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## Morbid Obesity Attenuates the Skeletal Abnormalities Associated with Leptin Deficiency in Mice

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### Abstract

Leptin-deficient *ob/ob* mice are morbidly obese and exhibit low total bone mass and mild osteopetrosis. In order to disassociate the skeletal effects of leptin deficiency from those associated with morbid obesity, we evaluated bone mass, architecture, gene expression, and indices of bone turnover in wild type (WT) mice, *ob/ob* mice fed *ad libitum* (*ob/ob*), and *ob/ob* mice pair fed to WT mice (pair-fed *ob/ob*). Mice were maintained at 32°C (thermoneutral) from 6 to 18 weeks of age to minimize differences in resting energy expenditure. *ob/ob* mice were heavier, had more abdominal white adipose tissue (WAT), and were hyperglycemic compared to WT mice. Femur length, bone mineral content (BMC) and bone mineral density, and midshaft femur cortical thickness were lower in *ob/ob* mice than in WT mice. Cancellous bone volume fraction was higher but indices of bone formation and resorption were lower in *ob/ob* compared to WT mice; reduced bone resorption in *ob/ob* mice resulted in pathological retention of calcified cartilage. Pair-fed *ob/ob* mice were lighter, had lower WAT, uterine weight, and serum glucose than *ob/ob* mice. Similarly, femoral length, BMC, and cortical thickness were lower in pair-fed *ob/ob* mice compared to *ob/ob* mice, as were indices of cancellous bone formation and resorption. In contrast, bone marrow adiposity, calcified cartilage and cancellous bone volume fraction were higher at one or more cancellous sites in pair-fed *ob/ob* mice compared to *ob/ob* mice. These findings demonstrate that the skeletal abnormalities caused by leptin deficiency are markedly attenuated in morbidly obese *ob/ob* mice.

### Keywords

obesity; histomorphometry; microcomputed tomography; dual energy absorptiometry

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## Introduction

Leptin, a polypeptide hormone secreted by adipocytes, is best known for its role in the regulation of appetite and energy metabolism (Mistry, et al. 1997; Myers 2004). Rodents deficient in leptin signaling, due to either an inability to generate leptin (*ob/ob* mice) or the signaling form of the leptin receptor (*db/db* mice and *fa/fa* rats), become morbidly obese (Clement 2000). Their excess weight is the result of a combination of hyperphagia and reduced thermogenesis (Hwa, et al. 1996). In addition to morbid obesity, leptin deficiency is associated with hypogonadism (Barkan, et al. 2005), elevated corticosteroid levels (Saito and Bray 1983), and impaired thermoregulation (Trayhurn and James 1978).

Leptin-deficient mice and leptin receptor-deficient mice and rats have reduced overall bone mass, reduced longitudinal bone growth (Kishida, et al. 2005; Steppan, et al. 2000), and decreased bone formation (Gat-Yablonski and Phillip 2008). In addition, impaired bone resorption in *ob/ob* and *db/db* mice results in mild osteopetrosis (Turner, et al. 2013), a condition which may contribute to the reported reductions in bone quality observed in these animals (Ealey, et al. 2006; Kimura, et al. 2012). Impaired bone resorption may also account for the age- and bone-dependent variation in cancellous bone volume observed in leptin signaling-deficient mice; the mutant mice are reported to have normal to increased cancellous bone volume fraction in lumbar vertebra and decreased to increased cancellous bone volume fraction in long bones (Hamrick, et al. 2004; Iwaniec, et al. 2007; Williams, et al. 2011). The skeletal abnormalities observed in mice and rats with defective leptin signaling suggest that leptin plays a role in normal skeletal growth, maturation, and turnover (Bartell, et al. 2011; Iwaniec et al. 2007; Kishida et al. 2005; Steppan et al. 2000; Turner et al. 2013). However, it is not clear to what extent the skeletal phenotype of *ob/ob* mice is impacted by additional factors associated with morbid obesity.

Although rare in humans, leptin signaling deficiency occurs as a result of loss of function mutations in the genes for leptin and the leptin receptor (Friedman and Halaas 1998). More common causes for low leptin signaling in humans include anorexia and starvation (Dardeno, et al. 2010; Muller, et al. 2009). At the other end of the spectrum, diet-induced leptin resistance, associated with chronically elevated leptin levels, mimics many of the metabolic consequences of leptin deficiency on energy metabolism and is thought to contribute to development of morbid obesity (Jung and Kim 2013). Thus, changes in leptin levels and/or leptin sensitivity could have physiological effects on bone metabolism.

In order to better understand the specific actions of leptin in regulating bone mass, density and architecture, we performed a study in which differences in body weight gain between rapidly growing wild type (WT) and *ob/ob* mice were prevented by housing all mice at thermoneutral temperature (32°C) to minimize differences in resting energy expenditure (Trayhurn 1979) and pair-feeding the mutant mice to WT mice to equalize food intake.

## Materials and Methods

### Experimental Design

The experimental protocols were approved by the Institutional Animal Care and Use Committee in accordance with the NIH Guide for the Care and Use of Laboratory Animals. Four-week-old female *ob/ob* (n=20) and homozygous WT (+/+) littermate (n=11) mice were purchased from Jackson Laboratory (Bar Harbor, ME, USA). At 6 weeks of age, all animals were transferred from normal room temperature (22°C) to thermoneutral temperature (32°C) and maintained at thermoneutrality on a 12-h light, 12-h dark cycle for the duration of study (12 weeks). Food (Teklad 8604, Harlan Laboratories, Indianapolis, IN) and water were provided *ad libitum* to WT mice (n=11) and a group of *ob/ob* mice (n=10). A second group of *ob/ob* mice (n=10) was pair-fed to the WT mice; the *ob/ob* mice were fed an amount of food equivalent to the group mean for WT mice.

Food consumption was measured daily in WT mice and weekly in the *ad libitum*-fed *ob/ob* mice. Body weight was recorded weekly in all mice. The fluorochrome declomycin (20 mg/kg, Sigma, St Louis, MO) was administered at study initiation and the fluorochrome calcein (20 mg/kg, Sigma, St Louis, MO) was administered 4 days and 1 day prior to sacrifice to label mineralizing bone. All mice were fasted overnight. The mice were then anesthetized with 2–3% isoflurane delivered in oxygen, bled by cardiac puncture, and glucose measured using a glucometer (Life Scan Inc, Milpitas, CA). Serum was collected and stored at –80°C for measurement of global markers of bone turnover. Uteri and abdominal white adipose tissue (WAT) were excised and weighed. Femora and lumbar vertebrae were removed and stored in 70% ethanol for analysis using dual energy absorptiometry (DXA), microcomputed tomography ( $\mu$ CT), and histomorphometry. Tibiae were removed, cleaned of soft tissue, frozen in liquid nitrogen, and stored at –80°C for mRNA isolation and gene expression analysis.

### Serum Chemistry

Serum osteocalcin was measured using mouse Gla-osteocalcin High Sensitive EIA kit from Clontech (Mountain View, CA). Serum CTx was measured using mouse CTx ELISA kit from Life Sciences Advanced Technologies (Saint Petersburg, FL).

### Dual Energy X-ray Absorptiometry

Femoral bone mineral content (BMC, mg), area ( $\text{cm}^2$ ), and bone mineral density (BMD,  $\text{g}/\text{cm}^2$ ) were determined *ex vivo* using DXA (Piximus, Lunar Corp., Madison, WI, USA).

### Microcomputed Tomography

$\mu$ CT was used for nondestructive three-dimensional evaluation of bone microarchitecture (Bouxsein, et al. 2010). Femora and lumbar vertebrae were scanned in 70% ethanol using a Scanco  $\mu$ CT40 scanner (Scanco Medical AG, Basserdorf, Switzerland) at a voxel size of  $12 \times 12 \times 12 \mu\text{m}$  (55 kV<sub>p</sub> x-ray voltage, 145  $\mu\text{A}$  intensity, and 200 ms integration time). Evaluations were conducted with filtering parameters sigma and support set to 0.8 and 1, respectively. Bone segmentation was performed at a threshold of 245 (scale, 0–1000) determined empirically. Total femur was evaluated followed by evaluation of cortical bone

in the midshaft femur and cancellous bone in the distal femur metaphysis and epiphysis. Automated contouring was used to delineate cortical bone from non-bone. Following, all cortical slices were examined visually for potential inclusion of cancellous struts originating from the endocortex (extremely rare at this site) and manually removed when present. For cortical bone, 20 slices (0.24 mm) of bone were evaluated, and cross-sectional tissue volume (cortical and marrow volume, mm<sup>3</sup>), cortical volume (mm<sup>3</sup>), marrow volume (mm<sup>3</sup>), and cortical thickness (μm) measured. Polar moment of inertia was determined as a surrogate measure of bone strength in torsion. Architectural parameters are expressed using standard 3-dimensional nomenclature. For the femoral metaphysis, 40 slices (480 μm) of cancellous bone were evaluated. The entire cancellous compartment was assessed in the femoral epiphysis. Analysis of lumbar vertebra included the entire region of cancellous bone between the cranial and caudal growth plates. Manual contouring was used to delineate cancellous from cortical bone in the femur metaphysis, femur epiphysis, and vertebral body. Cancellous bone measurements in femur and lumbar vertebra included bone volume fraction (bone volume/tissue volume, BV/TV, %), connectivity density (mm<sup>-3</sup>), trabecular number (mm<sup>-1</sup>), trabecular thickness (mm) and trabecular separation (mm) (Thomsen, et al. 2005).

### Histomorphometry

The histological methods used here have been described in detail (Iwaniec, et al. 2008). In brief, the distal femur was dehydrated in graded series of ethanol and xylene and embedded undecalcified in modified methyl methacrylate. Sections (4 μm thick) were cut with a vertical bed microtome (Leica/Jung 2165) and affixed to slides precoated with a 1% gelatin solution. Sections were mounted unstained for measurement of fluorochrome labels. For cell-based measurements, sections were stained for tartrate resistant acid phosphatase and counterstained with toluidine blue (Sigma, St Louis). All data were collected using the OsteoMeasure System (OsteoMetrics, Inc., Atlanta, GA). The sampling site for the distal femoral metaphysis was located 0.25–1.25 mm proximal to the growth plate. The entire cancellous compartment was evaluated in the distal femoral epiphysis and body of lumbar vertebra.

Fluorochrome-based measurements of bone formation included: (1) mineralizing perimeter (mineralizing perimeter/bone perimeter: cancellous bone perimeter covered with double plus half single label normalized to bone perimeter, %), (2) mineral apposition rate (the distance between two fluorochrome markers that comprise a double label divided by the 3 day interlabel interval, μm/day), and (3) bone formation rate (bone formation rate/bone perimeter: calculated by multiplying mineralizing perimeter by mineral apposition rate normalized to bone perimeter, μm<sup>2</sup>/μm/y).

Declomycin label retained in the regions of interest (mm label/mm<sup>2</sup> tissue area) in the distal femur epiphysis and lumbar vertebra was measured as a dynamic index of bone resorption in *ob/ob* and pair-fed *ob/ob* mice. This method is based on the principle that declomycin label was incorporated into bone at equivalent rates in both groups of *ob/ob* mice when the fluorochrome was administered prior to treatment initiation and differences measured at study termination reflect the effect of treatment on resorption of the fluorochrome-labeled bone. This method has been described (Westerlind, et al. 1997). We did not measure

declomycin label in WT mice because almost no label was retained in this group at termination of study.

Cell-based measurements included osteoblast perimeter, osteoclast perimeter, adipocyte number, and adipocyte area. Osteoblast perimeter was determined as a percentage of total bone perimeter lined by a palisade of plump cuboidal cells located immediately adjacent to the thin layer of osteoid in direct physical contact with the bone perimeter (osteoblast perimeter/bone perimeter, %). Osteoclast perimeter was determined as the percentage of cancellous bone perimeter covered by multinucleated (two or more nuclei) cells with acid phosphatase positive (red-stained) cytoplasm (osteoclast perimeter/bone perimeter, %). Cartilage area in the distal femur metaphysis and epiphysis was measured and expressed as % cancellous bone area. The intense metachromatic (purple) staining of proteoglycans and glycosaminoglycans with toluidine blue was used to identify cartilage within trabeculae. Adipocyte number and area were also measured and expressed as bone marrow adiposity (tissue area occupied by adipocytes: adipocyte area/tissue area, %), and adipocyte density (number of adipocytes/mm<sup>2</sup>). Adipocytes were identified morphologically as large (> 75 μm<sup>2</sup>) circular or oval-shaped cells bordered by a prominent cell membrane and lacking cytoplasmic staining due to alcohol extraction of intracellular lipids during processing. This method was previously validated by fat extraction and analysis (Menagh, et al. 2010). All histomorphometric data are expressed using standard 2-dimensional nomenclature (Dempster, et al. 2013) to distinguish 2-dimensional histomorphometric measurements from 3-dimensional μCT measurements.

### Gene Expression

Tibiae were pulverized with a mortar and pestle in liquid nitrogen and then further homogenized in Trizol (Invitrogen; Valencia, CA). Total RNA was isolated according to the manufacturer's protocol, and mRNA was reverse transcribed into cDNA using SuperScript III First-Strand Synthesis SuperMix for qRT-PCR (Invitrogen). The expression of 84 genes related to osteogenic differentiation was determined using the Mouse Osteogenesis RT<sup>2</sup> Profiler™ PCR Array (Qiagen; Carlsbad, CA) according to the manufacturer's protocol. Gene expression was normalized to GAPDH and relative quantification was determined using the Ct method using RT<sup>2</sup> Profiler PCR Array Data Analysis software version 3.5 (Qiagen).

### Statistics

Mean responses of individual variables were compared between the WT, *ob/ob*, and pair-fed *ob/ob* groups using separate one-way analysis of variance (ANOVA), with Tukey's procedure for pairwise multiple comparisons. The required conditions for valid use of ANOVA models were assessed using Levene's test for homogeneity of variance, plots of residuals versus fitted values, normal quantile plots, and the Anderson-Darling test of normality. A modified F test was used when the assumption of equal variance was violated, with Welch's two-sample t-test used for pairwise comparisons (Welch 1951). The Kruskal-Wallis nonparametric test was used when only the normality assumption was violated, in which case the Wilcoxon-Mann-Whitney test was used for pairwise comparisons. Longitudinal data on body weight were analyzed using a random intercept linear mixed

model that allowed distinct linear rates of change for the WT and pair-fed *ob/ob* groups and a quadratic time trend for the *ob/ob* group. The Hommel method for maintaining the family-wise error rate at 5% was used to adjust for multiple comparisons (Hommel 1988). Data analysis was performed using R version 2.12. Gene expression analysis was performed using RT<sup>2</sup> Profiler PCR Array Data Analysis version 3.5 (Qiagen). Gene expression is reported as mean fold change. All other data are reported as mean  $\pm$  SE.

## Results

The effects of pair feeding (to match food intake in WT mice) on body weight and cumulative food intake, and on abdominal WAT weight, blood glucose levels, uterine weight, and serum osteocalcin and CTx levels at sacrifice in leptin-deficient *ob/ob* mice housed at thermoneutral temperature are shown in Figure 1. Six-week-old *ob/ob* mice were heavier than their age-matched WT littermates at start of treatment and gained more weight than the WT mice during the 12-week study (Figures 1A and B). *ob/ob* mice were hyperphagic consuming, on average, twice as much food as WT mice (Figure 1C). At termination of study, the *ob/ob* mice had more abdominal WAT (Figure 1D), and were hyperglycemic (Figure 1E) and hypogonadal (Figure 1F) compared to WT mice.

Pair-fed *ob/ob* mice had lower body weight and body weight gain, lower cumulative food intake, lower abdominal WAT weight, lower blood glucose levels, and slightly lower uterine weight than *ob/ob* mice. Pair-feeding abolished differences in weight gain and blood glucose levels between WT and *ob/ob* mice. However, the pair-fed *ob/ob* mice were still heavier, had more WAT, and were hypogonadal compared to WT mice.

### Effects of pair feeding on serum markers of bone turnover in *ob/ob* mice

Serum levels of osteocalcin (Figure 1G) and CTx (Figure 1H) were lower in *ob/ob* mice than in WT mice. Also, pair-fed *ob/ob* mice had lower serum osteocalcin levels than either WT or *ob/ob* mice. Insufficient serum was available for measurement of CTx in pair-fed *ob/ob* mice.

### Effects of pair feeding on bone mass and architecture in *ob/ob* mice

The effects of pair feeding on femur BMC, area and BMD, and on femur length, bone volume, and cortical and cancellous bone architecture in *ob/ob* mice are shown in Table 1. *ob/ob* mice had lower femur BMC, bone area, and BMD than WT mice. Pair-fed *ob/ob* mice had lower BMC and bone area than WT or *ob/ob* mice. Significant differences in BMD were not detected between *ob/ob* and pair-fed *ob/ob* mice.

**Femur length and bone volume**—*ob/ob* mice had shorter femurs and lower femoral bone volume than WT mice. Pair-fed *ob/ob* mice had lower femur length and lower total femur bone volume than WT or *ob/ob* mice.

**Cortical bone in midshaft femur**—*ob/ob* mice had lower cortical bone volume and cortical thickness than WT mice. Pair-fed *ob/ob* mice had lower midshaft femur cortical bone volume and cortical thickness, and greater marrow volume than WT or *ob/ob* mice.



Significant differences in cross-sectional volume and polar moment of inertia were not detected among groups.

**Cancellous bone in distal femur metaphysis**—*ob/ob* mice had higher cancellous bone volume fraction, connectivity density, and trabecular number, and lower trabecular separation than WT mice. Differences in trabecular thickness were not detected between the two groups. Pair-fed *ob/ob* mice had higher cancellous bone volume fraction and trabecular number, and lower trabecular separation than WT mice or *ob/ob* mice, and higher connectivity density than WT mice. Trabecular thickness in pair-fed *ob/ob* mice was higher than in *ob/ob* mice but did not differ from WT mice.

**Cancellous bone in distal femur epiphysis**—*ob/ob* mice and pair-fed *ob/ob* mice had higher bone volume fraction, connectivity density, and trabecular number, and lower trabecular thickness and separation than WT mice. Differences between *ob/ob* mice and pair-fed *ob/ob* mice were not detected for any epiphyseal endpoint measured.

The effects of pair feeding on cancellous bone mass and architecture in lumbar vertebra in *ob/ob* mice are shown in Table 1. *ob/ob* mice had higher cancellous bone volume fraction, connectivity density, and trabecular number, and lower trabecular separation than WT mice. Pair-fed *ob/ob* mice had higher bone volume fraction than WT mice and a tendency ( $p < 0.1$ ) for higher bone volume fraction than *ob/ob* mice. Also, pair-fed *ob/ob* mice had higher trabecular thickness than WT or *ob/ob* mice. Connectivity density was lower in pair-fed *ob/ob* mice than in *ob/ob* mice but did not differ from WT mice. Trabecular number was higher and trabecular separation was lower in pair-fed *ob/ob* mice than WT mice. Differences in trabecular number and separation were not detected between *ob/ob* mice and pair-fed *ob/ob* mice.

### Effects of pair feeding on bone histomorphometry in *ob/ob* mice

**Cancellous bone in distal femur metaphysis**—The effects of pair feeding on dynamic and static cancellous bone histomorphometry and marrow adiposity in the distal femur metaphysis in *ob/ob* mice are shown in Figure 2. *ob/ob* mice had lower mineralizing perimeter (Figure 2A), mineral apposition rate (Figure 2B), bone formation rate (Figure 2C), and osteoblast perimeter (Figure 2D) compared to WT mice. Pair-fed *ob/ob* mice had lower mineralizing perimeter, mineral apposition rate, bone formation rate, and osteoblast perimeter than WT or *ob/ob* mice. Differences in osteoclast perimeter (Figure 2E) were not detected with treatment. However, cartilage area (Figure 2F), an index of impaired bone resorption, was higher in both groups of *ob/ob* mice compared to WT mice. Bone marrow adiposity (Figure 2G) tended ( $p < 0.1$ ) to be greater in *ob/ob* mice than WT mice and was greater in pair-fed *ob/ob* mice than in WT or *ob/ob* mice. Adipocyte density (Figure 2H) was higher in pair-fed *ob/ob* mice than in *ob/ob* mice. Representative photomicrographs of marrow adipocytes in the 3 groups of mice are shown in Figure 3A–C.

**Cancellous bone in distal femur epiphysis**—The effects of pair feeding on dynamic and static cancellous bone histomorphometry and marrow adiposity in the distal femur epiphysis in *ob/ob* mice are shown in Figure 4. *ob/ob* mice had lower mineralizing perimeter

(Figure 4A), bone formation rate (Figure 4C), and osteoblast perimeter (Figure 4D) than WT mice. Pair-fed *ob/ob* mice had lower mineralizing perimeter, bone formation rate, and osteoblast perimeter than WT or *ob/ob* mice. Osteoclast perimeter (Figure 4E) was higher in *ob/ob* mice and pair-fed *ob/ob* mice than WT mice. Retention of declomycin label (Figure 4F), an index of decreased bone resorption, was higher in pair-fed *ob/ob* mice than in *ob/ob* mice. *ob/ob* mice had greater cartilage area (Figure 4G), marrow adiposity (Figure 4H), and adipocyte density (Figure 4I) than WT mice. Pair-fed *ob/ob* mice had higher cartilage area, marrow adiposity, and adipocyte density than WT or *ob/ob* mice. Representative photomicrographs of cartilage and declomycin label in the 3 groups are shown in Figure 3D–F and G–I, respectively.

**Cancellous bone in lumbar vertebra**—The effects of pair feeding on dynamic and static cancellous bone histomorphometry and on bone marrow adiposity in lumbar vertebra are shown in Figure 5. *ob/ob* mice had lower mineralizing perimeter (Figure 5A), bone formation rate (Figure 5C), and osteoblast perimeter (Figure 5D) than WT mice. Pair-fed *ob/ob* mice had lower mineralizing perimeter, mineral apposition rate (Figure 5B), bone formation rate, and osteoblast perimeter than WT or *ob/ob* mice. Significant differences in osteoclast perimeter (Figure 5E) were not detected among the 3 groups. However, retention of declomycin label (Figure 5F) was higher in pair-fed *ob/ob* mice than in *ob/ob* mice. *ob/ob* mice had greater marrow adiposity (Figure 5G) and adipocyte density (Figure 5H) than WT mice. Pair-fed *ob/ob* mice had higher marrow adiposity and adipocyte density than WT or *ob/ob* mice.

### Effects of pair feeding on gene expression in tibia

The effects of pair feeding on expression of 84 genes related to osteogenesis in tibia are shown in Figure 6. Compared to WT mice, expression levels of 24 genes were altered in *ob/ob* mice (7 higher and 17 lower) and 26 genes in pair-fed *ob/ob* mice (10 higher and 16 lower). The expression levels of 14 genes were altered in common in *ob/ob* and pair-fed *ob/ob* mice. Compared to *ob/ob* mice, expression levels of 11 genes (3 higher and 8 lower) were altered in pair-fed *ob/ob* mice. *ob/ob* mice exhibited reductions in several key genes related to osteoblast and osteoclast differentiation and function, including *Sp7/osterix* (–2.5 fold), *NfκB1* (–1.2 fold), alkaline phosphatase (*Alpl*) (–1.5 fold), *Itgam* (*CD11b*) (–1.3 fold) and osteocalcin (*Bglap*) (–1.8 fold). *ob/ob* mice also exhibited reductions in genes related to cartilage formation and maturation, including *col2a1* (–1.5 fold) and *col10a1* (–1.7 fold). Genes whose expression were elevated include *Bmpr1a* (+1.3 fold) and *Smad4* (+1.3 fold). Compared to WT mice, pair-fed *ob/ob* mice expressed even lower levels of osteocalcin (–3.8 fold) than observed in *ob/ob* mice and, in addition, had lower mRNA levels for the osteoclast differentiation factor *Csf1* (–1.5 fold) and osteoblast differentiation factor *Runx2* (–1.4 fold). Furthermore, compared to WT mice, pair-fed *ob/ob* mice had higher gene expression levels for the cytokines/growth factors *TNFA* (+1.6 fold) and *TGFB2* (+1.3 fold).

## Discussion

We investigated the contribution of excessive weight gain to the skeletal phenotype of leptin-deficient *ob/ob* mice. Mice were housed at 32°C and a group of *ob/ob* mice was pair-



fed to WT mice. Housing mice at 32°C, which is within the thermoneutral range for WT and *ob/ob* mice, was expected to minimize differences in resting energy expenditure and pair feeding *ob/ob* mice to WT mice was expected to equalize energy intake (Knehans and Romsos 1982; Rafael and Herling 2000). The parallel increase in body weight gain in WT mice and *ob/ob* mice pair-fed to WT mice over the 12-week duration of study demonstrates the efficacy of this approach for preventing development of morbid obesity in leptin-deficient mice.

Consistent with earlier studies (Ealey et al. 2006; Hamrick et al. 2004; Iwaniec et al. 2007; Williams et al. 2011), *ob/ob* mice exhibited a mosaic skeletal phenotype associated with decreased bone formation due to reduced osteoblast number and activity, and defective osteoclast function. Femurs from *ob/ob* mice were smaller and had thinner cortices than those from WT mice. Furthermore, serum osteocalcin levels, mRNA levels for osteocalcin and SP7 (osterix), osteoblast perimeter and dynamic indices of bone formation were lower in *ob/ob* mice than in WT mice. However, cancellous bone volume fraction in both distal femur and lumbar vertebra was higher in the *ob/ob* mice. The higher cancellous bone volume fraction was associated with (1) greatly reduced serum CTx levels, (2) reductions in mRNA levels for NFκB, Itgam and Icam1, (3) retention of calcified cartilage, and (4) retention of fluorochrome label administered at study initiation, all indicative of a defect in bone resorption. Osteoclast perimeter was either unaffected (distal femur metaphysis and lumbar vertebra) or increased (distal femur epiphysis), indicating that reduced bone resorption in *ob/ob* mice was not due to an inability to generate osteoclasts. Taken together, these findings further support our previous observation that *ob/ob* mice exhibit mild osteopetrosis. Osteopetrosis due to a defect in osteoclast function may be responsible for the poor bone quality (Ealey et al. 2006) and defective tooth eruption (Batt 1978, 1992) reported in *ob/ob* mice.

Pair-feeding accentuated many of the abnormalities in bone size, mass, microarchitecture, gene expression, and turnover typically observed in *ob/ob* mice. Thus, chronic excessive weight gain typically present in *ob/ob* mice reduces the magnitude of skeletal effects associated with leptin deficiency. We did not adjust  $\mu$ CT-derived cortical bone data for body weight differences because the association between body weight and bone size and density in growing mice varies with bone and bone compartment and this association is altered by leptin (Iwaniec, et al. 2009). However, it is clear that the weight gain due to hyperphagia in *ob/ob* mice was associated with increases in cortical bone accrual but that these gains were not commensurate with the magnitude of body weight gain.

The mechanisms for the partial protective effect of morbid obesity on bone in *ob/ob* mice are likely to be multifactorial. We have previously reported a strong positive correlation between body weight and total femur bone mass in *ob/ob* mice (Iwaniec et al. 2009). This finding provides strong evidence that the higher total and cortical bone mass in *ad libitum*-fed compared to pair-fed *ob/ob* mice are, at least in part, attributable to higher body weight. Although there have been significant recent advances in our understanding of regulation of bone metabolism by mechanotransduction, the molecular mechanism for this positive effect of weight on bone mass is largely unknown. In addition to direct effects of increased

mechanical loading, changes in levels of hormones and/or growth factors may mediate additional effects of increased weight on the skeleton (Robling 2012).

We did not measure levels of insulin, glucocorticoids, or norepinephrine, factors known to be dysregulated in *ob/ob* mice (Dubuc, et al. 1985; Kim and Romsos 1990; Knehans and Romsos 1982; Rafael and Herling 2000) and known to influence bone metabolism. However, housing food-restricted *ob/ob* mice at thermoneutral temperature has been reported to reduce abnormalities in blood glucose, insulin, glucocorticoids, and norepinephrine turnover. In contrast, *ad libitum*-fed *ob/ob* mice were reported to remain hyperglycemic and hyperinsulinemic (Lindstrom 2007). We confirmed that blood glucose levels in pair-fed *ob/ob* mice did not differ from WT mice and were much lower than in *ad-libitum*-fed *ob/ob* mice. The contribution of excess insulin secretion and other factors associated with morbid obesity to the protective effects of excess body weight (Hickman and McElduff 1990; Ituarte, et al. 1988; Martineau-Doize, et al. 1986; Pun, et al. 1989; Yaturu 2009) on cortical bone microarchitecture in *ob/ob* mice requires further investigation.

Leptin is required for gonadotropin-releasing hormone secretion (Donato, et al. 2011). As a consequence, *ob/ob* mice are severely hypogonadal (Chehab, et al. 2002). The resulting deficiency in sex steroids would be expected to play a role in the pathological skeletal changes associated with leptin deficiency because sex steroids, especially estrogen, contribute to the sexual dimorphism of the female skeleton and regulate bone turnover balance in adults (Rickard, et al. 2008; Turner, et al. 1989; Turner, et al. 1994). Gonadal dysfunction typically results in elevated bone turnover and cancellous osteopenia. The difference between the skeletal responses to gonadal hormone insufficiency resulting from gonadectomy (high turnover) and from leptin-deficiency (low turnover) is perplexing. However, estrogen has a tonic effect on bone resorption and leptin is required for normal osteoclast function. As a consequence, high bone turnover generally associated with estrogen deficiency may be attenuated in *ob/ob* mice due to impaired osteoclast function. We speculate that the requirement for leptin in normal osteoclast function may serve as a counter regulatory mechanism to prevent excessive bone loss during fasting.

We did not measure sex steroid levels but the higher uterine weight in *ad libitum*-fed *ob/ob* mice compared to pair-fed *ob/ob* mice implies that morbid obesity results in a small increase in circulating estrogen. Higher estrogen levels in blood or adipose tissue may contribute to the protective skeletal effects of morbid obesity (Gennari, et al. 2011; Turner 1999; Turner et al. 1994). Aromatase, the enzyme responsible for conversion of androgens to estrogens, is expressed by adipocytes (Gennari et al. 2011). Thus, adipose-derived estrogen is a plausible mechanism contributing to the skeletal differences between pair-fed and *ad libitum*-fed *ob/ob* mice.

Initially heavier, *ob/ob* mice pair-fed to WT mice remained heavier than WT mice for the duration of study. We chose to equalize energy availability rather than body weight in the present study because bone accrual in rapidly growing rodents is strongly associated with energy availability and bone loss accompanies weight loss in adults (Devlin, et al. 2010; Turner and Iwaniec 2011). Low leptin levels induced by caloric restriction are associated with diminished bone mass accrual whereas partial leptin signaling deficiency induced in *ad*

*lib*-fed mice by treatment with a leptin antagonist increased body weight and bone mass accrual (Solomon, et al. 2014). However, administration of leptin increases bone formation in adult *ob/ob* mice in spite of weight loss (Bartell et al. 2011; Turner et al. 2013). Thus, both adequate energy availability and leptin play important roles in bone accrual in growing mice.

In contrast to peripheral fat depots, which act as dynamic energy reservoirs to maintain circulating triglyceride and free fatty acid levels during feeding and fasting, marrow fat increases with severe weight loss as well as weight gain (Bredella, et al. 2011; Devlin 2011). It is therefore unlikely that the bone marrow fat depot serves as an important energy reservoir for peripheral tissues. There is, however, evidence that bone marrow adipocytes function as negative regulators of hematopoietic lineage cell differentiation (Dorshkind, et al. 2009; Gimble, et al. 1996; Naveiras, et al. 2009). Bone marrow fat is increased in *ob/ob* mice and this increase is reversed by leptin treatment (Bartell et al. 2011; Hamrick, et al. 2005; Hamrick et al. 2004). Leptin stimulates hematopoiesis (Fantuzzi and Faggioni 2000; Trottier, et al. 2012), an action that may be due to inhibition of bone marrow adipogenesis, increased lipolysis, and direct stimulatory actions on hematopoietic cells. Adipocytes and osteoblasts are derived from the same bone marrow mesenchymal precursor and a reciprocal association between osteoblast number and adipocyte number has often been noted, giving rise to the hypothesis that bone marrow adipocytes may either antagonize osteoblast production or, alternatively, generation of adipocytes precludes generation of osteoblasts (Akune, et al. 2004). Other studies suggest that adipocyte and osteoblast number in bone marrow can be independently regulated (Iwaniec and Turner 2013; Menagh et al. 2010). Bartell et al. (2011) reported that intracerebroventricular administration of leptin to *ob/ob* mice increases expression of SP7, a transcription factor critical for osteoblast differentiation. Our findings of low expression of SP7 in *ob/ob* mice support the hypothesis that leptin acts to regulate bone formation by controlling mesenchymal cell decision.

In the present study, pair feeding to WT mice resulted in a further increase in bone marrow adiposity in leptin-deficient *ob/ob* mice. Of interest is our finding that pair-fed *ob/ob* mice had elevated mRNA levels for CD36; CD36 (fatty acid translocase) promotes adipocyte differentiation and adipogenesis (Christiaens, et al. 2012). The observed increase in CD36 may play a role in the increased bone marrow adiposity observed in the pair-fed *ob/ob* mice. Additional factors regulating bone marrow adiposity that may be influenced by leptin levels and obesity include growth hormone and IGF1. In rats, treatment with parathyroid hormone (whose actions on bone are largely mediated through IGF1) prevented the increase in bone marrow adiposity associated with severe energy restriction (Turner and Iwaniec 2011), whereas growth hormone treatment dramatically reduced excessive bone marrow adiposity in severely hypoleptinemic hypophysectomized rats (Menagh et al. 2010). In both of these studies, treatment had no effect on serum leptin levels. *ob/ob* mice, however, have decreased circulating growth hormone and impaired growth hormone signaling (Luque, et al. 2007), likely contributing to retention of fat within the bone marrow. High circulating insulin levels in morbidly obese *ob/ob* mice may further contribute to the suppression of growth hormone synthesis and release (Luque and Kineman 2006). However, IGF1 is produced and secreted by adipocytes (Kloting, et al. 2008) and higher levels of IGF1 are detected in obese subjects

(Frystyk, et al. 1999). Low leptin levels resulting from severe energy restriction are associated with increased growth hormone secretion, decreased circulating levels of IGF1, and end-organ resistance to growth hormone (Douyon and Scheingart 2002; Grinspoon, et al. 1995; Inagaki, et al. 2008; Munoz and Argente 2002; Thissen, et al. 1999). Thus, disturbances in growth hormone, IGF1 and insulin signaling as well as leptin deficiency may contribute to the increase in bone marrow adiposity in *ob/ob* mice. Chronic hyperphagia and excessive weight gain typically observed in *ob/ob* mice may partially compensate for leptin deficiency and reduced bone marrow adiposity by altering growth hormone and insulin levels and sensitivity.

In summary, leptin deficiency resulted in decreased femur length, BMC, BMD, and cortical bone volume and thickness. In contrast, cancellous bone volume fraction was increased in distal femur and lumbar vertebra. The profound changes in bone mass, density, and microarchitecture were associated with reduced linear bone growth, increased bone marrow adiposity, altered expression of osteogenic genes, decreased osteoblast number, and decreased osteoclast function. Prevention of morbid obesity in *ob/ob* mice further exacerbated many of the abnormalities associated with leptin deficiency. Because morbid obesity normally present in *ob/ob* mice attenuates the abnormal skeletal phenotype associated with leptin deficiency, the role of leptin in regulating postnatal bone growth, maturation, and turnover may have been underestimated.

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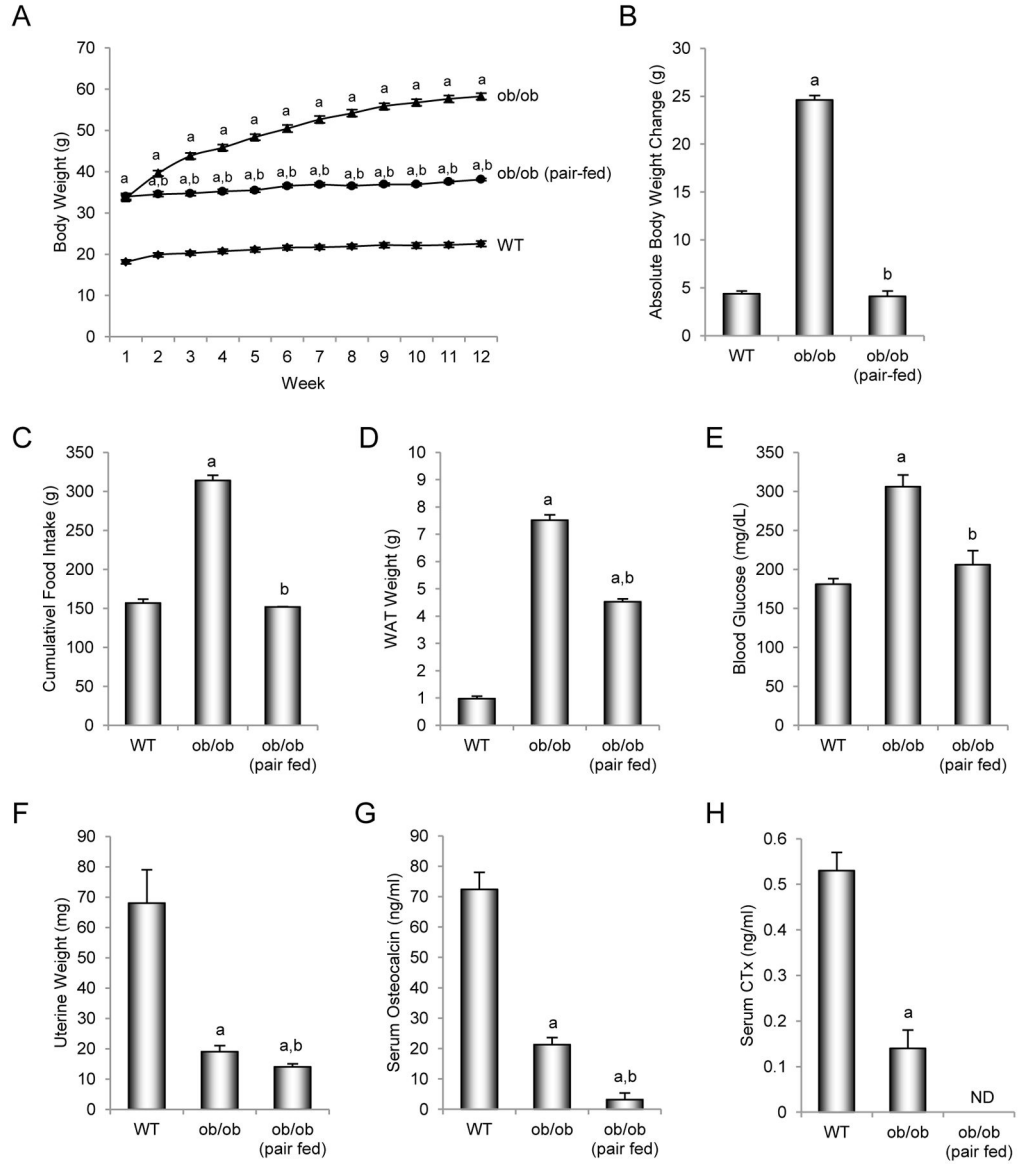
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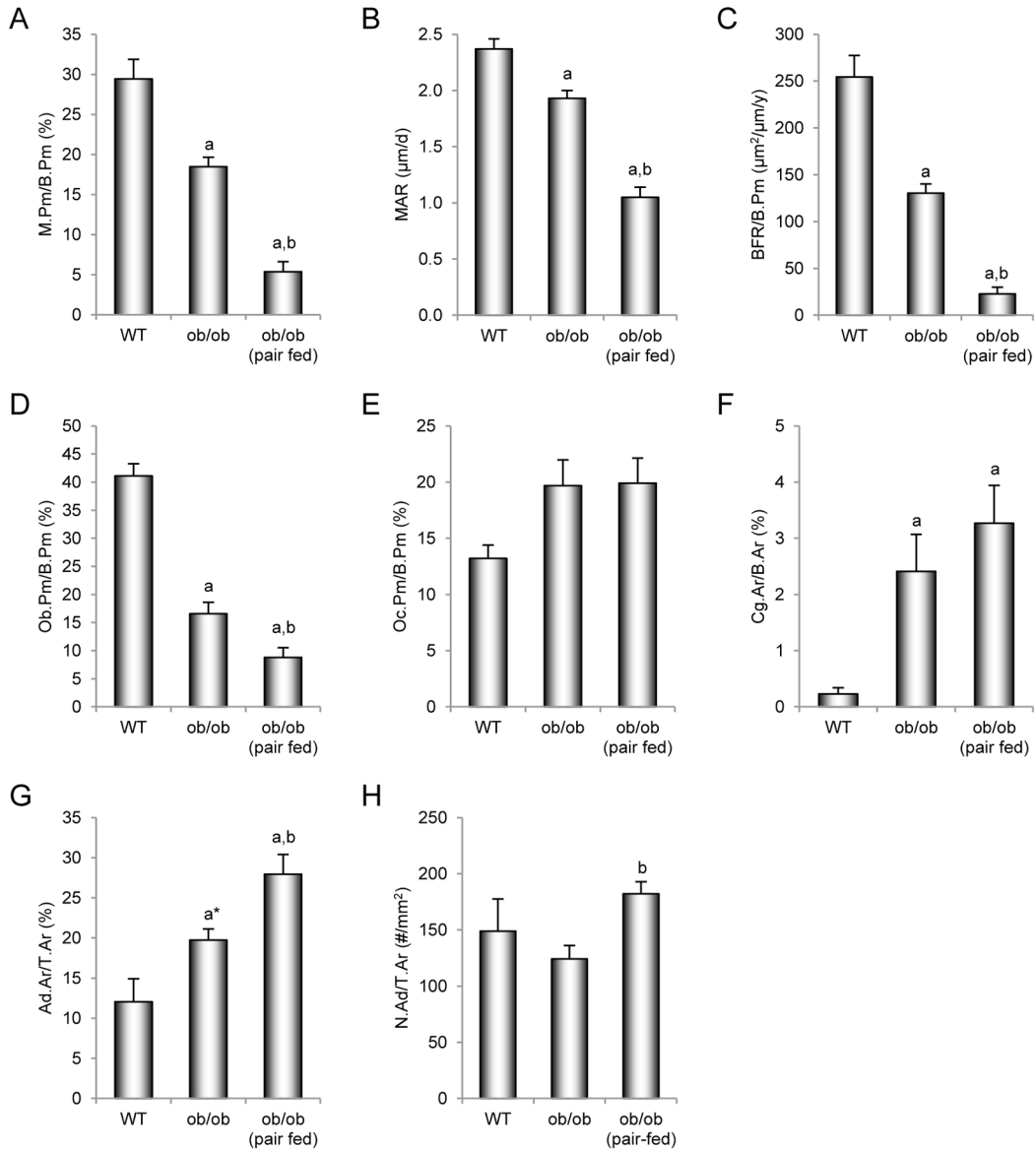
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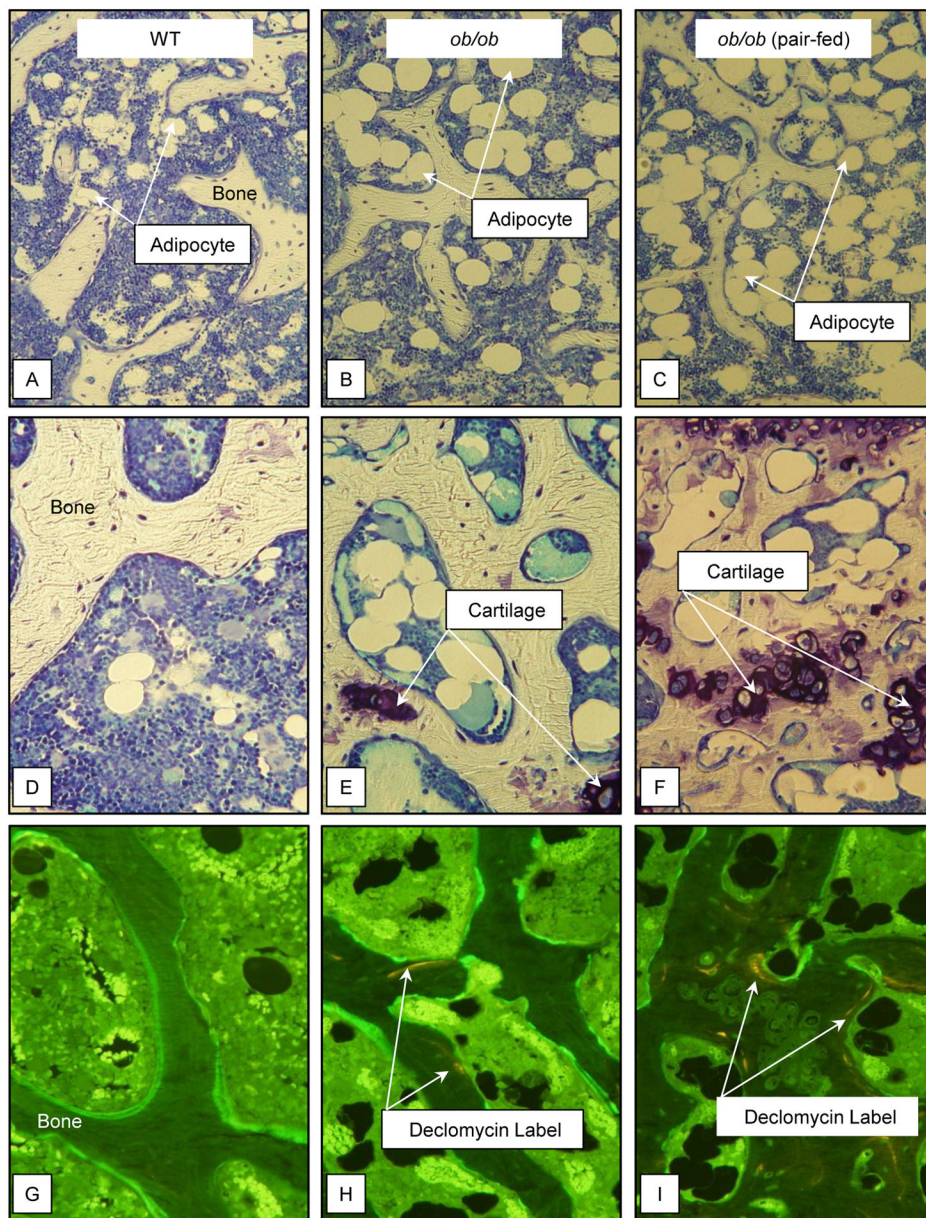
**Figure 1.**

Effects of morbid obesity on body weight (A), cumulative body weight change (B), cumulative food intake (C), abdominal white adipose tissue (WAT) weight (D), blood glucose (E), uterine weight (F), and serum osteocalcin (G) and CTx levels (H) in leptin-deficient *ob/ob* mice. Values are mean  $\pm$  SE, n=10–11/group. ND, no data due to insufficient serum for measurement of CTx in pair-fed *ob/ob* mice. <sup>a</sup>Different from WT mice,  $p < 0.05$ . <sup>b</sup>Different from *ob/ob* mice,  $p < 0.05$ .

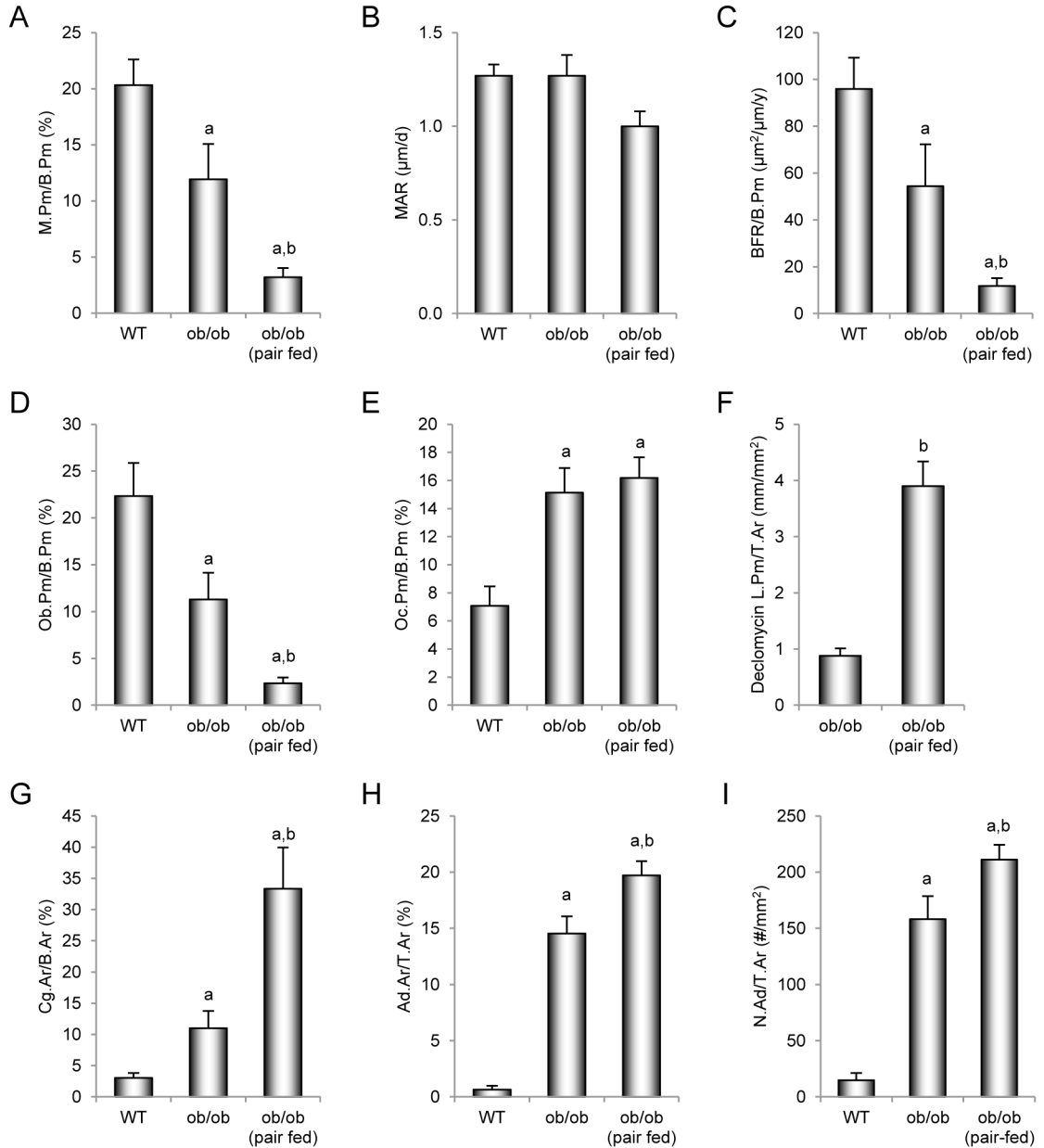


**Figure 2.**

Effects of morbid obesity on cancellous bone histomorphometry and marrow adiposity in distal femur metaphysis in leptin-deficient *ob/ob* mice. Shown are mineralizing perimeter (A, mineralizing perimeter/bone perimeter, M.Pm/B.Pm), mineral apposition rate (B, MAR), bone formation rate (C, bone formation rate/bone perimeter, BFR/B.Pm), osteoblast perimeter (D, osteoblast perimeter/bone perimeter, Ob.Pm/B.Pm), osteoclast perimeter (E, osteoclast perimeter/bone perimeter, Oc.Pm/B.Pm), cartilage area (F, cartilage area/bone area, Cg.Ar/B.Ar), marrow adiposity (G, adipocyte area/tissue area, Ad.Ar/T.Ar), and adipocyte density (H, number of adipocytes/tissue area, N.Ad/T.Ar). Values are mean  $\pm$  SE,  $n=10-11/\text{group}$ . <sup>a</sup>Different from WT mice,  $p < 0.05$ . <sup>a\*</sup>Different from WT mice,  $p < 0.1$ . <sup>b</sup>Different from *ob/ob* mice,  $p < 0.05$ .



**Figure 3.** Representative photomicrographs of 1) marrow adiposity in distal femur metaphysis in WT mouse (A), *ob/ob* mouse (B), and pair-fed *ob/ob* mouse (C), 2) cartilage remnants in distal femur epiphysis in WT mouse (D), *ob/ob* mouse (E), and pair-fed *ob/ob* mouse (F), and 3) declomycin label administered at study initiation in distal femur epiphysis in WT mouse (G), *ob/ob* mouse (H), and pair-fed *ob/ob* mouse (I).

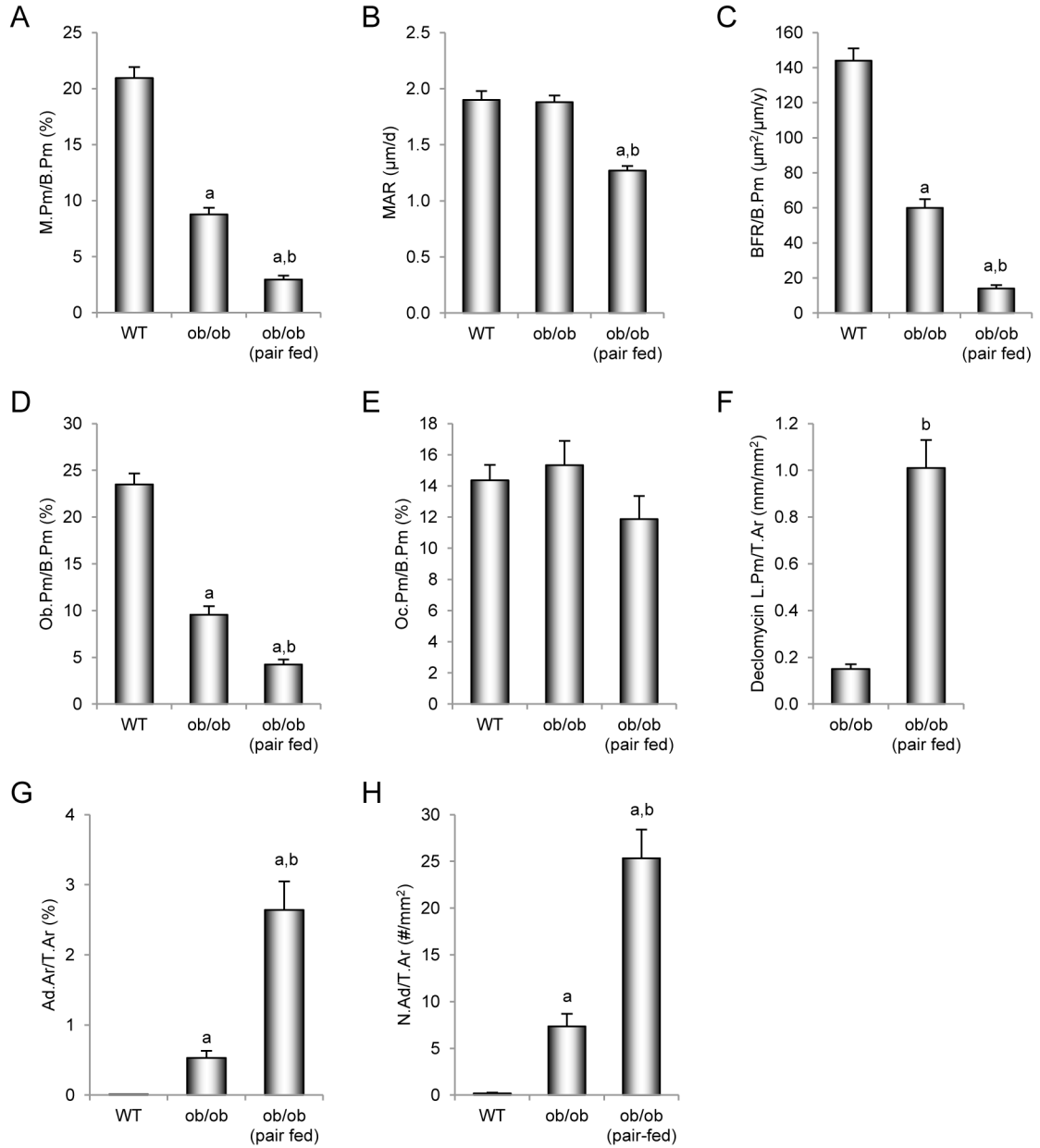


**Figure 4.**

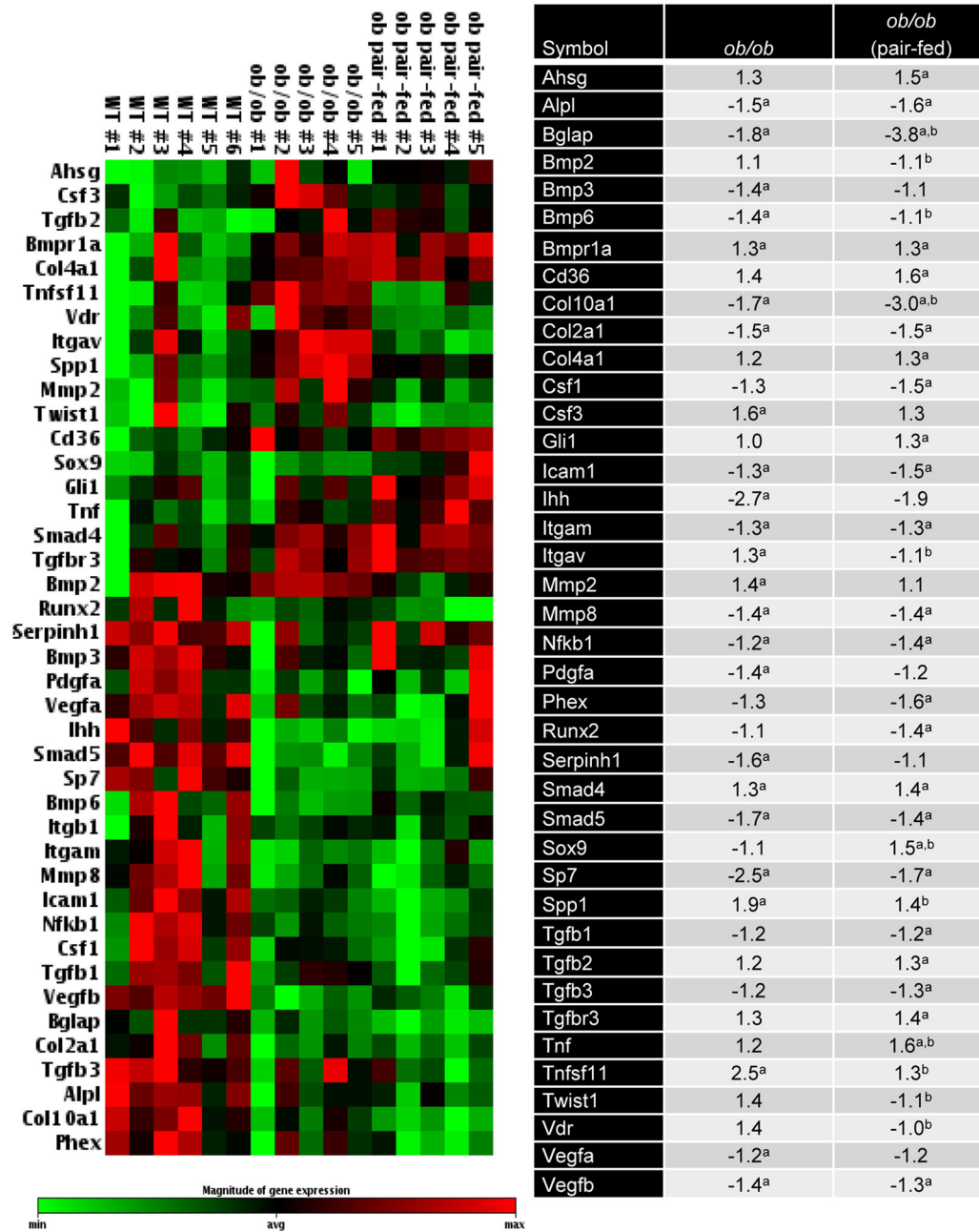
Effects of morbid obesity on cancellous bone histomorphometry and marrow adiposity in distal femur epiphysis in leptin-deficient *ob/ob* mice. Shown are mineralizing perimeter (A, mineralizing perimeter/bone perimeter, M.Pm/B.Pm), mineral apposition rate (B, MAR), bone formation rate (C, bone formation rate/bone perimeter, BFR/B.Pm), osteoblast perimeter (D, osteoblast perimeter/bone perimeter, Ob.Pm/B.Pm), osteoclast perimeter (E, osteoclast perimeter/bone perimeter, Oc.Pm/B.Pm), declomycin label perimeter (F, declomycin label perimeter/bone area, declomycin L.Pm/B.Ar), cartilage area (G, cartilage area/bone area, Cg.Ar/B.Ar), marrow adiposity (H, adipocyte area/tissue area, Ad.Ar/T.Ar), and adipocyte density (I, number of adipocytes/tissue area, N.Ad/T.Ar). Values are mean  $\pm$



SE, n=10–11/group. <sup>a</sup>Different from WT mice,  $p < 0.05$ . <sup>b</sup>Different from *ob/ob* mice,  $p < 0.05$ .

**Figure 5.**

Effects of morbid obesity on cancellous bone histomorphometry and marrow adiposity in lumbar vertebra in leptin-deficient *ob/ob* mice. Shown are mineralizing perimeter (A, mineralizing perimeter/bone perimeter, M.Pm/B.Pm), mineral apposition rate (B, MAR), bone formation rate (C, bone formation rate/bone perimeter, BFR/B.Pm), osteoblast perimeter (D, osteoblast perimeter/bone perimeter, Ob.Pm/B.Pm), osteoclast perimeter (E, osteoclast perimeter/bone perimeter, Oc.Pm/B.Pm), declomycin label perimeter (F, declomycin label perimeter/bone area, declomycin L.Pm/B.Ar), marrow adiposity (G, adipocyte area/tissue area, Ad.Ar/T.Ar), and adipocyte density (H, number of adipocytes/tissue area, N.Ad/T.Ar). Values are mean  $\pm$  SE, n=10–11/group. <sup>a</sup>Different from WT mice,  $p < 0.05$ . <sup>b</sup>Different from *ob/ob* mice,  $p < 0.05$ .



**Figure 6.**

Effects of morbid obesity on expression of osteogenesis-related genes in tibia. The expression of a panel of 84 osteogenesis-related genes in the tibia of WT mice (n=6), *ob/ob* mice (n=5), and *ob/ob* mice pair-fed to the WT mice (n=5) were determined using a mouse osteogenesis gene expression array. Left panel shows hierarchical clustering analysis of genes with significantly different gene expression. Magnitude of gene expression was represented by green and red bars, indicating decreased and increased expression, respectively. The normalized mean fold-change of *ob/ob* mice and pair-fed *ob/ob* mice

relative to the WT mice were shown in the right panel. <sup>a</sup>Different from WT mice,  $p < 0.05$ . <sup>b</sup>Different from *ob/ob* mice,  $p < 0.05$ .

**Table 1**

Effects of morbid obesity on femoral bone mineral content and density, femoral cortical and cancellous bone microarchitecture, and vertebral cancellous bone microarchitecture in leptin-deficient *ob/ob* mice.

End Point	WT	<i>ob/ob</i>	<i>ob/ob</i> (pair-fed)	ANOVA P*
<b>Dual Energy X-ray Absorptiometry</b>				
<b>Total Femur</b>				
BMC (g)	0.023 ± 0.000	0.019 ± 0.000 <sup>a</sup>	0.017 ± 0.000 <sup>a,b</sup>	0.0001
Bone area (cm <sup>2</sup> )	0.43 ± 0.01	0.40 ± 0.00 <sup>a</sup>	0.37 ± 0.01 <sup>a,b</sup>	0.0001
BMD (g/cm <sup>2</sup> )	0.053 ± 0.001	0.048 ± 0.000 <sup>a</sup>	0.047 ± 0.000 <sup>a</sup>	0.0003
<b>Microcomputed Tomography</b>				
<b>Total Femur</b>				
Length (mm)	14.9 ± 0.1	14.2 ± 0.1 <sup>a</sup>	13.4 ± 0.1 <sup>a,b</sup>	<0.0001
Bone volume (mm <sup>3</sup> )	18.5 ± 0.3	16.8 ± 0.3 <sup>a</sup>	15.0 ± 0.3 <sup>a,b</sup>	<0.0001
<b>Midshaft Femur (cortical bone)</b>				
Cross-sectional volume (mm <sup>3</sup> )	0.41 ± 0.01	0.40 ± 0.01	0.42 ± 0.01	0.4220
Cortical volume (mm <sup>3</sup> )	0.20 ± 0.00	0.18 ± 0.00 <sup>a</sup>	0.17 ± 0.00 <sup>a,b</sup>	0.0008
Marrow volume (mm <sup>3</sup> )	0.22 ± 0.00	0.23 ± 0.00	0.25 ± 0.01 <sup>a,b</sup>	0.0012
Cortical thickness (μm)	195 ± 3	176 ± 3 <sup>a</sup>	159 ± 2 <sup>a,b</sup>	0.0001
Polar moment of inertia (mm <sup>4</sup> )	0.36 ± 0.01	0.33 ± 0.01	0.33 ± 0.01	0.4220
<b>Distal Femur Metaphysis (cancellous bone)</b>				
Bone volume/tissue volume (%)	11.2 ± 0.8	13.2 ± 0.5 <sup>a</sup>	18.1 ± 0.5 <sup>a,b</sup>	0.0004
Connectivity density (1/mm <sup>3</sup> )	106 ± 9	173 ± 7 <sup>a</sup>	190 ± 8 <sup>a</sup>	<0.0001
Trabecular thickness (μm)	44 ± 1	41 ± 1	45 ± 0 <sup>b</sup>	0.0365
Trabecular number (mm <sup>-1</sup> )	4.8 ± 0.1	5.2 ± 0.1 <sup>a</sup>	5.7 ± 0.1 <sup>a,b</sup>	0.0001
Trabecular separation (μm)	210 ± 4	191 ± 4 <sup>a</sup>	173 ± 4 <sup>a,b</sup>	0.0013
<b>Distal Femur Epiphysis (cancellous bone)</b>				
Bone volume/tissue volume (%)	31.2 ± 0.6	36.1 ± 0.6 <sup>a</sup>	36.6 ± 0.9 <sup>a</sup>	0.0013
Connectivity density (1/mm <sup>3</sup> )	155 ± 4	254 ± 8 <sup>a</sup>	261 ± 7 <sup>a</sup>	<0.0001
Trabecular thickness (μm)	64 ± 1	60 ± 1 <sup>a</sup>	58 ± 1 <sup>a</sup>	0.0001
Trabecular number (mm <sup>-1</sup> )	5.5 ± 0.1	6.7 ± 0.1 <sup>a</sup>	6.7 ± 0.1 <sup>a</sup>	0.0004
Trabecular separation (μm)	179 ± 2	138 ± 3 <sup>a</sup>	138 ± 3 <sup>a</sup>	<0.0001
<b>Lumbar Vertebra (cancellous bone)</b>				
Bone volume/tissue volume (%)	21.8 ± 1.0	31.5 ± 0.8 <sup>a</sup>	34.2 ± 0.5 <sup>a,b*</sup>	<0.0001
Connectivity density (1/mm <sup>3</sup> )	198 ± 10	364 ± 15 <sup>a</sup>	191 ± 5 <sup>b</sup>	<0.0001
Trabecular thickness (μm)	49 ± 1	50 ± 0.5	54 ± 0.4 <sup>a,b</sup>	0.0010
Trabecular number (mm <sup>-1</sup> )	4.6 ± 0.1	6.0 ± 0.1 <sup>a</sup>	5.8 ± 0.1 <sup>a</sup>	0.0004
Trabecular separation (μm)	215 ± 5	160 ± 3 <sup>a</sup>	163 ± 3 <sup>a</sup>	0.0005

Data are mean ± SE

\* The Hommel method for maintaining the family-wise error rate at 5% was used to adjust for multiple comparisons

<sup>a</sup> Different from WT,  $p < 0.05$

<sup>b</sup> Different from *ob/ob*,  $p < 0.05$