Adipocyte Death Defines Macrophage Localization and Function in Adipose Tissue of Obese Mice and Humans


*Institute of Normal Human Morphology and ** Endocrinology Unit, Univ. of Ancona, Ancona, Italy
#Division of Medical Genetics, Research Centre, Hopital Ste.-Justine, Montreal, Canada
‡Obesity and Metabolism Laboratory, JMUSDA-HNRCA at Tufts Univ., Boston, MA

Nonstandard Abbreviations:
CLS, crown-like structures
MGC, multinucleate giant cell
WAT, white adipose tissue

††Correspondence:
Dr. Saverio Cinti MD, University of Ancona (Politecnica delle Marche), Via Tronto 10A 60020 Ancona-Italy, Ph +39 071 2206088, email: cinti@univpm.it
Dr. Andrew S. Greenberg, JMUSDA-HNRCA at Tufts Univ., 711 Washington Street, Boston, MA 02111. Ph 617 556-3144, email: andrew.greenberg@tufts.edu
ABSTRACT. Macrophage infiltration of white adipose tissue (WAT) is implicated in the metabolic complications of obesity. The precipitating event(s) and function(s) of macrophage infiltration into WAT are unknown. We demonstrate that >90% of all macrophages in WAT of obese mice and humans are localized to dead adipocytes, where they fuse to form syncitia that sequester and scavenge the residual “free” adipocyte lipid droplet and ultimately form multinucleate giant cells, a hallmark of chronic inflammation. Adipocyte death increases in obese (db/db) mice (30-fold) and humans and exhibits ultrastructural features of necrosis (but not apoptosis). These observations identify necrotic-like adipocyte death as a pathologic hallmark of obesity and suggest that scavenging of adipocyte debris is an important function of WAT macrophages in obese individuals. The frequency of adipocyte death is positively correlated with increased adipocyte size in obese mice and humans and in hormone sensitive lipase-deficient (HSL-/-) mice, a model of adipocyte hypertrophy without increased adipose mass. WAT of HSL-/- mice exhibited 15-fold increase in necrotic-like adipocyte death and formation of macrophage syncitia, coincident with elevated TNF-α gene expression. These results provide a novel framework for understanding macrophage recruitment, function and persistence in WAT of obese individuals.

Key Words: obesity, macrophage, inflammation, apoptosis, necrosis, adipocyte hypertrophy, multinucleate giant cell, insulin resistance, type 2 diabetes.
INTRODUCTION. Adipose tissue inflammation is now recognized as an important early event in the development of obesity complications, especially type 2 diabetes (1-7). Recent studies suggest that adipose tissue inflammation is due in large part to the pro-inflammatory actions of bone marrow-derived WAT macrophages. These studies demonstrate that 1) macrophage numbers and/or macrophage inflammatory gene expression in WAT are positively correlated with adipocyte size and BMI in mice and negatively correlated with weight loss in obese humans (1, 5, 8); 2) macrophages are the predominant source of TNF-α and a significant source of IL-6 and NO in WAT of obese (ob/ob, db/db) mice and humans (2, 5, 7, 9); 3) a “spike” in macrophage inflammatory gene expression in WAT immediately precedes or is coincident with the onset of hyperinsulinemia in murine diet-induced obesity (7); and 4) obese mice with genetically ablated macrophage inflammatory (i.e., NF-κB) signaling are protected from insulin resistance (10). Together, these observations implicate macrophage activation in the development of obesity-associated WAT inflammation and insulin resistance.

Despite these advances, we have yet to elucidate the underlying cause and function of macrophage infiltration into WAT of obese subjects. Macrophages are monocytic phagocytes that function in innate immunity and wound-healing by sequestering and clearing pathogens, dead cells and cell debris in an activation-dependent manner (11). Macrophage activation consists of biochemical, morphological and functional changes that result in the secretion of preformed and/or newly-synthesized constituents, such as cytokines and chemokines. These, in turn, switch on the inflammatory response (11). Macrophage activation at sites of inflammation is typically transient, giving way to repair processes that re-establish local tissue function. However, at sites of resistant infectious agents (i.e. tuberculosis) or poorly biodegradable tissue irritants (i.e. foreign bodies), macrophages remain activated and fuse to form multinucleated giant cells (MGCs) that can persist for weeks or months surrounding the unresolved site. At these sites, MGCs actively phagocytose debris and can acutely produce
pro-inflammatory cytokines (i.e. IL-1α, TNF-α) until the insult is either cleared by phagocytosis or encapsulated (12).

MGCs have been reported surrounding individual adipocytes in WAT of obese (but not lean) mice and humans (1, 5, 7), suggesting that in the obese state, individual adipocytes become focal and persistent sites of macrophage activation. Macrophage infiltration and MGC formation in adipose tissue are also observed in conjunction with adipocyte (“fat”) necrosis, which occurs in various human diseases (i.e. Weber-Christian disease, Rothmann-Makai syndrome, pancreatic panniculitis) (13) and follows failure of autologous adipose implants (14). These observations suggest the potential involvement of adipocyte cell death in macrophage recruitment and activation in WAT of obese individuals.

The present work demonstrates that 1) the overwhelming predominance (>90%) of WAT macrophages in obese mice and humans are localized selectively to sites of necrotic-like adipocyte death, where they form syncitia that sequester and ingest adipocyte debris, in particular the residual lipid droplet, and 2) the frequency of adipocyte death is dramatically elevated in obese mice and humans and in a mouse model of adipocyte hypertrophy without obesity. These results identify adipocyte death as an important modulator of obesity-associated macrophage responses in WAT.
MATERIALS AND METHODS

**Mice.** All animal procedures were in accordance with National Institute of Medical Research (INSERM) guidelines. *db/db* and C57BL/6 mice were purchased from Charles River Italy at five weeks of age and used for experimental procedures at 12 weeks of age. Mice were individually caged and maintained on a 12:12 light/dark cycle with free access to standard pellet food (Harlan, Italy) and water. Obese male *db/db* mice (n=5), and lean wild-type (C57BL/6) control mice (n=5) were used to investigate obesity-associated adipocyte necrosis. The generation of HSL/-/- mice has been previously described (15). HSL/-/- mice were backcrossed to C57BL/6 mice for 8 generations and used at 8 weeks of age for gene expression analysis and 12 weeks of age for histological analysis.

**Human adipose tissue.** Fat biopsies were obtained from male (n = 12) and female (n = 9) patients aged 44.7 ± 16.2 (mean ± SD) years undergoing institutionally approved elective surgery at Ancona General Hospital (Ancona, Italy). Fully informed and written consent was obtained in all cases. The subjects did not suffer from any ongoing disease (infection or cancer). A total of 28 human biopsies from subcutaneous (abdominal and gluteal) and visceral (omental) depots were examined from 12 obese patients (BMI ≥30, 5 male, 7 female), 2 overweight patients (BMI 25-29.9, 1 male, 1 female), and 7 lean subjects (BMI 20-24.9, 5 male, 2 female). Mean (± SD) BMI of male subjects (31.3 ± 8.7) was not significantly different from mean BMI of female subjects (35.1 ± 9.1) (p = 0.39).

**Light microscopy.** Mice were sacrificed in the fed state with an overdose of anesthetic (xilazine-ketamine) and immediately perfused with a 4% paraformaldehyde in 0.1M phosphate buffer, pH 7.4 for 2-5 min. Subcutaneous and visceral WAT depots were dissected under a
Zeiss OPM1 surgical microscope (Carl Zeiss, Germany) and assessed by light microscopy, transmission electron microscopy, immunohistochemistry and morphometry. The following WAT depots were examined: subcutaneous (inguinal) and visceral (perivesicular) WAT depots in db/db and lean control mice and visceral (perirenal) depots in HSL-/− and HSL+/+ mice. After dissection, WAT depots were further fixed by immersion in the perfusion fixative (overnight, 4°C), dehydrated, cleared and then embedded in paraffin. Five µm-thick serial sections were obtained, the first stained by haematoxylin/eosin to assess morphology, the following processed for immunohistochemistry (see below).

**Electron Microscopy.** Small fragments of tissue were fixed in 2% glutaraldehyde-2% paraformaldehyde in 0.1 M phosphate buffer pH 7.4 for 4 h, postfixed in 1% osmium tetroxide and embedded in an Epon-Araldite mixture. Semi-thin sections (2 µm) were stained with toluidine blue, and thin sections were obtained with an MT-X ultratome (RMC, Tucson, Arizona), stained with lead citrate and examined with a CM10 transmission electron microscope (Philips, Eindhoven, Netherlands).

**Immunohistochemistry.** Immunohistochemistry was performed on de-waxed 5 µm serial sections, using 3% hydrogen peroxide to inactivate endogenous peroxidases followed by normal goat or horse serum to reduce non-specific staining. Consecutive serial sections were incubated overnight (4°C) with the following primary antibodies: anti-MAC-2/galectin-3 (1:3800, Cedarlane Laboratories), anti-mouse F4/80 (1:100, Serotec, Oxford), anti-perilipin (“PREK” 1:300) (16), anti-S-100B (1:250, Dako, Denmark), and a polyclonal antibody raised against a C-terminal peptide (TEVKNKASLKCVQEVKAQ) of mouse ADRP (1:100). Primary antibody was omitted from negative controls. Appropriate positive controls where also used to test specificity of all antibodies. Biotinylated, HRP-conjugated second antibodies included goat anti-rabbit IgG (ADRP-C; PREK; S100B), rat anti-goat IgG (F4/80) and horse anti-mouse IgG (MAC2/galectin-
3) (Vector Laboratories; Burlingame, CA). Histochemical reactions were performed using the Vectastain ABC Kit (Vector Labs) and Sigma Fast 3,3’-diaminobenzidine as substrate (Sigma, St Louis, MO). Sections were counterstained with haematoxylin.

**Morphometry.** Tissue sections were observed under a Nikon Eclipse E800 light microscope using an x20 objective, and digital images were captured with a DXM 1200 camera. CLS density (CLS per 400 adipocytes), adipocyte surface area and adipocyte volume were determined using a drawing tablet and the Nikon Lucia IMAGE Version 4.61 morphometric program. The mean cell volume was derived from the mean surface area and, positing that adipose cell density is the same as trioline (0.915 µg/µL), mean cell weight (µg lipid per cell) was mathematically derived from mean cell volume (17).

**Gene expression analysis.** Total WAT RNA was extracted from frozen perirenal fat of 2 month-old male wild type or HSL-/- mice using RNeasy Lipid Tissue kit (Qiagen, Mississauga, ON). Total bone marrow macrophage RNA was extracted after washing in 5 ml PBS and direct lysis of confluent cells in the Petri dish, following the manufacturer’s protocol (RNeasy kit, Qiagen). RNA served as template for first strand synthesis using poly dT primers and Superscript II reverse transcriptase (Invitrogen, Burlington, ON). For quantitative real-time PCR (SmartCycler, Cepheid), we used the QuantiTect SYBR green PCR kit (Quiagen). Primer sets used were: Mac-2a (5’ATGAAGAAGCTCCGGGAAT 3’, 3’ GCTTAGACTATGGCGTGGTT 5’), TNF-α (5’CCA GTGTGGGAAGCTGTCTT3’, 3’AAGCAAAAGAGGAGGCAACA 5’) and β-actin (5’CGTTGACATCCGTAAGACCT3’, 3’GCAGTAATCTCCTCTGCATCC 5’). Each reaction yielded amplicons of ~100 base pairs. PCR conditions were: 15 sec, 94°C; 20 sec, 60°C and 20 sec, 72°C, for 45 cycles. After amplification, a melting curve (0.01°C/sec) was used to confirm product purity. Results are expressed relative to β-actin.
Statistical analysis. Results are presented as mean ± standard deviation (SD). Data analysis was performed using Systat (v10) or Instat GraphPad. Differences between groups were assessed either by full interaction General Linear Models and protected post-hoc tests, paired or unpaired t-tests, or by Fisher exact tests, as appropriate. Differences between groups were considered statistically significant when $p \leq 0.05$. 
RESULTS

WAT macrophages localize almost exclusively to sites of necrotic-like adipocyte death in obese db/db mice and in obese humans. We initially assessed the frequency and distribution of mature macrophages in fixed WAT of lean (wild-type) and obese db/db mice, an established model of obesity-associated WAT inflammation (7). Macrophages were identified based on immunoreactivity for both F4/80 (not shown) and MAC-2 (also known as galectin 3), a lectin expressed by activated macrophages that mediates macrophage phagocytic and inflammatory responses (18-20). Surprisingly, macrophages were non-randomly distributed in WAT of both lean and obese mice. Rather than being dispersed throughout the tissue, macrophages were instead aggregated in “crown-like structures” (CLS), which contained up to 15 macrophages surrounding what appeared to be individual adipocytes (Fig. 1). CLS formation was a rare event in lean wild-type mice (0.34 ± 0.28 CLS per 100 adipocytes) (Fig. 1A), but increased ~30-fold (10.50 ± 1.05 CLS per 100 adipocytes) in obese db/db mice (p < 0.001) (Fig. 1B). Notably, the preponderance of MAC-2 positive macrophages in WAT of lean wild-type and obese db/db mice were localized to CLS (lean: 82.3 ± 5.3%; obese: 90.6 ± 1.8% [mean ± SD]). CLS contained only an occasional (<1%) neutrophil or lymphocyte and no preadipocytes, based on EM morphology (21) and absence of immunoreactivity for the preadipocyte marker S-100B (22, 23) (data not shown). CLS macrophages formed MAC-2 positive multinucleate giant cells (MGCs) (Fig. 1 C,D), indicating that individual adipocytes were acting as chronic sites of macrophage activation (12, 24). Together, these observations demonstrate that 1) the vast majority of WAT macrophages are aggregated in CLS that form around individual adipocytes and 2) the number of such “inflammatory” adipocytes increases dramatically in obese db/db mice.

We next used electron microscopy to examine ultrastructural features of those adipocytes that were surrounded by CLS. Remarkably, in all visceral and subcutaneous WAT depots of multiple obese mice, CLS were localized exclusively to dead adipocytes (Fig. 2A).
Adipocyte death in obese db/db mice exhibited none of the classical ultrastructural features of apoptosis—i.e., chromatin condensation, plasma membrane blebbing (zeiosis) or membrane-bound apoptotic bodies with nuclear fragments (25-27) (Fig. 2A,B). However, in all instances, obesity-associated adipocyte death exhibited features of necrosis, including ruptured plasma membranes, dilated ER, cell debris in the extracellular space, and the appearance of small lipid droplets in the cytoplasm (26-29) (Fig. 2A,B). Disrupted plasma membranes, small cytoplasmic lipid droplets and interstitial cell debris were recently noted as in vivo features of experimentally-induced adipocyte necrosis in mice (26). We also detected degenerating adipocytes that were not surrounded by CLS macrophages (Fig. 2B), indicating that sequestration of adipocytes within CLS is secondary to adipocyte demise. These adipocytes exhibited ultrastructural features of necrosis (i.e. ruptured basal membranes, organelle degeneration and small cytoplasmic lipid droplets) but no detectable features of apoptosis (Fig. 2B).

The appearance of small cytoplasmic lipid droplets suggested that the unilocular adipocyte lipid droplet degenerated during the process of obesity-associated adipocyte death. Lipid droplet degeneration was confirmed by immunohistochemistry for the essential lipid droplet-associated protein, perilipin, a key regulator of adipocyte lipolysis (Fig. 2C,D) (16, 30). No perilipin immunoreactivity was detected on lipid droplets of adipocytes surrounded by CLS (Fig. 2D). In contrast, perilipin was detected in adjacent, viable adipocytes (Fig. 2C, D). These results confirm that nonviable adipocytes are foci of macrophage localization and syncitia formation in WAT of obese db/db mice. In conjunction with ultrastructural studies of obesity-associated adipocyte death and CLS formation (Fig. 2A, B), these observations tell us that each CLS designates a site of necrotic-like adipocyte death. Given that >90% of macrophages in WAT of obese db/db mice are present in CLS (Fig. 1B), our results additionally demonstrate that the preponderance of macrophages in WAT of obese mice are selectively localized to individual dead adipocytes.
CLS macrophages persist at sites of obesity-associated adipocyte death to scavenge the exposed “free” lipid in the interstitium (Fig. 2A, E). The scavenging of “free” lipid by CLS macrophages is likely to be a protective response, at least initially (see Discussion). Internalization of adipocyte lipid (or its re-esterified derivatives) is associated with lipid droplet formation and intense macrophage immunoreactivity for adipocyte differentiation-related protein (ADRP) (Fig. 2E), a lipid droplet-associated protein that increases in cells in response to the accumulation of neutral lipid (31). Upregulated ADRP expression in CLS macrophages provides one explanation for the previous report of upregulated ADRP gene expression in WAT of obese db/db mice (7). The persistence of lipid-scavenging macrophages and the formation of MGCs at “free” lipid droplets of dead adipocytes (Fig. 1C, D) suggest that these droplets act as chronic sites of macrophage activation in WAT. However, we currently do not know if or how scavenging of adipocyte lipid and/or MGC formation by WAT macrophages promotes or prolongs the pro-inflammatory macrophage gene expression that characterizes obesity and its complications (5, 7).

We next investigated the relationship between WAT macrophage localization and obesity-associated adipocyte death in human subjects. We used CLS formation around perilipin-negative adipocytes as a marker of adipocyte death (Fig. 3A-D) and electron microscopy to assess necrotic and apoptotic features (Fig. 3E). As in db/db mice, CLS present in human WAT were localized exclusively to degenerate adipocytes, based on the absence of perilipin staining (Fig. 3 B, D). Again, adipocyte death exhibited only necrotic features, including plasma membrane rupture, presence of numerous small lipid droplets in the cytoplasm, and dilated endoplasmic reticulum and mitochondria (Fig. 3E).

Adipocyte death in human subjects was positively correlated with obesity, with adipocyte death detected in 13 of 19 biopsies from obese subjects (BMI >30-45), but in only 2 of 9 biopsies from lean or overweight individuals (BMI <30) (p = 0.04, Fisher Exact Test). Adipocyte death was also correlated with mean (± SD) adipocyte size, which increases in obesity (see
Adipocyte size in biopsies containing one or more dead adipocytes (0.37 ± 0.08 µg lipid per cell, n = 15) was twice that of biopsies containing no dead adipocytes (0.17 ± 0.09 µg lipid, n = 13) (p < 0.001). This association was observed in both subcutaneous and visceral WAT depots (Fig. 3F). Interestingly, adipocyte death was detected in a lean subject (BMI = 24) with large adipocytes (average = 0.40 µg lipid per cell), whereas no dead adipocytes were detected in an obese subject (BMI = 34) with relatively small adipocytes (average = 0.23 µg lipid per cell). These observations suggest that rates of adipocyte death increase as adipocytes enlarge.

Adipocyte hypertrophy in the absence of obesity is associated with necrotic-like adipocyte death and CLS formation. Increases in WAT mass reflect adipocyte hypertrophy, adipocyte hyperplasia or both (32). The correlation of adipocyte death with adipocyte size in humans (above) and associations of adipocyte hypertrophy with dysregulated adipocyte metabolism and elevated adipocyte stress (1, 5, 33, 34) suggested that adipocyte hypertrophy might promote obesity-associated adipocyte death. To test this hypothesis, we assessed the frequency of adipocyte death and CLS formation in hormone sensitive lipase (HSL) knockout mice (HSL-/- mice). HSL is a major lipase in mature adipocytes, and HSL abrogation results in increased adipocyte lipid storage and thus, adipocyte hypertrophy. Adipocyte hypertrophy in HSL-/- mice is not associated with increased adipose mass (15, 35-37). Thus, HSL-/- mice provide a genetic model with which to examine the association of adipocyte hypertrophy and adipocyte death in the absence of obesity.

In the present study, mean cell size of viable (perilipin-positive) adipocytes in HSL-/- mice (0.51 ± 0.14 µg lipid per cell) was >3-fold that of wild-type controls (0.15 ± 0.05 µg lipid per cell) (p < 0.001). As early as eight weeks of age, increased adipocyte size in HSL-/- mice was associated with macrophage infiltration and inflammation, manifested as elevated levels of transcripts for MAC-2 (7-fold) and TNF-α (5-fold) in perirenal WAT (p ≤ 0.05) (Table 1; see also
TNF-α gene expression was not elevated in HSL-/- macrophages that were differentiated in vitro from bone marrow-derived precursors (data not shown). Thus, elevated TNF-α gene expression is not an intrinsic property of non-activated macrophages in HSL-/- mice. These data indicate that adipocyte hypertrophy in HSL-/- mice is coincident with macrophage recruitment and activation, and with WAT inflammation in the absence of increased adipose mass.

As in obese db/db mice and obese humans, the overwhelming preponderance (98 ± 0.7%) of macrophages (F4/80, MAC-2 positive cells) in WAT of lean HSL-/- mice were aggregated in CLS that selectively surrounded nonviable (perilipin-negative) adipocytes (Fig. 4A). By 12 weeks of age, the frequency of adipocyte death (CLS formation around perilipin-negative adipocytes) in WAT of HSL-/- mice was ~15-fold greater (29.0 ± 2.0 CLS per 100 adipocytes) than the frequency of adipocyte death in wild-type controls (2.0 ± 0.4 CLS per 100 adipocytes). Adipocyte death in lean HSL-/- mice exhibited ultrastructural features of necrosis, including basal membrane rupture and ER dilatation (Fig. 4B). As observed in WAT of obese mice and humans (Figs. 2C, D and Fig. 3E), lipid droplet degeneration was a characteristic early feature of adipocyte death in HSL-/- mice (Fig. 4B). Moreover, as with adipocyte death in obese mice and humans, we detected no chromatin or nuclear condensation (Fig. 4B), consistent with a previous report that the frequency of apoptotic (i.e., TUNEL-positive) cells is not increased in WAT of HSL-/- mice (37). CLS macrophages in WAT of HSL-/- mice actively ingested exposed adipocyte lipid (Fig. 4C), elaborated numerous lipid droplets, and expressed intense ADRP immunoreactivity (Fig. 4D). Thus, adipocyte hypertrophy in the absence of increased adipose mass is associated with 1) increased rates of necrotic-like adipocyte death, 2) selective macrophage recruitment to sites of adipocyte death and 3) formation of lipid-scavenging macrophage syncitia around “free” adipocyte lipid droplets. This is identical to events observed in WAT of obese mice and humans (Figs. 1-3).
DISCUSSION

Bone marrow-derived macrophages accumulate in WAT of obese mice and humans, where they promote adipose and systemic inflammation and the development of obesity-related metabolic complications (5, 7, 38). However, neither the underlying cause nor the function of macrophage recruitment in WAT of obese subjects is known. In the present study we demonstrate that 1) the preponderance of WAT macrophages in lean and obese mice and humans is localized to individual dead adipocytes, 2) at these sites, macrophages express activation markers (MAC-2), form syncitia (CLS) that surround and scavenge residual adipocyte lipid, and ultimately form multinucleate giant cells, a hallmark of chronic inflammatory states, and 3) the frequency of adipocyte death is increased >30-fold in a mouse (db/db) model of obesity-associated WAT inflammation (7) as well as in obese humans (BMI > 30). These observations suggest that adipocyte death promotes macrophage recruitment, accumulation and persistence in WAT of obese individuals.

The mechanism of adipocyte death has potentially significant implications for the development of obesity-associated WAT inflammation. Broadly speaking, apoptosis is non-inflammatory, reflecting the packaging of cell constituents into inflammation-suppressive apoptotic bodies (26, 27, 39). In contrast, during necrosis, cell contents are released into the extracellular space where they evoke inflammatory responses (27, 39, 40). Electron microscopy is the most reliable technique for assessing classical apoptosis, which is defined by stereotypic morphological changes, including chromatin condensation, plasma membrane blebbing and the formation of membrane-bound apoptotic bodies containing nuclear fragments (26, 27, 39). In the present study, obesity-associated adipocyte death exhibited none of these apoptotic hallmarks. Rather, moribund and dead adipocytes invariably displayed features of necrosis, most notably, rupture of the plasma membrane (Figs. 2A,B and 4B), dilatation of the endoplasmic reticulum (Figs. 3E and 4B) and release of cell contents to the extracellular space.
Fig. 2A) (26, 27, 40). These observations suggest that obesity-associated adipocyte death occurs by necrosis.

Arguing against this conclusion and supporting an apoptotic mechanism of obesity-associated adipocyte death is the relative absence of neutrophils at sites of adipocyte death (present study) and in adipose tissue of obese mice and humans (7). Whereas apoptosis can activate macrophages without the participation of other pro-inflammatory cells (i.e. neutrophils), the inflammatory response to necrosis is typically a sequential process in which neutrophils initially participate. Neutrophil infiltration into WAT is observed in response to adipocyte necrosis in human pathologic conditions as well as acutely in response to experimentally-induced adipocyte necrosis in mice (13, 14, 29, 41). Thus, obesity-associated adipocyte death appears to occur by an alternative death pathway sharing features of both necrosis and apoptosis. A number of such alternative death pathways (e.g., “necrosis-like programmed cell death”, “paraptosis”) have recently been described in which apoptotic signaling pathways and proteolytic cascades may participate, but the morphology of cell execution is predominantly necrotic (27, 40). These necrosis-like forms of apoptotic death typically lack chromatin condensation and apoptotic bodies (as in the present study). A critical unresolved question is whether this alternative pathway of adipocyte death promotes pro-inflammatory activation of CLS macrophages and MGCs, thereby contributing to the chronic pro-inflammatory tone in WAT of obese individuals (5, 7).

The development of obesity is associated with release by adipocytes of macrophage chemoattractants, arrestants and activators (5, 38, 42, 43). Elevated release of these substances by metabolically dysregulated or otherwise “distressed” adipocytes or by adjacent cells could provide one mechanism to account for the preferential localization of WAT macrophages to dead adipocytes. However, it is conceivable that events unrelated to obesity-associated adipocyte death induce initial macrophage extravasation from the blood into WAT, and that these macrophages are subsequently recruited to moribund adipocytes. Irrespective of
the factors promoting initial macrophage recruitment to WAT, the relative paucity of macrophages that were not localized to dead adipocytes in the present study suggests that macrophages are recruited to degenerating or dead adipocytes soon after extravasation into WAT.

Our data for human subjects and for HSL-/- mouse model of adipocyte hypertrophy without obesity suggest that adipocyte hypertrophy per se promotes adipocyte death, macrophage aggregation and CLS formation. Therefore, hypertrophy-induced adipocyte death is a plausible mechanism underlying the reported association between mean adipocyte size and the magnitude of macrophage infiltration and inflammatory gene expression in WAT (7). In addition, the association between adipocyte size and adipocyte death provides a potential explanation for the fact that, during the development of obesity, adipocytes get only “so large,” after which adipocyte hypertrophy gives way to adipocyte hyperplasia (32). The present study suggests that the initial hypertrophic response of adipocytes renders them prone to necrotic-like death, after which only adipocyte hyperplasia can efficiently maintain or increase the lipid storage capacity of adipose tissue.

The mechanism(s) by which hypertrophy potentially promotes adipocyte death is unclear, but cell stress is implicated. Hypertrophic adipocytes are subjected to multiple cytotoxic stressors, including ER stress, hypoxia, elevated TNF-α, reactive oxygen species, and free fatty acids (1, 4, 33, 34, 44, 45). These cytotoxic stresses activate inflammatory signaling cascades that regulate stress-induced cell death (46, 47) and downregulate adipocyte insulin signaling and PPARγ-regulated gene expression (34, 48, 49). Interestingly, experimental depletion of systemic insulin levels or targeted disruption of PPARγ gene expression in mature adipocytes has been shown to induce adipocyte necrosis and subsequent inflammatory cell infiltration in mouse WAT (29, 41). Thus, compromised insulin signaling and/or downregulated expression of PPARγ-regulated genes in hypertrophic adipocytes may contribute to obesity-associated adipocyte death. Stress-induced mechanisms of adipocyte death may be complemented and
locally amplified by cytotoxins released from activated WAT macrophages (i.e., TNF-α, NO, reactive oxygen species). Thus, WAT macrophages could conceivably participate in adipocyte “execution” (7), thereby contributing to a feed-forward mechanism of adipocyte death and macrophage recruitment.

MAC-2 expression in CLS macrophages and MGCs is consistent with the lectin’s roles in leukocyte recruitment, phagocytosis, lipid scavenging, and inflammatory progression (18-20, 50, 51). MAC-2 expression at sites of adipocyte necrosis may be functionally linked to macrophage aggregation and CLS formation, as MAC-2 is a potent macrophage chemoattractant that promotes macrophage aggregation (19). MAC-2 also possesses anti-apoptotic activity that is implicated in macrophage survival at sites of inflammation (51). The chemoattractant and anti-apoptotic activities of MAC-2 are likely to facilitate and prolong macrophage recruitment, aggregation and function at sites of adipocyte death.

Clearance of “free” lipid appears to be an important function of MAC-2-expressing macrophages in WAT (present study) as well as in the arterial wall (52) and in the regenerating peripheral nervous system following injury (53). In WAT, “free” lipid droplets released from necrotic adipocytes must be sequestered and removed from the interstitium, since they are insoluble and provide an unprotected source of cholesterol and cytotoxic fatty acids that could damage WAT cells (44, 54, 55). In addition, clearance of “free” lipid (and other adipocyte debris) must precede remodeling of the extracellular matrix and the recruitment of adipocyte precursors required for new fat cell differentiation at sites of adipocyte loss (56). This and other remodeling functions of WAT macrophages (11) may be particularly important in maintaining and expanding the lipid storage capacity of obese individuals despite dramatically increased rates of adipocyte death. However, it is currently undetermined if and how lipid-laden macrophage syncitia are “cleared” from WAT to make room for new adipocytes.

In summary, we demonstrate for the first time that 1) the preponderance of macrophages in WAT of lean and obese mice and humans is selectively localized to dead
adipocytes; 2) rates of adipocyte death increase dramatically in obesity, potentially reflecting cytotoxic effects of adipocyte hypertrophy; 3) obesity-associated adipocyte death appears to involve an alternative death pathway exhibiting morphological features of necrosis and the leukocyte-eliciting profile of apoptosis; 4) “free” lipid droplets of dead adipocytes act as persistent sites of macrophage fusion, lipid uptake and MGC formation. These observations provide a novel framework for understanding macrophage recruitment and function in WAT and for elucidating underlying causes of obesity-associated WAT inflammation.

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FIGURE LEGENDS

Figure 1. WAT macrophages localize to crown-like structures (CLS) around individual adipocytes, which increase in frequency with obesity. Light microscopy of visceral WAT of lean (A) and obese db/db (B) mouse showing MAC-2 immunoreactive macrophages (brown color) aggregated to form rare (A: lean) or numerous (B: obese) crown-like structures (CLS) among unilocular adipocytes. Note that almost all MAC-2 immunoreactive macrophages are organized to form CLS. (C) Enlargement of the bottom right corner of panel B showing that almost all mononuclear cells in CLS are MAC-2 immunoreactive (i.e., activated macrophages). Note the multinucleate giant cell (MGC), which stains intensely for MAC-2. (D) Serial section consecutive to panel C confirming the presence of multiple nuclei (blue) in the MGC. Bar length: panels A and B =100 µm; C = 28 µm; D = 10 µm.

Figure 2. CLS form exclusively at sites of adipocyte death and scavenge the residual adipocyte lipid droplet. (A) Electron micrograph showing CLS macrophages (M) surrounding the residual lipid droplet (L) of a dead adipocyte. Evidence of necrosis includes disrupted basal membrane (BM), cytoplasmic remnants (CR), cell debris (CD) in the interstitium, and the apparent degeneration of the unilocular lipid droplet into small droplets. White arrows indicate lipid-laden phagolysosomes. Inset: enlargement of the squared area showing small adipocyte lipid droplets engulfed by a macrophage. (B) Necrotic features in db/db mouse adipocyte before recruitment of CLS macrophages. Note rupture of basal membrane (arrowheads) and loss of lipid droplet (L) integrity manifest as small lipid droplets in the cytoplasm (arrows). (C,D) Lipid droplet degeneration in necrotic adipocytes can be detected at the light microscope level by the loss of lipid droplet-associated proteins. All adipocytes of lean mice (C) are perilipin immunoreactive (brown), whereas adipocytes surrounded by CLS in obese mice (D) are not immunoreactive for perilipin (or ADRP [not shown]). Note that viable adipocytes (not
surrounded by CLS) in obese mice (D) retain perilipin immunoreactivity (arrows). (E)

Scavenging of residual adipocyte lipid droplet by CLS macrophages is associated with
upregulated ADRP expression (brown) and formation of ADRP-positive lipid droplets in CLS
macrophages. Bar length: panel A = 5.3 µm; inset = 2.7 µm; B = 0.8 µm; C = 50 µm; D = 43
µm; E = 16 µm.

Figure 3. CLS formation at sites of adipocyte death in human WAT and correlation of
adipocyte death and adipocyte size. Human subcutaneous (A,B) and visceral (omental)
(C,D) fat depots demonstrating MAC-2-positive macrophages (A,C) organized exclusively in
CLS surrounding degenerate (perilipin-free) (B,D) lipid droplets (asterisk). Arrows in panel A
denote MAC-2 immunoreactivity. Inset in A more clearly shows surface staining for MAC-2 in a
CLS macrophage from the same patient. Arrows in panels C and D denote perilipin
immunoreactivity of lipid droplets in viable adipocytes. (E) Representative electron micrograph
showing necrotic degenerative features of obesity-associated adipocyte death in human WAT.
Evidence of necrosis includes dilated endoplasmic reticulum (ER) and mitochondria (not shown)
and the presence of numerous small lipid droplets (arrows, enlarged in the bottom right panel).
Note that the nucleus (L) exhibits no signs of apoptosis (i.e., chromatin condensation, pyknosis).
(F) Adipocyte death is correlated with mean adipocyte size in human subcutaneous (SC) and
visceral (V) adipose tissue. Fat biopsies from human subjects (BMI 20-45) were scored for the
presence (+) or absence (-) of CLS (i.e., MAC-2 positive cells surrounding a perilipin-negative
lipid droplet). *, p < 0.05 for within-depot comparison. Bar length: panels A-D = 27 µm; E = 3.7
µm; inset = 1.5 µm.
Figure 4. Adipocyte death, CLS formation and lipid scavenging in HSL-/- mice, a model of adipocyte hypertrophy without obesity. (A) Perilipin immunohistochemistry demonstrating CLS formation selectively around perilipin-free adipocyte-like structures (*), but not around viable adipocytes expressing perilipin (arrows). (B) Electron micrograph demonstrating degenerative aspects (early signs of necrosis) and necrosis in HSL-/- mouse. Degenerative aspects include dilatation of the endoplasmic reticulum (ER) and lipid infiltration (arrows) into the nucleus (L). Cell death is indicated by the ruptured basal membrane (BM) and free lipid droplets (FL) in the interstitium. The absence of chromatin or nuclear condensation suggests degeneration in the absence of “classical” apoptosis. M, macrophage; L, adipocyte lipid droplet. (C) High power electron micrograph showing internalization of free lipid (FL) derived from the degenerating adipocyte lipid droplet (L) by CLS macrophage (M). (D) CLS macrophages of HSL-/- mice elaborate numerous intracellular lipid droplets and are intensely immunoreactive for the lipid droplet-associated protein, ADRP. Bar length: panel A = 36 µm; B = 1.25 µm; C = 0.45 µm; D = 15 µm.
Table 1. Evidence of macrophage infiltration and inflammatory activation in WAT of HSL-/- mice. Relative levels of MAC-2 and TNF-α mRNA in HSL-/- and HSL+/+ mice were determined by real-time PCR of perirenal WAT from 2 month-old male mice (n = 6) of each genotype. Gene expression is expressed relative to that measured for beta-actin (see Materials and Methods). Ratios comparing HSL-/- and HSL+/+ mice are shown.

<table>
<thead>
<tr>
<th>mRNA</th>
<th>Genotype</th>
<th>Ratio</th>
<th>P</th>
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<tbody>
<tr>
<td>MAC-2</td>
<td>HSL-/-</td>
<td>105.2 ± 26.7</td>
<td>14.8 ± 5.1</td>
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<tr>
<td></td>
<td>HSL+/+</td>
<td>14.8 ± 5.1</td>
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<tr>
<td>TNF-α</td>
<td>HSL-/-</td>
<td>0.9 ± 0.3</td>
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<tr>
<td></td>
<td>HSL+/+</td>
<td>0.2 ± 0.1</td>
<td></td>
</tr>
</tbody>
</table>
Figure 1.
Figure 2.
Figure 3.
Figure 4