

## ApoE Gene Deficiency Enhances the Reduction of Bone Formation Induced by a High-Fat Diet Through the Stimulation of p53-Mediated Apoptosis in Osteoblastic Cells

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**ABSTRACT:** Osteoblast apoptosis increased in the tibias of apoE<sup>-/-</sup> mice fed with a high-fat diet, decreasing bone formation. The expression of p53 mRNA in marrow adherent cells increased. LDL or oxidized LDL increased apoptosis in the calvarial cells of apoE<sup>-/-</sup> mice. The increase in p53-mediated apoptosis is apparently related to a high-fat diet-induced osteopenia in apoE<sup>-/-</sup> mice.

**Introduction:** The effects of high-fat loading and the *apolipoprotein E* (*apoE*) gene on bones have not been elucidated. We hypothesized that *apoE* gene deficiency (apoE<sup>-/-</sup>) modulates the effects of high-fat loading on bones.

**Materials and Methods:** We assessed this hypothesis using wildtype (WT) and apoE<sup>-/-</sup> mice fed a standard (WTS and ApoES groups) or a high-fat diet (WTHf and ApoEHf groups). The concentration of serum lipid levels and bone chemical markers were measured. Histomorphometry of the femurs was performed using  $\mu$ CT and a microscope. Bone marrow adherent cells from the femurs were used for colony-forming unit (CFU)-fibroblastic (CFU-f) assay and mRNA expressions analysis. The apoptotic cells in the tibias were counted. TUNEL fluorescein assay and Western analysis were performed in cultures of calvarial cells by the addition of low-density lipoprotein (LDL) or oxidized LDL.

**Results:** In the ApoEHf group, the values of cortical bone volume and trabecular and endocortical bone formation of the femurs decreased, and urinary deoxyypyridinoline increased. Subsequent analysis revealed that the number of apoptotic cells in the tibias of the ApoES group increased, and more so in the ApoEHf group. The ratio of alkaline phosphatase-positive CFU-f to total CFU-f was decreased in the ApoEHf group. p53 mRNA expression in adherent cells of the apoE<sup>-/-</sup> mice increased and had a significantly strong positive correlation with serum LDL. TUNEL fluorescein assay of osteoblastic cells revealed an increase of apoptotic cells in the apoE<sup>-/-</sup> mice. The number of apoptotic cells in the apoE<sup>-/-</sup> mice increased with the addition of 100  $\mu$ g/ml LDL or oxidized LDL. The p53 protein expression in apoE<sup>-/-</sup> cells exposed to 100  $\mu$ g/ml LDL or oxidized LDL increased.

**Conclusions:** We concluded that *apoE* gene deficiency enhances the reduction of bone formation induced by a high-fat diet through the stimulation of p53-mediated apoptosis in osteoblastic cells.

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**Key words:** apolipoprotein E, low-density lipoprotein, lipid peroxide, apoptosis, osteopenia

### INTRODUCTION

THE EFFECT OF lipid metabolism on bones has been shown in previous studies. For instance, inverse correlations were clinically observed between the levels of plasma total cholesterol and lumbar spine BMD<sup>(1)</sup> and those of plasma low-density lipoprotein (LDL) and BMD of the radius.<sup>(2)</sup> In vitro studies showed that oxidized LDL

inhibited the differentiation of osteoprogenitors into osteoblasts,<sup>(3,4)</sup> and an atherogenic high-fat diet reduced bone mineralization in mice.<sup>(5)</sup> Thus, it could be suggested that impairments of lipid metabolism, such as high serum levels of LDL and/or oxidized LDL, may decrease bone mass by reducing bone formation.

The effect of apolipoprotein E (apoE) on bones, on the other hand, has been controversial. ApoE is a structural component in the triglyceride-rich lipoprotein particles including very LDL (VLDL) and chylomicrons and their

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remnants. The apoE component is a ligand for the LDL receptor, consisting of three isoforms including E2, E3, and E4,<sup>(6)</sup> and belonging to the family of LDL receptor-related proteins (LRPs).<sup>(7)</sup> Accordingly, deficiency of the *apoE* gene (*apoE*<sup>-/-</sup>) in mice has a considerable effect on cholesterol transport; as a result, an increase in the plasma level of VLDL because of the failure of LDL receptor and LRP-mediated clearance from blood causes an increase in that of LDL.<sup>(8)</sup> The expression of the *apoE* gene has been observed in osteoblasts,<sup>(9)</sup> and some studies have shown that the apoE genotype is related to fracture risk and BMD.<sup>(10-14)</sup> However, others did not confirm this relationship.<sup>(15-18)</sup>

In rodents, the calcification of vessels did not occur in *apoE*<sup>-/-</sup> mice given a standard diet during the growth period.<sup>(8)</sup> When *apoE*<sup>-/-</sup> mice were given a high-fat diet, however, the vascular atherosclerotic lesions dramatically increased, as well as the plasma levels of LDL and VLDL.<sup>(19)</sup> These data indicate that the effect of *apoE*<sup>-/-</sup> on atherosclerosis in the vascular walls could depend on the amount of dietary fat given to the animals.

We hypothesized that *apoE*<sup>-/-</sup> modulates the effects of high-fat loading on bones. In this study, we examined the bone mass, turnover, and differentiation of osteogenic cells in four groups of mice, with or without the *apoE* gene, given a standard or high-fat diet during the age period of 5-17 wk. We also assessed the effects of *apoE*<sup>-/-</sup> and lipid loading on apoptosis of osteoblastic cells obtained from the calvariae of wildtype (WT) and *apoE*<sup>-/-</sup> mice.

## MATERIALS AND METHODS

### *Animal preparation*

Homozygous *apoE*<sup>-/-</sup> mice on a C57BL/6 background (N11) were obtained from Taconic (Hudson, NY, USA). Male mice homozygous for the *apoE*<sup>-/-</sup> were mated to female C57BL/6J mice (Clea Japan, Tokyo, Japan). Three hundred eggs of the subsequent heterozygous generation were collected, and after *in vitro* fertilization with heterozygous spermatozoa, the embryos were transferred into 50 foster mothers (ICR; Clea Japan), as described previously.<sup>(20)</sup> Using RT-PCR, the mice were genotyped at 3 wk of age to identify the homozygous *apoE*<sup>-/-</sup> and WT mice in the litter.

### *Experimental design*

Thirteen male *apoE*<sup>-/-</sup> and 15 WT littermate mice were used in this study. Those mice were weaned at 4 wk of age and acclimatized for 1 wk. At the age of 5 wk, the mice were fed a standard diet (1.25% calcium, 1.06% phosphorus, 2.0 IU/g vitamin D<sub>3</sub>; Clea Japan) or atherogenic high-fat diet (1.25% cholesterol, 7.5% cacao butter, 7.5% casein, 0.5% sodium cholate; Oriental Yeast Co., Tokyo, Japan), divided into four body weight-matched groups (WT mice fed with the standard diet, WTS group; WT mice fed with a high-fat diet, WTHf group; *apoE*<sup>-/-</sup> mice fed with the standard diet, ApoES group; *apoE*<sup>-/-</sup> mice fed with a high-fat diet, ApoEHf group) and maintained on the diet for 12 wk.

Food intake was assessed as food weight (g) per mouse per day and matched among the four groups. All mice were housed individually in cages.

At the age of 17 wk, urine samples were collected for 24 h before death. After exsanguination, the bilateral humeri, femurs, and tibias were harvested under pentobarbital sodium anesthesia. Bone labeling with intramuscular injection of calcein (8 mg/kg body weight) was performed at 7 and 3 days before death in the mice used for histomorphometry. For examination of serum and urine, morphometry, analysis of apoptotic cells and CFU-f assay, seven *apoE*<sup>-/-</sup> mice (ApoES, *n* = 3; ApoEHf, *n* = 4) and nine WT mice (WTS, *n* = 5; WTHf, *n* = 4) were used. For serum LDL and lipid peroxides and extraction of mRNA, six *apoE*<sup>-/-</sup> mice (ApoES, *n* = 3; ApoEHf, *n* = 3) and six WT mice (WTS, *n* = 4; WTHf, *n* = 2) were used. The experimental protocol was approved by the Ethics Review Committee for Animal Experimentation of the University of Occupational and Environmental Health.

### *Examination of serum and urine*

Serum levels of LDL, high-density lipoprotein (HDL), total cholesterol, osteocalcin, and calcium were determined by routine laboratory methods (Panapharm Laboratories, Kumamoto, Japan). Lipid peroxidize production in the serum was determined by measuring thiobarbituric acid-reactive substances (TBARS). The amount of TBARS was calculated based on the standard curve using malondialdehyde (MDA, 0-30 nmol) and expressed as nmoles of MDA per milliliter. Urinary deoxyypyridinoline (D-pyr) levels were determined with an enzyme immunoassay kit (Pyrilinks-D; Metra Biosystems, Mountain View, CA, USA), and the results are expressed as nM/mM of creatinine.

### *Structural analysis using $\mu$ CT*

The trabecular bones of the distal metaphysis and the cortical bones of the mid-diaphysis of the right femurs were analyzed by the  $\mu$ CT system ( $\mu$ CT40; Scano Medical, Bassersdorf, Switzerland), as reported previously.<sup>(21)</sup> On 3D analysis of the distal and middle femurs, the tissue volume (TV, mm<sup>3</sup>) and the trabecular and the cortical bone volume (BV, mm<sup>3</sup>) were measured directly, and the trabecular and the cortical bone volume fraction (BV/TV, %) were calculated.

### *Bone histomorphometry*

Undecalcified sections were obtained from the site of the mid-diaphysis and the distal metaphysis of the right femurs. The specimens were dehydrated and embedded in methylmethacrylate (MMA) after toluidine blue staining to yield 5- $\mu$ m-thick sections. The sections were further stained with alkaline phosphatase (ALP)/TRACP-double staining (Kureha Special Laboratory Co., Tokyo, Japan). For each section, the area of the secondary spongiosa was measured. To exclude the primary spongiosa, the regions within 500  $\mu$ m of the growth plate and one cortical shell-width of the endocortical surface were not measured. For the structural parameters of the distal femur, the trabecular BV/TV (%) and trabecular bone surface (BS,  $\mu$ m) were obtained. For

bone formation parameters, the bone formation rate (BFR/BS, %- $\mu\text{m}/\text{d}$ ) was obtained.<sup>(22,23)</sup>

Paraffinized 5- $\mu\text{m}$ -thick section replicates of the right proximal tibias were stained for TRACP, and trabecular osteoblast surface (Ob.S/BS, %) and osteoclast surface (Oc.S/BS, %) were obtained, as described previously.<sup>(24)</sup>

For each section of the middle femurs, the total cross-sectional area ( $\text{mm}^2$ ), bone marrow area ( $\text{mm}^2$ ), and cortical bone area ( $\text{mm}^2$ ) were measured, and cortical thickness ( $\mu\text{m}$ ) was calculated. BFR/BS (%- $\mu\text{m}/\text{d}$ ) and eroded surface (ES/BS, %) were measured in the endocortical envelopes.

#### *Analysis of apoptotic cells in vivo*

The right tibial section replicates were examined for apoptosis by TUNEL staining<sup>(25)</sup> using the APO-BRDU-IHCTM kit (CHEMICON International, Temecula, CA, USA) in accordance with the manufacturer's instructions. For ALP activity, the serial sections were stained using an alkaline phosphatase kit (86-R; Sigma-Aldrich, St Louis, MO, USA) according to the manufacturer's instructions.

At a magnification of  $\times 400$ , ALP<sup>+</sup> and apoptotic cells were counted from five random visual fields regardless of the trabecular and endosteal surface area, except the regions within 300  $\mu\text{m}$  of the growth plate.

#### *Evaluation of bone marrow cells*

*Preparation of bone marrow cells:* Bone marrow from the bilateral humeri and left femurs was flushed with a total of 6 ml of  $\alpha$ -MEM (Nacalai tesque, Kyoto, Japan). To assay ALP<sup>+</sup> colony forming unit-fibroblastic (CFU-f) formation, bone marrow cells were plated at  $1 \times 10^6$  cells/well in 6-well plates in  $\alpha$ -MEM containing 10% FCS (GIBCO, New York, NY, USA), 10 mg/ml of streptomycin and 10,000 U/ml of penicillin mixed solution (Nacalai tesque), 1.25 U/ml of nystatin (Sigma-Aldrich), and 50  $\mu\text{g}/\text{ml}$  of ascorbic acid (Sigma-Aldrich). The cells were cultured at 37°C in a humidified atmosphere of 5% CO<sub>2</sub> in air with a medium, which was changed at 2-day intervals.

*ALP<sup>+</sup> CFU-f:* On day 13 of culture, the CFU-fs were fixed and stained, using an alkaline phosphatase kit (86-R; Sigma-Aldrich). A colony comprising >50 cells was defined as CFU-f. We counted the total and ALP<sup>+</sup> CFU-fs with the culture dishes backlit and calculated the ratio of ALP<sup>+</sup> CFU-fs to total CFU-fs (ALP<sup>+</sup> CFU-f/total CFU-f).<sup>(26,27)</sup>

#### *RNA isolation from adherent cells and first-strand cDNA synthesis*

The remaining bone marrow cells for each of three samples per group (only the WThf group had two samples) were cultured in the  $\alpha$ -MEM for extraction of total RNA. On day 3 of culture, the nonadherent cells were discarded, and the adherent cells were scraped with 1 ml of TRIZOL Reagent (Invitrogen, Carlsbad, CA, USA) and collected. The RNA was isolated from the adherent cells in accordance with the manufacturer's instructions and cleaned up using DNase I (Roche Applied Science, Penzberg, Germany). First-strand cDNA was reverse-transcribed from total RNA (1  $\mu\text{g}$ ) using Moloney murine leukemia

virus reverse transcriptase (SuperScript; Invitrogen) and Oligo(dT)12–18 Primer (Invitrogen).

#### *Quantitative RT-PCR*

Quantitative RT-PCR analysis was performed using an iCycler apparatus (Bio-Rad Laboratories, Hercules, CA, USA) associated with the iCycler Optical System Interface software (version 3.1; Bio-Rad). The quantitative RT-PCRs for *type I collagen (colla1)*, *osteocalcin*, *p53*, *bax*, *bcl-2*, and  *$\beta$ -actin* were performed in 20  $\mu\text{l}$  with  $\sim 7.5$  ng cDNA, 0.5  $\mu\text{M}$  of primers, and 10  $\mu\text{l}$  iQ SYBR Green Supermix (Bio-Rad). PCR products of mouse *colla1* (170 bp), *osteocalcin* (231 bp), *p53* (232 bp), *bax* (173 bp), *bcl-2* (183 bp), and  *$\beta$ -actin* (639 bp) were amplified using primers for *colla1* (5'-acgtcc tggtagtggtc-3' and 5'-cagggaagcctcttctcct-3'), *osteocalcin* (5'-tgacaaagccttcatgtcca-3' and 5'-tgccagagttt-ggctttagg-3'), *p53* (5'-agagaccgacctacagaaga-3' and 5'-ctgtagcatggcat ccttt-3'), *bax* (5'-tgacagagatgattgctgac-3' and 5'-gatcag ctcgggacttttag-3'), *bcl-2* (5'-ctggcatctt-ctcttccag-3' and 5'-ga cggtagcgcagagagaag-3'), and  *$\beta$ -actin* (5'-ttgagacctcaacac cccag-3' and 5'-acttgcgctcaggaggagca-3'). These primers were designed using Primer3 software (Whitehead Institute for Biomedical Research, Cambridge, MA, USA) and synthesized at the Sigma-Aldrich Japan K.K. Genosys Division (Hokkaido, Japan).  *$\beta$ -actin* was used as an internal control. The amplification conditions were an initial 3 min at 95°C, 45 cycles of denaturation at 95°C for 30 s, annealing at 62°C for 30 s, and extension at 72°C for 30 s. Only  *$\beta$ -actin* was annealed at 65°C for 30 s. The mRNA expression levels were normalized with  *$\beta$ -actin* mRNA expression and expressed as relative value (fold change) to the expression of the WTS group, respectively.

#### *Analysis of apoptotic cells in vitro*

Preparation of osteoblastic cells from 19.5-day fetal calvariae of the respective genotypes, as described previously,<sup>(28)</sup> was performed. The cells were seeded at  $2 \times 10^4$  cells/well (48-well plates) and cultured for 24 h, and subsequently, either human LDL (Athens Research & Technology, Athens, GA, USA) or human oxidized LDL (Biomedical Technologies, Stoughton, MA, USA) was added to the fresh medium at different concentrations (0, 10, 30, 60, and 100  $\mu\text{g}/\text{ml}$ ). TUNEL assay was performed using the In Situ Cell Death Detection Kit, Fluorescein (Roche Applied Science) according to the manufacturer's instructions, at the 24-h time-point, and the cells were counted at 100-fold magnification. The results are expressed as the ratio of apoptotic cell numbers to osteoblast numbers per microscopic field.

#### *Western blot analysis*

The osteoblastic cells were seeded at  $2 \times 10^5$  cells in 6-cm dishes and cultured for 24 h, and subsequently, either 100  $\mu\text{g}/\text{ml}$  human LDL or oxidized LDL was added to the fresh medium. The cells were lysed in lysis buffer and harvested at 0- and 24-h time-points. Aliquots equivalent to  $2 \times 10^4$  cells were subjected to SDS-PAGE on 10% polyacrylamide

TABLE 1. BODY WEIGHT, SERUM LIPID LEVELS, AND BONE CHEMICAL MARKERS IN WT AND apoE<sup>-/-</sup> MICE FED WITH A STANDARD OR A HIGH-FAT DIET

	WT		ApoE <sup>-/-</sup>	
	Standard diet (WTS)	High-fat diet (WTHf)	Standard diet (ApoES)	High-fat diet (ApoEHf)
Body weight (g)	25.2 ± 2.2	27.6 ± 2.8	23.5 ± 3.7	24.0 ± 2.1
Serum				
Total cholesterol (mg/dl)	102 ± 5.8	312 ± 34	1195 ± 155*	2443 ± 407 <sup>†‡</sup>
LDL (mg/dl)	7.2 ± 2.8	84.7 ± 16	258 ± 29*	727 ± 124 <sup>†‡</sup>
Lipid peroxides (nmol MDA/ml)	83.6 ± 27	56.4 ± 0.8	1210 ± 125*	1649 ± 173 <sup>†‡</sup>
HDL (mg/dl)	47.6 ± 3.9	51.5 ± 5.1	10.0 ± 0.0*	6.7 ± 2.9 <sup>†</sup>
Osteocalcin (ng/ml)	121.4 ± 32	72.8 ± 19	111 ± 11	65.7 ± 31
Calcium (mg/dl)	8.4 ± 0.5	8.2 ± 0.5	7.8 ± 0.2	8.2 ± 0.8
Urine				
D-pyr (nM/mM Cre)	14.4 ± 3.0	18.7 ± 2.9	11.2 ± 0.3	38.9 ± 2.6 <sup>†‡</sup>

Results are expressed as mean ± SD over all animals in each diet group. These data are 17-wk-old mice.

\*  $p < 0.05$  vs. WT on standard diet.

<sup>†</sup>  $p < 0.05$  vs. WT on high-fat diet.

<sup>‡</sup>  $p < 0.05$  vs. apoE<sup>-/-</sup> on standard diet.

gel and transferred to a PVDF transfer membrane (Hybond-P; GS Healthcare Bio-Science Corp., Piscataway, NJ, USA). The membrane was blocked with 5% nonfat milk in PBS containing 0.1% Tween 20 for 1 h at room temperature. p53 antibody (Sigma-Aldrich) was used. The membrane was incubated overnight at 4°C with the primary antibody diluted to 1:200. After washing, the membrane was incubated with the appropriate second antibody conjugated to horseradish peroxidase. Protein was detected by the ECL Plus Western Blotting Detection System (GS Healthcare Bio-Science Corp.).

### Statistical analysis

All values are expressed as mean ± SD. They were assessed by the Tukey-Kramer posthoc test after one-way ANOVA. In the study of apoptotic cell analysis in vitro, statistical analysis was done by Student's *t*-test and Dunnett's method on the WT control group and Dunnett's method on the apoE<sup>-/-</sup> control group. Correlation studies were done using Pearson's correlation coefficient. Two-factor factorial ANOVA was used to detect the potential effects of apoE<sup>-/-</sup> and a high-fat diet. Statistically significant differences were set at <0.05. Calculation was performed with Stat View 5.0 software (SAS Institute, Cary, NC, USA) on a Macintosh computer.

## RESULTS

### Body weight, serum lipid levels, and bone chemical markers

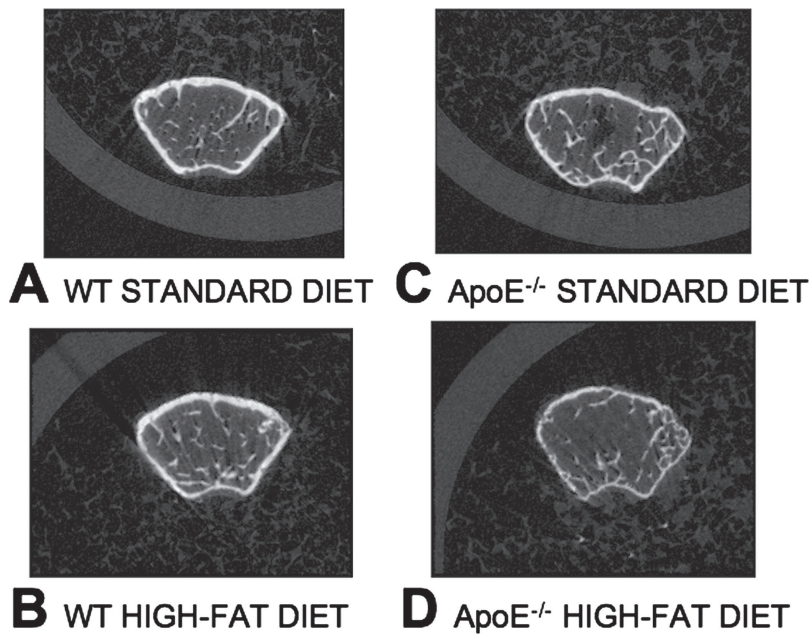
No significant differences in body weight were found among the four groups at the end of the experiment, at 17 weeks of age after being given either a standard or high-fat diet for the duration of 12 weeks (Table 1). In the apoE<sup>-/-</sup> mice fed with the standard diet (ApoES), the values of total cholesterol, LDL and lipid peroxides significantly increased, compared with those in the WT mice fed with

the standard diet (WTS). The values further significantly increased in the apoE<sup>-/-</sup> mice fed with a high-fat diet (ApoEHf), compared with those in the WT mice fed with a high-fat diet (WTHf) and the ApoES group. The levels of serum HDL in the ApoES and ApoEHf groups significantly decreased compared with those in the WTS and WTHf groups, respectively. Serum osteocalcin and calcium levels did not significantly differ among the four groups of mice. The values of urinary D-pyr significantly increased in the ApoEHf group compared with those in the WTHf and ApoES groups (Table 1).

### Morphological and morphometrical assessments of bone mass and turnover in the femurs

Trabecular BV/TV in the distal femur and cortical BV/TV in the midfemur were measured by  $\mu$ CT (Fig. 1). The values of trabecular BV/TV did not significantly differ among the groups. However, the values of cortical BV/TV in the ApoEHf group were significantly smaller than those in the WTHf group (Table 2).

In the samples of mid- and distal femur, histological and histomorphometrical assessments under light microscopy were also performed on undecalcified sections of the specimens after fluorescence labeling. The values of cortical thickness of the mid-femur in the ApoEHf group were significantly smaller compared with the respective values in the WTHf and ApoES groups. The BFR/BS values in the trabecular bone surface in distal femoral metaphysis were reduced significantly in the ApoEHf group compared with the WTHf group (Table 2). In the samples of the proximal tibias, the trabecular Ob.S/BS values decreased, and the trabecular Oc.S/BS values increased in the WTHf and ApoEHf groups insignificantly (Table 2). The BFR/BS values in endosteum at the mid-femur were also reduced in the ApoEHf group. The ES/BS values in the ApoEHf group



**FIG. 1.** 2D axial image of the middle femur from 17-wk-old mice using  $\mu$ CT in each diet group (A–D). Cortical thickness of the apoE<sup>-/-</sup> mice (C and D) decreased compared with the WT mice (A and B).

TABLE 2. MORPHOLOGICAL AND MORPHOMETRICAL ASSESSMENTS OF BONE MASS AND TURNOVER IN THE FEMURS

	WT		ApoE <sup>-/-</sup>	
	Standard diet (WTS)	High-fat diet (WTHf)	Standard diet (ApoES)	High-fat diet (ApoEHf)
<b>Bone mass</b>				
Distal femur trabecular BV/TV (%)	5.94 ± 0.7	4.15 ± 0.9	6.21 ± 1.0	3.92 ± 1.6
Mid-femur cortical BV/TV (%)	29.8 ± 1.7	30.9 ± 1.9	25.9 ± 1.6	24.2 ± 2.2*
Cortical thickness (μm)	137.3 ± 7.5	136.3 ± 4.1	133.5 ± 5.6	112.5 ± 6.5 <sup>†</sup>
<b>Bone turnover</b>				
<b>Trabecular bone</b>				
BFR/BS (%-μm/d)	41.6 ± 7.1	40.5 ± 2.3	32.2 ± 4.2	20.1 ± 5.5*
Ob.S/BS (%)	21.2 ± 4.2	12.7 ± 3.7	21.9 ± 11.6	8.8 ± 2.6
Oc.S/BS (%)	10.7 ± 5.1	14.9 ± 4.7	8.1 ± 1.2	15.1 ± 2.7
<b>Cortical endosteum</b>				
BFR/BS (%-μm/d)	59.5 ± 17.6	49.2 ± 6.9	32.8 ± 11.4	4.3 ± 7.5*
ES/BS (%)	3.28 ± 1.4	7.49 ± 4.5	4.75 ± 2.2	12.3 ± 1.9 <sup>†</sup>

Results are expressed as mean ± SD over all animals in each diet group. These data are 17-wk-old mice.

\*  $p < 0.05$  vs. WT on high-fat diet.

<sup>†</sup>  $p < 0.05$  vs. apoE<sup>-/-</sup> on standard diet.

significantly increased compared with those in the ApoES group (Table 2).

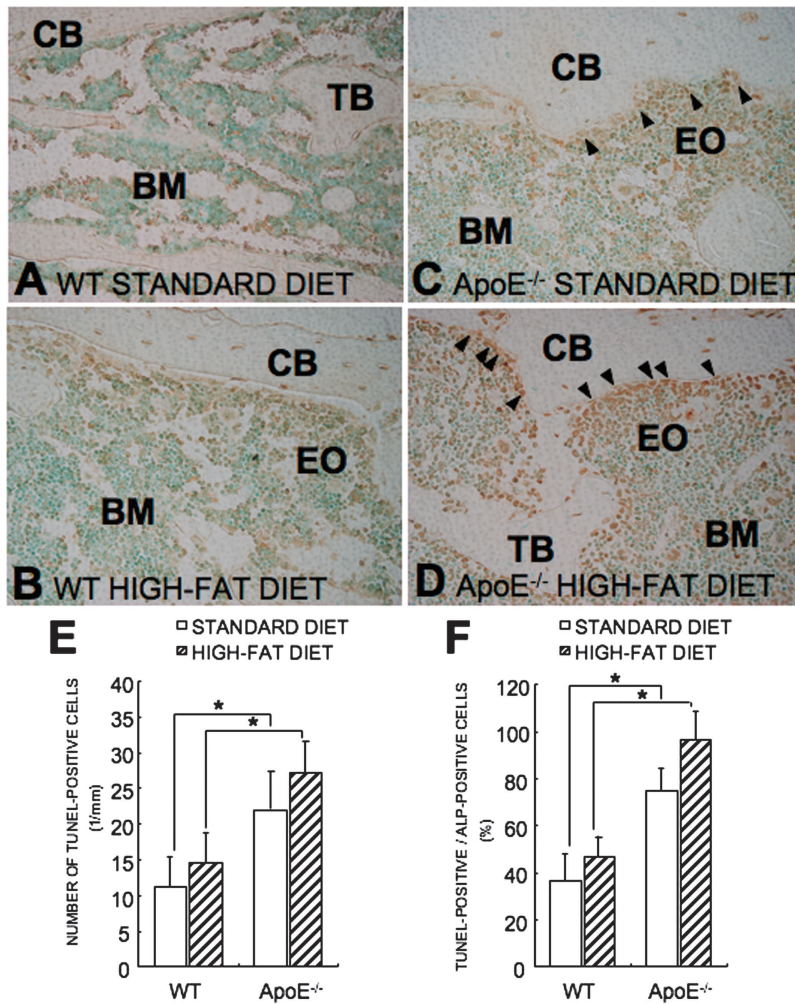
#### Alkaline phosphatase stain and TUNEL stain of bone and bone marrow

As reductions in bone formation and bone mass by a high-fat diet and apoE<sup>-/-</sup> were confirmed, we proceeded to evaluate the osteogenic cells in the bones and bone marrow of the mice. The number of ALP<sup>+</sup> cells in the sections of decalcified specimens of the proximal tibia did not significantly differ among the four groups of mice (data not shown). However, the number of TUNEL<sup>+</sup> cells in the bone and bone marrow significantly increased in the ApoES and ApoEHf groups compared with the WTS and WTHf groups, respectively (Figs. 2A–2E). The percent ra-

tios of the number of TUNEL<sup>+</sup> cells to ALP<sup>+</sup> cells also significantly increased in the groups of apoE<sup>-/-</sup> mice compared with those in the groups of WT mice (Fig. 2F).

#### Assessments of bone marrow cells for total CFU-f and ALP<sup>+</sup> CFU-f

To evaluate the bone marrow capacity for developing osteogenic cells, we assessed the numbers of total CFU-f and ALP<sup>+</sup> CFU-f from bone marrow cells of the femur. The number of total CFU-f in the WTHf and ApoEHf groups was apparently smaller than the respective values of those in the WTS and ApoES groups, but the differences were not statistically significant (Fig. 3A). The ratios of ALP<sup>+</sup> CFU-f to total CFU-f in the ApoEHf group were significantly smaller than those of the ApoES group (Fig. 3B).



**FIG. 2.** TUNEL<sup>+</sup> cells in the proximal tibiae (magnification,  $\times 400$ ). TUNEL stain showed an increase of apoptotic cells (arrowheads) at the endosteum in the apoE<sup>-/-</sup> mice (C and D) compared with the WT mice (A and B). BM, bone marrow; CB, cortical bone; EO, endosteum; Tb, trabecular bone. Cell quantification using serial sections in the proximal tibiae. (E) The number of TUNEL<sup>+</sup> cells significantly increased in the ApoES and ApoEHf groups compared with the WTS and WTHf groups, respectively. (F) The ratio of TUNEL<sup>+</sup> cells to the ALP<sup>+</sup> cells also significantly increased in the ApoES and ApoEHf groups compared with the WTS and WTHf groups, respectively (\* $p < 0.05$ ).

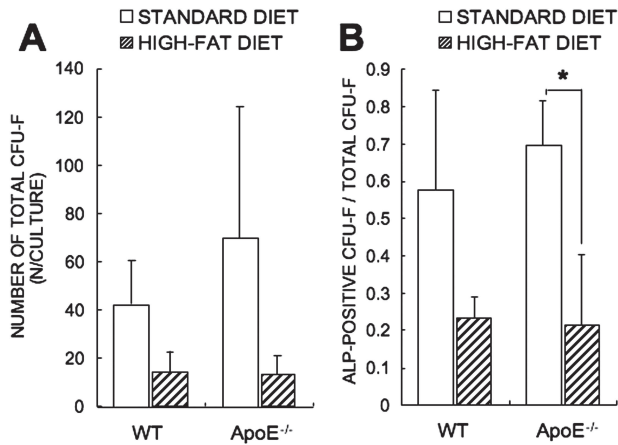
#### Assessments of mRNA expression of osteocalcin, type I collagen, p53, bax, and bcl-2 genes in bone marrow cells

Increases of apoptotic cells in ALP<sup>+</sup> cells in histology and reductions of ALP<sup>+</sup> CFU-f developments in bone marrow cells prompted us to study the mRNA expressions of genes relating to the development of osteogenic cells and apoptosis in bone marrow cells of the femur. We measured the mRNA expression of *osteocalcin*, *colla1*, *p53*, *bax*, and *bcl-2* genes. The expression of *osteocalcin* mRNA did not significantly differ among the groups (data not shown). The expression of *colla1* mRNA did not significantly differ either (Fig. 4A). The expression of *p53* mRNA in the ApoES and ApoEHf groups significantly increased compared with those in the WTS and WTHf groups, respectively (Fig. 4B). The expression of *bax* mRNA (data not shown) and the ratio of *bax* to *bcl-2* mRNA expression (Fig. 4C) had upward trends in the ApoES and ApoEHf groups, and the expression of *bcl-2* mRNA had a downward trend in the ApoES and ApoEHf groups (data not shown), but the differences were not statistically significant.

#### Statistical evaluations for the effect of a high-fat diet and apoE<sup>-/-</sup> on the development and apoptosis of osteogenic cells in bone marrow cells

The contributions of a high-fat diet and apoE<sup>-/-</sup> to the development and apoptosis of osteogenic cells in bone marrow cells were evaluated by two-factor factorial ANOVA. It showed that a high-fat diet significantly affected all the parameters of bone marrow CFU-fs, but apoE<sup>-/-</sup> did not significantly relate to these parameters of bone marrow cells (Table 3). Both a high-fat diet and apoE<sup>-/-</sup> significantly affected the ratios of the numbers of TUNEL<sup>+</sup> cells to ALP<sup>+</sup> cells. The expression of *osteocalcin* mRNA was not affected by either a high-fat diet or apoE<sup>-/-</sup>. The expression of *colla1* mRNA, however, was significantly affected by apoE<sup>-/-</sup>. The expression of *p53* mRNA was significantly affected by both of these factors (Table 3).

Between the expression of *p53* mRNA in bone marrow cells and serum levels of LDL and lipid peroxides, significant linear correlations were found in apoE<sup>-/-</sup> mice, combined with the ApoES and ApoEHf groups (Figs. 5A and 5B). However, these correlations were not significant in the groups of WT mice, WTS and WTHf (data not shown). We



**FIG. 3.** The results of CFU-f assay using bone marrow cells. (A) The number of total CFU-f in the WTHf and ApoEHf groups was apparently smaller than the respective values of those in the WTS and ApoES group, but the differences were not statistically significant. (B) The ratio of ALP<sup>+</sup> CFU-f to total CFU-f significantly decreased in the ApoEHf group compared with the ApoES group (\* $p < 0.05$ ).

calculated the ratio values of mRNA expression of *osteocalcin* to *colla1* (mRNA *osteocalcin*/mRNA *colla1*) as the parameter of the signals representing the capacity of developing osteogenic cells in bone marrow cells. A significant inverse relationship was found between the values of *p53* mRNA expression and the ratio values of mRNA *osteocalcin*/mRNA *colla1* (Fig. 5C).

#### TUNEL fluorescein assay and Western blot analysis in cultures of osteoblastic cells obtained from fetal mice calvaria by the addition of LDL or oxidized LDL

To compare the effects of LDL and oxidized LDL on the development of apoptosis between WT and apoE<sup>-/-</sup> osteoblastic cells, we obtained fetal calvarial cells. In the cells from WT fetal mice, TUNEL fluorescein assays revealed a minimum increase in apoptotic cells in the culture media with ascending doses of LDL or oxidized LDL (Figs. 6A–6C). In the cells from apoE<sup>-/-</sup> fetal mice, however, the number of apoptotic cells was significantly larger than that of WT cells in the medium without LDL or oxidized LDL (Figs. 6D–6F) and significantly increased with the addition of 100  $\mu$ g/ml LDL or lipid or oxidized LDL (Fig. 6G). Two-factor factorial ANOVA showed that the ratios of apoptotic cells in the culture media to which was added LDL or oxidized LDL were significantly affected by the respective lipid concentrations and apoE<sup>-/-</sup> as well (Table 4).

To examine the protein level of p53 in osteoblastic cells, we performed Western blot analysis. The p53 expression in the apoE<sup>-/-</sup> cells exposed to 100  $\mu$ g/ml LDL or oxidized LDL at 0 and 24 h increased compared with that in the WT cells (Fig. 4D).

## DISCUSSION

Our study showed that a high-fat diet and apoE<sup>-/-</sup> synergistically increased apoptosis in osteoblastic cells, leading

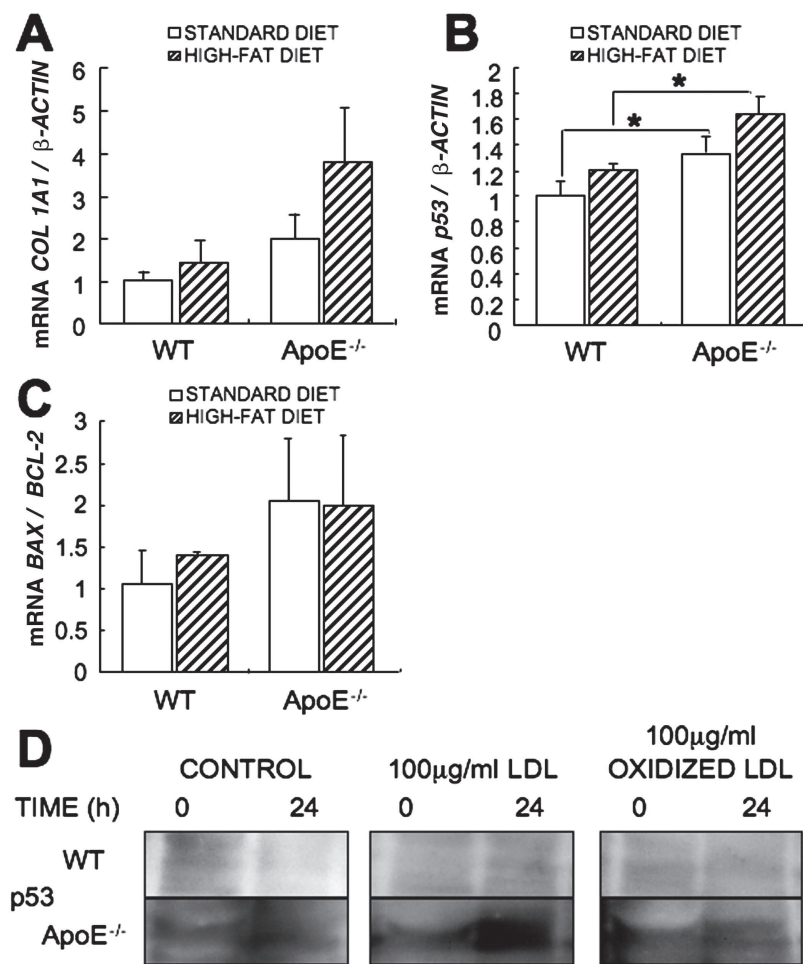
to decreased trabecular and endosteal bone formation in mice. *p53* mRNA expression significantly increased in the two groups of apoE<sup>-/-</sup> mice compared with those in the respective groups of WT mice. There were significant linear correlations between the expression of *p53* mRNA in adherent stromal cells and serum levels of LDL and lipid peroxides. In the apoE<sup>-/-</sup> mice, the ratio of the numbers of ALP<sup>+</sup> CFU-f to total CFU-f significantly decreased in the group of a high-fat diet compared with that in the group of standard diet. Thus, it was revealed that, in the apoE<sup>-/-</sup> mice, the osteoblastic cells were vulnerable to apoptosis induced by either LDL or oxidized LDL.

Several previous studies have reported that bone volume and bone strength decrease in animals given a high-fat diet.<sup>(29–31)</sup> A high-fat diet reduces intestinal absorption of calcium and causes a negative balance between the intake and the output of serum calcium.<sup>(32,33)</sup> There is no difficulty in thinking that the deficient state of serum calcium is compensated for by osteoclastic bone resorption. Thus, we considered that a high-fat diet increases bone resorption consequent to a negative balance of calcium.

The number of ALP<sup>+</sup> cells in vivo did not significantly differ among the four groups of mice. However, the percent ratios of the number of TUNEL<sup>+</sup> cells to ALP<sup>+</sup> cells in vivo increased in the WTHf and ApoEHf groups compared with the WTS and ApoES groups, respectively (Fig. 2F). These in vivo data on histomorphometry indicate that the disappearance of ALP<sup>+</sup> cells increased in the WTHf and ApoEHf groups. Thus, we think that the apparent reductions in ALP<sup>+</sup> CFU-f assays in vitro (Fig. 3B) are consistent with the histomorphometry data in vivo in these groups. Also, we found that the BFR/BS and Ob.S/BS values decreased in the ApoEHf group (Table 2). These data are compatible with the reduction in life span of functional osteoblasts in vivo. All these data in vitro and in vivo consistently indicate that the apoptotic disappearance of osteoblastic lineage cells was enhanced in the condition of the deleted *apoE* gene under high-fat loading.

A previous study showed that the degree of hyperlipidemia correlated with the degree of osteoclastic functional activity in the marrow.<sup>(34)</sup> In our study, the apoE<sup>-/-</sup> mice fed with a high-fat diet revealed excessive hyperlipidemia; as a result, it increased bone resorption, excessively, compared with apoE<sup>-/-</sup> mice fed with the standard diet. On the other hand, another previous study suggested that increased dietary lipids interfered with osteoblast maturation and reduced bone mineralization in vivo,<sup>(5)</sup> and our results of trabecular BV/TV and serum osteocalcin are compatible with this previous study. Consequently, in the apoE<sup>-/-</sup> mice fed with a high-fat diet, bone resorption increased, and bone formation decreased remarkably, compared with the apoE<sup>-/-</sup> mice fed with the standard diet. Thus, we considered that the abnormal bone phenotype is only observed in apoE<sup>-/-</sup> mice fed with a high-fat diet, because of the uncoupling of bone resorption and formation.

*p53* mRNA expression increased in adherent stromal cells by either a high-fat diet or apoE<sup>-/-</sup>. The ratio of the *osteocalcin* mRNA to the *colla1* mRNA negatively correlated with *p53* mRNA expression. TUNEL stain revealed that apoE<sup>-/-</sup> and a high-fat diet increased ALP<sup>+</sup> apoptotic



**FIG. 4.** Evaluation of mRNA expression in bone marrow adherent cells. (A) The expression of *colla1* mRNA had an upward trend in the ApoEHf group, but the difference did not achieve statistical significance. (B) The expression of *p53* mRNA in the ApoES and ApoEHf groups significantly increased compared with those in the WTS and WTHf groups, respectively ( $*p < 0.05$ ). (C) The ratio of *bax* to *bcl-2* mRNA expression had upward trends in the ApoES and ApoEHf groups. (D) Western blot analysis for p53 in cultures of osteoblastic cells with the addition of 100  $\mu$ g/ml LDL or oxidized LDL. The p53 expression in the apoE<sup>-/-</sup> cells exposed to 100  $\mu$ g/ml LDL or oxidized LDL at 0 and 24 h increased compared with that in the WT cells.

**TABLE 3.** STATISTICAL EVALUATIONS FOR THE EFFECT OF A HIGH-FAT DIET AND APOE<sup>-/-</sup> ON THE DEVELOPMENT AND APOPTOSIS OF OSTEOGENIC CELLS IN BONE MARROW CELLS

	ApoE <sup>-/-</sup>	High-fat diet
Bone marrow cells		
Total CFU-f	0.427	0.008*
ALP <sup>+</sup> CFU-f	0.345	0.015*
ALP <sup>+</sup> /total	0.588	0.003*
Proximal tibia		
TUNEL <sup>+</sup> /ALP <sup>+</sup>	<0.0001*	0.013*
Bone marrow adherent cells		
<i>osteocalcin</i> mRNA	0.156	0.968
<i>colla1</i> mRNA	0.01*	0.059
<i>p53</i> mRNA	0.001*	0.01*

Data represent *p* values for the relationship. The significant differences were determined by two-factor factorial ANOVA and set at  $p < 0.05$ .

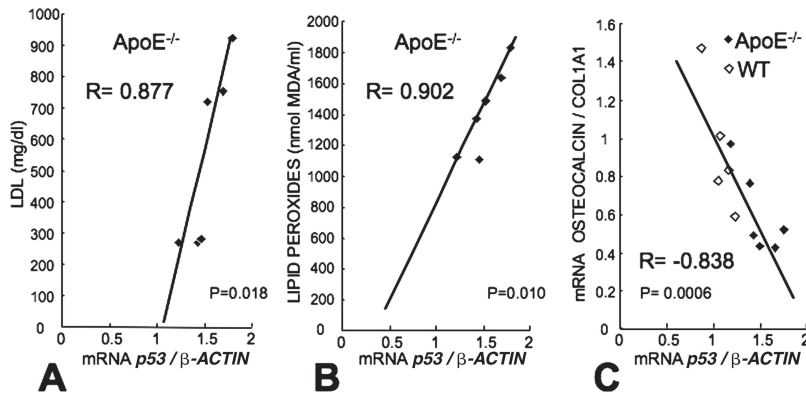
\* Significant.

cells. Apoptosis in osteoblasts and osteoblast precursors was caused by a high-fat diet and further enhanced by apoE<sup>-/-</sup>. It was previously reported that LDL or oxidized LDL caused apoptosis in endothelial cells,<sup>(35,36)</sup> vascular smooth muscle cells (VSMCs),<sup>(37,38)</sup> and osteoblastic Saos2 cells<sup>(39)</sup> and that apoE<sup>-/-</sup> also induced apoptosis of VSMCs<sup>(40)</sup> and neurons.<sup>(41)</sup> Although *p53* mRNA expres-

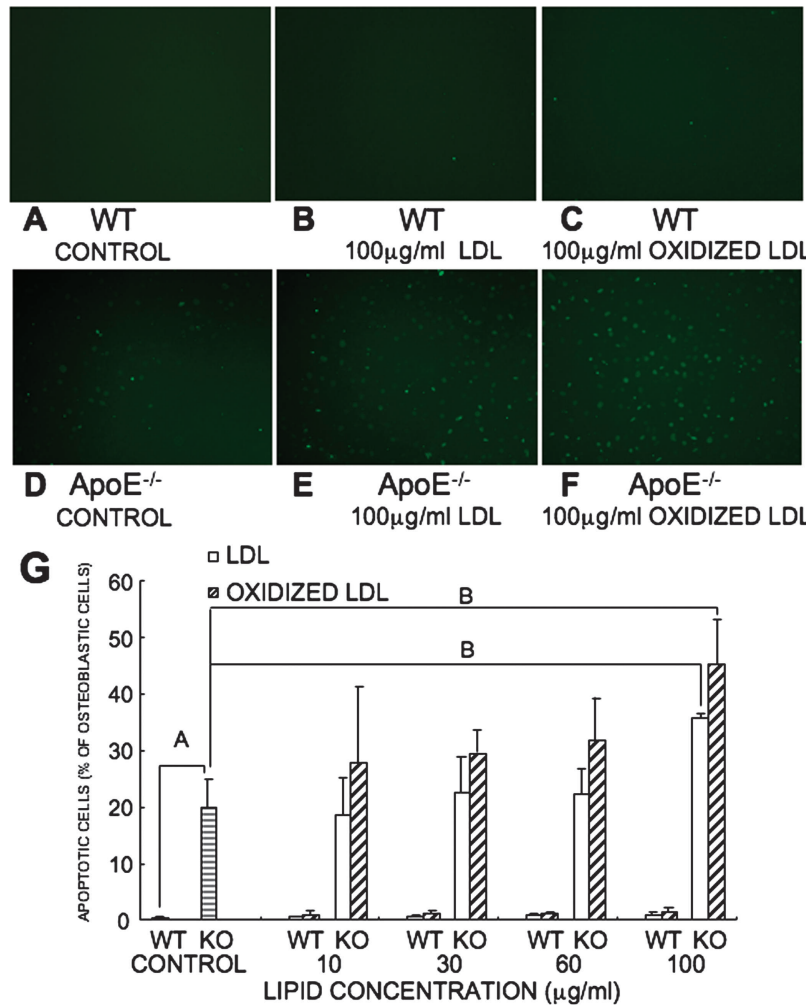
sion in the adherent stromal cells did not statistically correlate with serum LDL or lipid peroxides in the WT mice, they significantly correlated with it in the apoE<sup>-/-</sup> mice. Thus, the *apoE* gene in osteoblastic cells could prevent apoptosis induced by LDL or lipid peroxides. Those synergistic apoptotic actions of apoE<sup>-/-</sup> and LDL or lipid peroxides were obvious in the calvarial osteoblastic cell study. A high-fat diet and/or apoE<sup>-/-</sup> caused apoptosis in osteoblastic cells, and this could be a major reason for decreased bone formation in trabecular bone and endosteum.

In our study, apoE<sup>-/-</sup> mice fed with the standard diet could maintain constant levels of serum osteocalcin, the numbers of ALP<sup>+</sup> cells in vivo, and the numbers of ALP<sup>+</sup> colonies in vitro despite the increases of osteoblastic cell apoptosis. We measured the Ob.S/BS values, finding that the parameter apparently reduced under high-fat loading in both the WT and apoE<sup>-/-</sup> mice. These data suggest that the reduction in the number of bone-forming cells, mediated by apoptosis, was induced by high-fat loading, and the effects were enhanced in the *apoE* gene deleted condition. Regarding the serum osteocalcin levels, we also found that the ALP<sup>+</sup> cells in vivo and signal for bone-forming cell lineages such as mRNA *colla1* in vitro did not decrease in the face of high-fat loading (Fig. 4A). Thus, we can assume that the recruitment of bone-forming cells was not reduced in either





**FIG. 5.** Correlation between the expression of *p53* mRNA and lipid levels and the ratio values of mRNA expression of *osteocalcin* to *coll1a1*. The expression of *p53* mRNA in the apoE<sup>-/-</sup> mice had a significantly strong correlation with serum LDL levels (A) and a much stronger correlation with serum lipid peroxide levels (B), but not in the WT mice (data not shown). (C) Correlation between the mRNA *osteocalcin* / mRNA *coll1a1* ratio and the *p53* mRNA expression in bone marrow adherent cells had a significantly strong negative correlation. *r* = Pearson's correlation coefficient.



**FIG. 6.** Effects of the addition of LDL and oxidized LDL in cultures of osteoblastic cells from fetal mice calvaria using TUNEL fluorescein assay. Representative fluorescence micrographs of osteoblastic cells (A–F; magnification, ×100). In the cells from WT mice, there was a slight increase in apoptotic cell number by the addition of LDL and oxidized LDL (A–C). In the cells from the apoE<sup>-/-</sup> mice, in contrast, there were apoptotic cells in the apoE<sup>-/-</sup> control (D), and the number of apoptotic cells increased with addition of both 100 μg/ml LDL (E) and 100 μg/ml oxidized LDL (F). Quantification of the absolute number of apoptotic cells expressed as a percentage of the total number of osteoblastic cells. The percentage of apoptotic cells in the apoE<sup>-/-</sup> control significantly increased compared with that in the WT control. Addition of 100 μg/ml LDL and 100 μg/ml oxidized LDL in the apoE<sup>-/-</sup> significantly increased the percentage of apoptotic cells compared with the apoE<sup>-/-</sup> control. <sup>A</sup>*p* < 0.05 compared with the WT control. <sup>B</sup>*p* < 0.05 compared with the apoE<sup>-/-</sup> control.

group in this experiment, maintaining serum osteocalcin levels in the face of an apparent reduction in bone formation surface in vivo.

Apoptosis of osteoblastic cells has been reported as a possible mechanism of bone loss in a few kinds of disorders, such as rheumatoid arthritis,<sup>(42)</sup> glucocorticoid-induced osteoporosis,<sup>(43)</sup> and unloading-induced osteoporosis.<sup>(26,44)</sup> The mechanism of osteoblast apoptosis involves the modu-

lation of mitochondrial functions, intracellular reactive oxygen species, and Bcl-2 protein.<sup>(45)</sup> Recently, it has been reported that high plasma homocysteine, which has been associated with an increased risk of fracture, enhances apoptosis in human bone marrow stromal cells through the intracellular reactive oxygen species-mediated mitochondrial pathway and NF-κB activation.<sup>(46)</sup> Our report is the first to describe that osteoblast apoptosis induced by lipid

TABLE 4. STATISTICAL EVALUATIONS FOR THE EFFECT OF LIPID CONCENTRATIONS OR apoE<sup>-/-</sup> IN CULTURES OF OSTEOBLASTIC CELLS FROM FETAL MICE CALVARIA

Ratio of apoptotic cells	ApoE <sup>-/-</sup>	Lipid concentration
Addition of LDL	<0.0001*	0.003*
Addition of oxidized LDL	<0.0001*	0.016*

Data represent *p* values for the relationship. The significant differences were determined by two-factor factorial ANOVA and set at *p* < 0.05.

\* Significant.

loading through p53 signal is related to the reduced bone formation in apoE<sup>-/-</sup> mice.

This study suggests that the *apoE* gene prevented a high-fat diet from inducing apoptosis in osteoblasts and osteoblast precursors. In conclusion, *apoE* gene deficiency enhances the reduction of bone formation induced by a high-fat diet through the stimulation of p53-mediated apoptosis in osteoblastic cells.

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