Cafeteria Diet Is a Robust Model of Human Metabolic Syndrome With Liver and Adipose Inflammation: Comparison to High-Fat Diet

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Obesity has reached epidemic proportions worldwide and reports estimate that American children consume up to 25% of calories from snacks. Several animal models of obesity exist, but studies are lacking that compare high-fat diets (HFD) traditionally used in rodent models of diet-induced obesity (DIO) to diets consisting of food regularly consumed by humans, including high-salt, high-fat, low-fiber, energy dense foods such as cookies, chips, and processed meats. To investigate the obesogenic and inflammatory consequences of a cafeteria diet (CAF) compared to a lard-based 45% HFD in rodent models, male Wistar rats were fed HFD, CAF or Chow control diets for 15 weeks. Body weight increased dramatically and remained significantly elevated in CAF-fed rats compared to all other diets. Glucose- and insulin-tolerance tests revealed that hyperinsulinemia, hyperglycemia, and glucose intolerance were exaggerated in the CAF-fed rats compared to controls and HFD-fed rats. It is well-established that macrophages infiltrate metabolic tissues at the onset of weight gain and directly contribute to inflammation, insulin resistance, and obesity. Although both high fat diets resulted in increased adiposity and hepatosteatosis, CAF-fed rats displayed remarkable inflammation in white fat, brown fat and liver compared to HFD and controls. In sum, the CAF provided a robust model of human metabolic syndrome compared to traditional lard-based HFD, creating a phenotype of exaggerated obesity with glucose intolerance and inflammation. This model provides a unique platform to study the biochemical, genomic and physiological mechanisms of obesity and obesity-related disease states that are pandemic in western civilization today.

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INTRODUCTION

In the United States, obesity has recently reached pandemic levels for a variety of reasons including inactivity and poor diet (1,2). The most recent National Health and Nutrition Examination Survey (NHANES) showed that 68.3% of those studied were considered overweight (BMI ≥25) and 33.9% were obese (BMI ≥30) (1). Childhood obesity is particularly important as more children are developing what were previously considered adult diseases, like type II diabetes, high blood pressure, elevated serum insulin and dyslipidemia; with obese children often progressing to obese adults (2,3).

Genetic models of obesity, such as ob/ob mice, db/db mice, Zucker fa/fa obese rats, Agouti yellow mice, and melanocortin 4 receptor knockout mice, have contributed significantly to the understanding of genetic anomalies underlying mechanisms that control energy homeostasis (4). Other investigations into the aberrant homeostasis responsible for most human obesity were carried out using rodent models of diet-induced obesity (DIO) through the administration of high-fat diets (HFD) (4). Traditionally, these diets consist of a simple exchange of carbohydrate-derived calories with fat-derived calories when compared to low fat or chow control diets. However, another experimental rodent diet model exists that more accurately reflects the variety of highly palatable, energy dense foods that are prevalent in Western society and associated with the current obesity pandemic: the “cafeteria diet” (CAF) (5,6). In this model, animals are allowed free access to standard chow and water while concurrently offered highly palatable, energy dense, unhealthy human foods ad libitum. This diet promotes voluntary hyperphagia that results in rapid weight gain, increases fat pad mass and prediabetic parameters such as glucose and insulin intolerance (7–10). Moreover, this diet

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engages hedonic feeding, consistent with the observed voluntary hyperphagia, which produces long-lasting neuronal alterations predicted to underlie non-homeostatically regulated feeding behaviors associated with some forms of human obesity (11,12). Current US Department of Agriculture labeling requirements combined with meticulous laboratory record keeping enables a research model of DIO that models progression to insulin resistance and diabetes.

To date, however, direct comparisons between the CAF model and other DIO animal models have not been adequately evaluated. Therefore, we embarked on a comprehensive comparative study, which demonstrated that the CAF diet is a more robust model for human obesity compared to HFD. Our hypothesis was that CAF diet would be a more representative model of human metabolic syndrome; indeed, we provide supportive evidence of severe obesity, nonalcoholic steatohepatitis (NASH), and glucose intolerance compared to a traditional lard-based HFD. In addition, as the role of chronic low-grade systemic inflammation is increasingly recognized to play a part in metabolic syndrome, we demonstrated dramatic CAF-induced inflammation in liver and white and brown adipose tissues (BAT). Use of this rapid and potent CAF DIO model may be useful to identify novel pathways for preventative intervention or therapeutics in the treatment of human obesity.

**METHODS AND PROCEDURES**

**Animals**

Male Wistar rats (~200 g, 7–8 weeks old) (Harlan Laboratories, Dublin, VA) were housed 2 rats/cage in a 12 h light/dark cycle and acclimated to the Duke animal housing facility on ad libitum undefined standard chow 7001 (“SC,” Harlan Teklad) for 2 weeks before assignment to one of four experimental diet groups. Upon initiation of experimental diets, rats (avg. 300 g, 9–10 weeks old) were either maintained on ad libitum SC as controls or switched to ad libitum defined 45% fat chow (“HFD,” Research Diets D06011802), the matched low-fat chow control (low-fat diet (LFD) Research Diets D07010502), or provided a CAF with three human snack foods varied daily in addition to ad libitum SC (see Supplementary Table S1 online). The CAF diet foods included cookies, cereals, cheese, processed meats, crackers, etc. and were provided in excess (see Supplementary Table S2 online). Snack items were weighed before and after consumption, corrected for drying, and varied daily according to the fat, protein, and carbohydrate content as listed in Supplementary Table S2 online. Fat intake was the largest macronutrient alteration in CAF rats, however, simple carbohydrate consumption was also elevated over HFD and SC rats groups. There are no added sugars to the SC diet; any sugars present are derived from whole-grains (corn, oats, etc.). The manufacturer Harlan Teklad estimates that simple sugars are <5% of total carbohydrates in SC7001. It should be noted that the LFD-fed rats also consumed the most simple sugars as a result of sucrose substitution for fat in the diet formula (see Supplementary Table S3 online).

Two experimental groups of SC and CAF-fed rats (n = 8–9 per diet per group) along with one experimental group of SC, LFD, and HFD-fed rats (n = 8 per diet) were initiated in parallel overlapping by 4–6 weeks. Food consumption and weight gain was monitored daily for 10 weeks (Figure 1a–d) to carefully characterize food intake on CAF diet. A subsequent group of experimental animals (n = 8–10 per diet) were fed the four respective diets for 15 weeks and analyzed for glucose tolerance, insulin sensitivity, metabolic alterations, and tissue mRNA, metabolite and phenotypic analysis (Figures 1e,f,2,3,4,5,6, and 7, Supplementary Figures S1 and S2 online). Nonesterified fatty acids in plasma were measured enzymatically (Wako, Richmond, VA) using the Hitachi 911 clinical chemistry analyzer (Hitachi, Tokyo, Japan).

**GTT and ITT**

For 1 week prior to the glucose tolerance test (GTT) or insulin-tolerance test (ITT), animals were acclimated daily to the procedure room and handling for injections and blood draws. Rats were fasted overnight (16h) and a basal blood draw from the saphenous vein was collected for plasma glucose and insulin measurements 1h before glucose administration. The GTT was performed after 7 weeks on respective diets; rats were given an intraperitoneal injection of glucose (1 g/kg of body weight; 45% glucose). Blood was obtained from the tail vein via nick immediately before glucose injection and every 30 min thereafter for a period of 180 min, and glucose levels were measured with BD Logic Glucose Meter (Becton-Dickinson, Franklin Lakes, NJ). ITT was performed after 11 weeks where rats were given an intraperitoneal injection of insulin (0.75IU/kg body weight; Humulin Regular; Eli Lilly, Indianapolis, IN) and blood glucose was measured as in GTT. The homeostasis model assessment was used to calculate approximate insulin resistance (glucose (mg/dl) × insulin(405) (13–16). Plasma insulin was measured with a mouse insulin enzyme-linked immunosorbent assay (Millipore, Billerica, MA).

**Tissue collection**

After 15 weeks of experimental diet, food was removed for 4–6 h before sacrifice under nembutal anesthesia for tissue collection. Cardiac blood was collected and divided into plasma (1.5 ml blood + 10 μl 1.7 mg/ml EDTA) and serum (clotted for 2 h at room temperature) aliquots, collecting the supernatant fractions after centrifugation (3,800g, 10 min, 4 °C). Pancreas, liver, epididymal white adipose tissue (eWAT), and BAT were rapidly isolated, washed in ice-cold phosphate-buffered saline, patted dry and weighed before freeze clamping and snap freezing in liquid nitrogen. A portion of eWAT, BAT, liver, and pancreas was placed in a cassette and fixed in Bouin’s fixative overnight, washed with 70% ethanol, and paraffin-embedded for histologic analysis. The remaining portion of the eWAT fat pads was digested to obtain the stromal-vascular fraction (SVF) (17). Briefly, tissue was immediately immersed into 25 mmol/l HEPES-buffered Dulbecco’s modified Eagle’s medium supplemented with 1% fatty acid free low endotoxin BSA (A8806; Sigma-Aldrich, St Louis, MO), and minced into <10 mg pieces, and filtered through nylon mesh. An lipopolysaccharide-depleted collagenase cocktail (Liberase 3; Roche Diagnostics, Indianapolis, IN) at a concentration of 0.03 mg/ml with 0.2 mg/ml DNAse I (Sigma-Aldrich) was added to each tissue filtrate. The samples were incubated in a shaking water bath at 37 °C for 20–30 min. Once digestion was complete, samples were passed through 250-μm nylon mesh (Sepaf, Depew, NY). The suspension was centrifuged at 200g for 5 min to pellet the SVF. The SVF was isolated after lysing red blood cells and a final spin to remove liquid.

**Histology**

Paraffin-embedded tissues were sectioned at 5 microns and mounted for histological staining. Fixative was removed using a graduated series of xylenes (Sigma, St Louis, MO), tissues were rehydrated and then flushed for an additional 15 min in deionized water to remove any remaining fixative. Tissue sections were stained using Harris hematoxylin and eosin (H&E) staining or prepared for immunostaining of adjacent serial sections. Briefly, immunohistochemistry was carried out using serial sections corresponding to H&E-stained tissue sections, processed for heat-induced epitope-retrieval using the Dako Target Retrieval Solution in conjunction with Dako PT Link (Dako North America, Carpenteria, CA). Samples were washed once in Tris-buffered saline/Tween 20 and then exposed to 3% hydrogen peroxide to ablate the innate peroxidase activity of the tissue. Tissues were then blocked in a commercial protein blocking solution (Dako, X0909) and exposed to primary antibodies at 4 °C overnight with gentle agitation (1:250). The sections were washed three times in Tris-buffered saline/Tween 20
for 15 min each. Biotinylated secondary antibody was added for 1 h at room temperature (1:500), and the tissues were washed. Visualization of the target epitopes recognized by primary and secondary antibody interactions were established using an avidin/biotinylated horseradish peroxidase kit (Vectastain Elite ABC; Vector Laboratories, Burlingame, CA; PK6100, 1:50) incorporating DAB as the chromogen. The primary antibodies used were raised against macropore markers F4/80 (Santa Cruz, Santa Cruz, CA, sc-25830, 1:100) and CD163 (AbD Serotech, Raleigh, NC, 1:100), and the tissue sections briefly counterstained with Harris hematoxylin to highlight tissue architecture and nuclear location. Digital photomicrographs of stained tissues were captured using a Nikon Eclipse TS100 equipped with a digital camera using TSlide2 software, Version 6.0.2.1 (Tucson Imaging Technology, Fujian, China).

Pancreatic islets of Langerhans were visualized by performing immunohistochemistry with an insulin antibody. Briefly, paraffin-embedded sections of pancreata were deparaffinized, rehydrated, quenched of endogenous peroxidase activity, blocked, incubated with rabbit anti-insulin primary antibody (Santa Cruz) overnight at 4 °C, and incubated with ImmPRESS reagent (peroxidase conjugated anti-rabbit Ab; Vector Laboratories) for 30 min NovaRed (Vector Laboratories) was used as the substrate for colorimetric detection. Sections were lightly counterstained with hematoxylin. Images were acquired using an Axio-Observer Z1 microscope (Zeiss, St Louis, MO) fitted with an AxioCam HRC camera and operated with Axio-Vision Software, v. 4.8 (Zeiss). Representative images for each tissue (eWAT, liver, BAT, pancreas) from each treatment group were selected. To determine the inflammatory state of the eWAT, three randomly selected 10× fields were assessed for crown-like structures (CLS) from both H&E stain and F4/80 macrophage staining, per study animal. Liver inflammation was determined by enumerating the inflammatory loci per 10× field using three fields H&E and three fields F4/80 stained for each animal (randomly selected). Data are presented as the mean ± s.e.m. for each treatment group.

RNA isolation
QIAzol Lysis Reagent was used to isolate mRNA from snap-frozen SVF pellets with DNase treatment (RNaseq Lipid Tissue Mini Kit mRNA; Qiagen, Valencia, CA). Samples were isolated in duplicate on two columns and eluted into the same tube. mRNA quantity and quality were analyzed by Nanodrop (Thermo Scientific, Wilmington, DE) and Bioanalyzer 2100 (Agilent, Wilmington, DE), respectively. For quantitative PCR analysis, cDNA was synthesized using iScript (BioRad, Hercules, CA). Real-time quantitative PCR was completed using the Roche LightCycler 480 using LightCycler 480 SW 1.5 software and TaqMan Assay on Demand primers for tumor necrosis factor α mRNA; Qiagen, Valencia, CA). Samples were isolated in duplicate on SVF pellets with DNAse treatment (RNeasy Lipid Tissue Mini Kit mRNA; Qiagen, Valencia, CA). QIAzol Lysis Reagent was used to isolate mRNA from snap-frozen tissue sections of pancreata were deparaffinized, rehydrated, quenched of endogenous peroxidase activity, blocked, incubated with rabbit anti-insulin primary antibody (Santa Cruz) overnight at 4 °C, and incubated with ImmPRESS reagent (peroxidase conjugated anti-rabbit Ab; Vector Laboratories) for 30 min NovaRed (Vector Laboratories) was used as the substrate for colorimetric detection. Sections were lightly counterstained with hematoxylin. Images were acquired using an Axio-Observer Z1 microscope (Zeiss, St Louis, MO) fitted with an AxioCam HRC camera and operated with Axio-Vision Software, v. 4.8 (Zeiss). Representative images for each tissue (eWAT, liver, BAT, pancreas) from each treatment group were selected. To determine the inflammatory state of the eWAT, three randomly selected 10× fields were assessed for crown-like structures (CLS) from both H&E stain and F4/80 macrophage staining, per study animal. Liver inflammation was determined by enumerating the inflammatory loci per 10× field using three fields H&E and three fields F4/80 stained for each animal (randomly selected). Data are presented as the mean ± s.e.m. for each treatment group.

Statistics
Data are shown as means ± s.e.m. Analysis was performed using ANOVA followed by a Tukey–Kramer honestly significant difference, and when appropriate a two-tailed Student’s t-test * P < 0.05 using JMP (SAS Institute, Cary, NC).

RESULTS
CAF diet-induced hyperphagia and weight gain
Prior to the start of the study, there were no differences in food intake or body weight between animals assigned to experimental diet groups. We first evaluated food intake, fat intake, and body weight changes in male Wistar rats fed four different diets (Figure 1 and Supplementary Table S1 online): (i) “SC”: control rats were continued on standard chow (SC7001) fed from birth, which contains ~12% kcal derived from fat (porcine fat and linoleic acid); (ii) “CAF”: continued SC7001 access plus a daily assortment of three human snack items; (iii) “LFD”: a defined standard chow (SC502) with 10% kcal derived from fat (lard and soybean oil); (iv) “HFD”-matched to SC502 with decreased carbohydrate content and increased lard content (45% kcal derived from fat). Rats on the CAF diet consumed standard chow at ~15% of that consumed by rats fed standard chow only.

Average weekly nutrient intake per rat in a 10-week food intake study is summarized in Supplementary Table S3 online. All rats consumed ~100 kcal/day/rat except for CAF-fed rats who significantly consumed about 30% more calories (Supplementary Table S3 and Figure 1a). HFD-fed rats, unlike CAF-fed rats, significantly decreased total gram intake

Figure 1 Cafeteria diet (CAF) drives food and fat consumption. (a) Male Wistar rats on CAF have elevated food intake in terms of kcal compared to standard chow (SC) control, high-fat diet (HFD), and low-fat diet (LFD). (*HF P < 0.05 vs. all; **CAF P < 0.001 vs. all). (b) Upon initiation of diets, only HFD-fed rats decreased food intake in grams (g) with an acute increase in CAF-intake. (*CAF P < 0.005 vs. all; **CAF P < 0.005 vs SC and HF; ***CAF P < 0.005 vs. all; **CAF P < 0.005 vs SC and CAF). (c) CAF-fed rats consume more fat when compared to animals receiving HFD, LFD, or SC. (**CAF P < 0.001 vs SC and LF; **CAF P < 0.05 vs HF). (d) CAF-diet fed rats exhibited significant weight gain starting at 2 weeks and persisting throughout the study. Data are normalized to SC. (*CAF, LFD, or HFD P < 0.05 vs. SC, **CAF P < 0.05 vs SC). (e) After 15 weeks on diet, CAF-fed rats exhibited greater total weight gain when compared to SC, LFD, and HFD. Although HFD-fed rats also gained weight, there is no difference in the weight gain between the HFD and LFD groups, however weight gain did reach statistical significance when compared to the SC control group. (**HF P < 0.05 vs SC; **CAF P < 0.05 vs all). (f) There were no changes in body length after 15 weeks on diet. (a–d) were conducted on a set of rats fed diet for 10 weeks. (n = 23–24, SC, n = 8 LFD, HFD, n = 17 CAF) and (e, f) were conducted on a group of rats fed diet for 15 weeks (n = 8, SC, LFD, and HFD, n = 10 CAF).
of the calorically dense diet to normalize caloric consumption by week 2, an effect commonly seen in rodents placed on HFD (Figure 1b). Fat intake consumption paralleled percent fat content of diet with SC and LFD consuming ~10 kcal/day/rat, the HFD group at 45 kcal/day/rat, and the CAF-diet group at 55 kcal/day/rat (Figure 1c). CAF-fed rats gained the most weight on the diet, almost double the weight gain evident in SC controls, whereas LFD and HFD-fed rats demonstrated similar and intermediate weight gain (Figure 1d). A subsequent group was placed on the four experimental diets for 15 weeks, demonstrating similar and continued weight gain (Figure 1e and Supplementary Figure S1 online) with no statistical differences in body length among treatment groups (Figure 1f).

**Figure 2** Cafeteria diet (CAF)-fed rats displayed hyperglycemia and hyperinsulinemia. Insulin sensitivity was evaluated on rats at weeks 7 and 11 during the 15-week study. (a,b) Fasting glucose levels were not different at 7 weeks of age while only standard chow (SC)-fed rats maintained low insulin levels, which were significantly different than CAF- and high-fat diet (HFD)-fed rats. (c) Homeostasis model assessment of insulin resistance (HOMA\(_{IR}\)) calculations at 7 weeks on diets indicated that CAF and HF groups demonstrated statistically significant elevations in insulin resistance compared to SC-fed rats. (*P < 0.05 vs. SC). (d) CAF-fed rats had significantly elevated blood glucose after a 6h light cycle fast at 15 weeks on diet (time of sacrifice) compared to SC, while low-fat diet (LFD) and HFD maintained normal glucose levels. (n = 8, SC, LFD, and HFD, n = 10 CAF). (*P < 0.05 SC to CAF).

**CAF-diet induced hyperglycemia and insulin resistance**

At an intermediate time point (7 weeks) on diets, glucose homeostasis was maintained irrespective of diet, as blood glucose measured in obese and lean rats fasted overnight was not significantly different amongst the four diet groups (Figure 2a). However, insulin was approximately doubled in CAF- and HFD-fed rats compared to SC controls (Figure 2b). Consistent with the observed intermediate body weight phenotype, LFD-fed rats displayed a modest, nonsignificant elevation in fasting insulin compared to SC. Similarly, homeostasis model assessment of insulin resistance HOMA\(_{IR}\) calculations as a measure of insulin sensitivity indicated that CAF- and HFD-fed rats were twofold less insulin sensitive than SC controls at just 7 weeks into the diets (Figure 2c). After extended exposure to diets (15 weeks), hyperglycemia became evident in CAF-fed rats only, as 3 of 10 exhibited hyperglycemia after a 6h fast before sacrifice with no differences between LFD, HFD, and SC control groups (Figure 2d). Based on HOMA\(_{IR}\) findings, intraperitoneal GTT (IP-GTT) and ITT (IP-ITT) were administered to determine the effect of the diets on glucose tolerance and insulin action. In line with weight gains, IP-GTT at 7 weeks on diets demonstrated that glucose intolerance and insulin resistance increased from SC < LFD < HFD < CAF (least resistant to most resistant) (Figure 3a,b), and insulin levels were significantly elevated for all diets relative to SC (Figure 3c). IP-ITT performed at 11 weeks on diets demonstrated that the insulin-mediated drop in glucose was not significantly different between diet exposures (Figure 3d,e).
Obese humans with metabolic syndrome accumulate massive fat depots, which are shown to be chronically inflamed (18). To determine whether adiposity in the rodent models of obesity described here parallel physiologic indications of metabolic syndrome described above, we measured eWAT and BAT fat pad masses. CAF diet induced nearly a tripling of epididymal white and brown adipose mass compared to SC controls. HFD induced no change relative to LFD, but both were significantly different than SC and CAF (Figure 4a). Similar results were found for omental fat in an alternate set of animals after 10 weeks on experimental diets (data not shown). Elevations in plasma nonesterified fatty acids, another hallmark of metabolic syndrome, paralleled only in CAF-fed adipose. Weisberg et al. and Xu et al. demonstrated that macrophages present in the SVF contribute to macrophage infiltration and inflammation in cafeteria diet (CAF)-fed rodents. (a–h) ×10 and ×20 hematoxylin and eosin (H&E) stain of eWAT from (a,e) low-fat diet (LFD), (b,f) high-fat diet (HFD), (c,g) standard chow (SC), and (d,h) CAF. The box indicates areas of interest magnified to ×20 imaging when appropriate. Asterisks indicate crown-like structures (CLS), or areas of macrophage infiltration around adipocytes. Arrow indicates inflammatory loci. (i–l) Macrophage marker F4/80 was used to identify CLS (*) and areas of cells aggregating within inflammatory loci (arrow) (×40). (m) Quantification of CLS revealed a significant twofold increase in HFD- vs. LFD-fed adipose whereas CAF-fed adipose displayed a 17-fold increase over SC. (*CAF P < 0.05 vs. SC, **HFD P < 0.005 vs. all) (n) Pooled stromal vascular fractions (SVF) from eWAT enriched in adipose-associated macrophages demonstrated higher expression of tumor necrosis factor α (TNFα) normalized to 18S in CAF-fed rats when compared to SC control animals as determined by qPCR. Representative images shown. (n = 4, SC, LFD, and HFD, n = 5 CAF). See Supplementary Table S4 online for detailed description.
Little is known about macrophage infiltration into BAT other than increased macrophage infiltration of BAT in a transgenic lipodystrophy model (25). We investigated macrophage infiltration of BAT and indeed, compared to the normal architecture of SC-fed rats, increasing obesity causes enlarged fat droplets around vasculature in BAT with LFD and HFD amassing intermediate lipid droplets, and CAF diet inducing substantial fat accumulation. Only CAF-fed rats demonstrated large fat droplet accumulation in the BAT parenchyma extending from the vasculature (Figure 6a–h). This fat deposition is distinct from the WAT surrounding BAT (Figure 7m–o).

**CAF-diet induced hepatosteatosis and inflammation**

To investigate the effect of the human unhealthy food diet on the formation of nonalcoholic fatty liver disease (NAFLD), we evaluated serial sections of livers from each diet group. H&E staining of liver sections demonstrated normal liver structure in SC controls with some macrovesicular fat accumulation in LFD-fed rats. HFD and CAF-fed rat livers display minor to major microvesicular steatosis, respectively (Figure 7b and d, respectively). Steatosis associated with HFD was mixed macro- and microvesicular, localized to the periportal (Figure 7j) to midzonal regions and distinctly absent in the pericentral regions where normal tissue architecture was conserved (Figure 7f). Conversely, the CAF diet invoked hepatosteatosis that encompassed the entire liver lobule and was microvesicular in nature (Figure 7h,l). Next, we evaluated the ability of each diet to promote NASH by visually documenting F4/80 positive macrophage/Kupffer cell content and quantifying inflammatory loci, a collection of inflammatory cells primarily consisting of macrophages. LFD- and SC-fed rat livers displayed zero to very few inflammatory loci (Figure 7m,o, respectively) and q). Although livers from rats fed HFD and CAF diets contained increased inflammatory loci vs. controls, there were dramatic differences in the size and localization of loci between HFD and CAF animals (Figure 7n,p, respectively) and q). In livers from HFD-fed rats, F4/80 positive macrophages were small and limited to perivascular regions;
whereas loci from CAF-fed rats were found in both perivascular regions and within intralobular regions and were notably larger compared to HFD-fed animals (Figure 7).

**CAF-diet induced islet dysfunction**

Significantly elevated glucose and insulin levels and evidence of insulin resistance by GTT and ITT in the CAF animals led us to examine the pancreata by immunohistochemistry. LFD- and SC-fed rats displayed normal pancreatic islets with defined borders, whereas HFD-fed rats contained slightly hypertrophic and disfigured islets. Islets from CAF-fed rats were much larger in size, with dramatically distorted architecture (see Supplementary Figure S2 online).

**DISCUSSION**

**CAF: A model for human obesity and metabolic syndrome**

DIO is a common tool used to examine metabolic syndrome in rodents. The study presented herein was a comprehensive physiologic and histologic analysis between a commonly used model for DIO (a lard-based HFD), an alternative dietary model of energy dense highly palatable cafeteria-style foods and controls fed LFD and SC. The CAF diet provides a highly relevant model in terms of examining the human diet in rodents and comes at a time when the poor eating habits of Americans are in the spotlight. Current worldwide increases in obesity cannot be accounted for by changes in genetic backgrounds alone. Therefore, diet combined with previous evolutionary selective pressure for a “thrifty,” energy-conserving phenotype from when humans survived in a feast or famine world is an obvious contributing factor to this epidemic (26). Recent NHANES reports demonstrate that 33.6% of the American diet is comprised of fat (%kcal) (27). A study by the US Department of Agriculture on the US diet indicated that ~16% of Americans had overly poor diets and that 74% needed to improve their diets, based on the US Department of Agriculture recommended daily allowances (RDA) for grains, vegetables, fruits, milk, meat, total fat, saturated fat, cholesterol, sodium, and dietary variety (28). Current research indicates that childhood obesity and dietary intake statistics also follow these trends. Increased snacking on energy dense foods parallels the dramatic increase in childhood obesity, with unhealthy snacks currently comprising >27% of an American child’s daily caloric intake (29).

With these striking dietary trends in mind, we embarked on a study to examine a human diet in rodents compared to the traditional rodent diet most researchers use to model DIO. Our study has revealed that rats fed human nutrient-poor foods develop severe metabolic syndrome which is more robust than the effect of traditional HFD exposure. CAF-diet fed rats exhibited voluntary hyperphagia and grossly elevated fat intake which resulted in dramatic and rapid weight gain. Furthermore, CAF diet feeding promoted a prediabetic condition with elevated glucose, insulin, and nonesterified fatty acids, accompanied by decreased glucose and insulin tolerance. In addition, chronically inflamed liver and adipose tissues and distorted pancreatic islet architecture were prevalent.

**Choice of control diet in animal studies**

Interestingly, our results show that HFD did not induce a significant weight gain compared to LFD controls for up to 15 weeks of this diet. This suggests that either a longer diet is necessary to develop overt metabolic syndrome on the traditional high-fat lard-based diet or that the diet was started at an age that was too late to induce a divergent phenotype in our rats. It is likely that if we extended the study further or perhaps started in younger rats, a larger difference between the LFD and HFD could have emerged. An alternative explanation is that compared to standard chow SC, the LFD is very high in sucrose, with a simple exchange of fat for sugar to engineer the lard-based HFD. Indeed, compared to the SC, the LFD caused weight gain and moderate insulin resistance despite the fact that animals consumed relatively similar grams of food and fat calories. One reason could be that SC is undefined and contains carbohydrates derived from barley, oats, wheat, and corn rather than highly processed carbohydrate sources such as cornstarch, sucrose, and maltodextrin comprising the LFD diet. The whole-grains in SC contain phytoestrogens and high fiber which are protective against insulin resistance and obesity (30,31). In addition, the protein and fat sources vary between the SC groups. In sum, our varied results between SC and LFD chow “controls” demonstrate that chow diets vary widely in their effects, which emphasizes the relevance of chow sources and compositions, especially with respect to diet and obesity studies. A defined diet such as the LFD included in our study, which is commonly used as a control for HFD studies, is helpful in terms of knowing the composition but also has poor nutritional additives such as processed carbohydrates that can affect baseline measurements.

Interestingly, our study revealed that while HFD-fed animals decreased food intake in terms of total grams to maintain caloric intake comparable to LFD and SC control, CAF-fed animals lacked this autoregulatory mechanism. The highly palatable diet with varied flavors and textures seems to have disabled the normal homeostatic mechanism for energy balance possibly by activating the reward system (32). Energy-dense diets have been shown to alter normal appetite regulation (33,34). Another finding from this study is that nutritional components of the CAF diet have robust effects beyond the fat content alone. The CAF and HFD have similar fat contents at about 45–50% while their appropriate controls contained about 10–12% kcal derived from fat. Despite these relative similarities, the CAF-fed rats were hyperphagic, eating 40% more calories/day compared to the HFD. Although the CAF diet is high in fat, it is also high in sodium, saturated fatty acid, trans-fatty acids, and cholesterol and low in fiber and micronutrients such as vitamin A and calcium. All of these experimental components limit the interpretation of the CAF data such that no causal conclusions can be made as to the role or mechanism of a specific food component such as high-fat or high-sucrose exposure to weight gain associated with CAF diet. However, this experimental platform, which more accurately reflects early onset obesity and its associated pathologies, continues to be improved upon from its inception in 1982. Careful selection...
of unhealthy food components with known nutritional content could facilitate a more mechanistic approach to understanding the relevant nutrients associated with rapid weight gain and progression toward disease states such as metabolic syndrome, NASH and type II diabetes.

Severe adipose and liver inflammation in CAF

Histologic analysis revealed that CAF-fed rats developed pathologic abnormalities in epididymal white adipose, brown adipose, liver, and pancreas consistent with metabolic syndrome. Chronic low-grade systemic inflammation is now considered to be a key component of metabolic syndrome with white adipose known to secrete many relevant adipokines (19,21). Visceral, not subcutaneous, fat, is associated with metabolic syndrome in humans and rodents (35,36). It is well-established that macrophages infiltrate into obese adipose and are necessary and sufficient to perpetuate the obese state and insulin resistance (21). The CAF diet resulted in dramatic macrophage infiltration of eWAT and significantly increased the formation of CLS, hallmarks of inflammatory adipose. Histologic measures of inflammation were supported by quantitative PCR analysis of the macrophage-enriched SVF of eWAT that demonstrated increased expression of tumor necrosis factor α, a cytokine proven to be involved in insulin resistance and adipose lipolysis (24). In BAT, we discovered that large lipid droplets accumulate in the cells surrounding vessels, which correlated with weight gain in the LFD and HFD-fed rats, with CAF-fed BAT demonstrating the most profound lipid droplet accumulation. Importantly, this fat accumulation was distinct from any VAT that would be on the periphery of the BAT depot. Furthermore, we showed for the first time that the macrophage infiltration pattern of BAT differed from that of VAT. There were no CLS evident in BAT; however, infiltration of macrophages was detected in the perivascular region where large lipid droplet accumulation was prevalent. Currently, the role of perivascular macrophage accumulation in BAT is unclear.

Another contributing component of metabolic syndrome is fatty liver. NAFLD affects over one-third of the American population and ranges from hepatosteatosis to steatohepatitis, and could result in cirrhosis or hepatocellular carcinoma (37). NAFLD is strongly associated with visceral adiposity (38). Hepatosteatosis and steatohepatitis lead to liver insulin insensitivity and the inability to blunt gluconeogenesis, which is another contributing factor to metabolic syndrome (39). Our data indicate that Sc rats have healthy livers with no steatosis or inflammation and histologically normal hepatic cords and other architecture. Hepatosteatosis developed in LFD livers with macrovesicular accumulations noted around the portal triad regions. In HFD livers, increased steatosis surrounding portal triads progressed to a microvesicular phenotype while the central vein architecture was preserved. The most extensive steatosis and distorted cord architecture was found in CAF-fed livers with microvesicular steatosis evident spanning from the portal triads to the central veins (panlobular). Macrosteatosis is thought to be benign and reversible while microsteatosis is considered malignant and precedes liver fibrosis (40). Prolonged steatosis can often lead to steatitis or activation of inflammatory cells in the liver. NAFLD, which is recognized as the hepatic manifestation of metabolic syndrome, is thought to occur in a two-hit manner with lipid accumulation followed by chronic inflammation, such as occurs with obesity (41). Like in adipose, it is thought that hepatosteatosis promotes the Th1 proinflammatory response suggesting that Kupffer cells could be polarized toward a proinflammatory phenotype, contributing to insulin resistance (42). Kupffer cells are resident liver macrophages and the primary source of liver inflammatory cytokines (41). Several groups have recently demonstrated that Kupffer cell activation is necessary to produce hepatic insulin resistance (42,43). Large Kupffer cells are thought to localize in the perivascular zone and exhibit higher phagocytosis, protease activity, and inflammatory potential than smaller Kupffer cells in midzonal and pericentral regions (41). Interestingly, we detected numerous small loci of inflammatory cells present in HFD-fed livers, which accumulated around the portal triad regions. In the CAF-fed model however, inflammatory loci were larger and infiltrates were evident not only in the portal triad regions but also in the midzonal, intralobular regions. During steatosis, Kupffer cells display a diffuse localization whereas in NASH patients, Kupffer cells migrate to and aggregate around the pericentral regions (44). In the present study, the CAF liver phenotype with inflammatory loci both at the portal triads and within midzonal regions indicated macrophage migration from the portal triad to central vein and suggests the malignant progression from NAFLD to NASH.

Taken together, we have demonstrated in a comparative study that the CAF is a more robust model of metabolic syndrome than Lard-based HFD and that the rapid-onset of weight gain, obesity, multiorgan dysfunctions and pathologies observed in the CAF model more closely reflect the modern human condition of early onset obesity. The CAF diet, while labor intensive and “untraditional,” may be the best model to emulate modern human obesity trends.

SUPPLEMENTARY MATERIAL

Supplementary material is linked to the online version of the paper at http://www.nature.com/obry

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DISCLOSURE

The authors declared no conflict of interest.

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