL)
HOMA-IR (mM) × fasting	φ	0.700	0.622	0.164	0.673	0.302	0.336
ınsulın (µ∪/mL)	р	$0.027^{*}$	$0.039^{*}$	0.605	0.044	0.295	0.266
BMI (kg/m <sup>2</sup> )	φ	0.811	0.665	0.655	0.164	0.516	0.238
	d	$0.007^{*}$	$0.021^{*}$	$0.039^{*}$	0.624	0.074	0.430
Total body fat (%)	θ	0.699	0.791	0.418	0.588	0.582	0.189
	d	$0.020^{*}$	$0.006^*$	0.186	0.078	$0.044^*$	0.531
Truncal fat (%)	φ	0.727	0.797	0.455	0.636	0.567	0.217
	р	$0.016^*$	$0.006^*$	0.151	$0.049^{*}$	$0.049^{*}$	0.472
Waist circumference (cm)	φ	0.733	0.618	0.217	0.333	0.700	0.491
	d	$0.028^*$	$0.049^{*}$	0.540	0.378	$0.027^{*}$	0.141
Plasma IL-6 (pg/mL)	φ	I	0.790	0.552	0.552	0.476	0.154
	d		$0.009^{*}$	0.098	0.098	0.115	0.610
Plasma CRP (ng/mL)	φ	0.790	Ι	0.627	0.758	0.495	0.126
	d	$0.009^{*}$	Ι	$0.047^{*}$	$0.023^{*}$	0.087	0.676

 $^{*}_{P < 0.05.}$