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Osteoclast formation, survival and morphology are highly dependent on exogenous cholesterol/ lipoproteins

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Abstract

Osteoporosis is associated with both atherosclerosis and vascular calcification. No mechanism yet explains the parallel progression of these diseases. Here, we demonstrate that osteoclasts (OCL) depend on lipoproteins to modulate cellular cholesterol levels and that this controls OCL formation and survival. Removal of cholesterol in OCL via high-density lipoprotein or cyclodextrin treatment dose-dependently induced apoptosis, with actin disruption, nuclear condensation and caspase-3 activation. One mechanism linked to the induction of OCL apoptosis was the cell-type-specific failure to induce HMG-CoA reductase mRNA expression, suggesting an absence of feedback regulation of de novo cholesterol biosynthesis. Furthermore, cyclodextrin treatment substantially suppressed essential M-CSF and RANKL-induced survival signaling pathways via Akt, mTOR and S6K. Consistent with these findings, cholesterol delivery via lowdensity lipoprotein (LDL) significantly increased OCL viability. Interestingly, OCLs from the LDL receptor (LDLR)-/- mouse exhibited reduced size and lifespan in vitro. Remarkably, LDLR + I + OCL in lipoprotein-deficient medium phenocopied LDLR-/- OCL, while fusion and spreading of LDLR-/- OCL was rescued when cholesterol was chemically delivered during differentiation. With hyperlipidemia being associated with disease of the vascular system and bone, these findings provide novel insights into the selective lipoprotein and cholesterol dependency of the bone resorbing cell.

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Abbreviations: OCL, osteoclast; HDL, high-density lipoprotein; LDL, low-density lipoprotein; LDLR, low-density lipoprotein receptor; LPDS, lipoprotein-deficient serum; MBCD, methyl- β cyclodextrin; S6, ribosomal protein S6; S6K, ribosomal protein S6 kinase; mTOR, mammalian target of rapamycin

Introduction

Epidemiological evidence links osteoporosis and cardiovascular disease. Patients with low bone mineral density have a greater risk of mortality in later life, especially from stroke¹ and cardiovascular disease,^{2,3} a higher plaque score of the artery wall,^{4–7} and higher total cholesterol levels.^{8,9} Although a role for estrogen is suggested, the specific mechanisms connecting osteoporosis and cardiovascular disease remain unknown. Both high low-density lipoprotein (LDL)- and low high-density lipoprotein (HDL)-cholesterol levels were recently found to correlate with low bone mass in postmenopausal women,¹⁰ suggesting a possible link between plasma lipids and the maintenance of bone mass.

The paradox of osteoporosis accompanied by atherosclerotic calcification has been the subject of several experimental studies. The role of LDL oxidation products and their accumulation in the vessel wall during atherosclerotic lesion formation is well established.^{11,12} Others have shown effects of oxidized lipids on osteoblastic differentiation *in vitro*. These lipids direct progenitor marrow stromal cells to favor adipogenic over osteogenic differentiation.^{13,14} It was further speculated that oxidized lipids might inhibit osteoblastic activity and bone mineral formation *in vivo*.¹⁵ Indeed, a mouse strain with a susceptibility to atherosclerosis (C57BL6) did lose greater bone mass in response to a high fat diet when compared to atherosclerosis-resistant strains.¹⁶

In contrast to profound inhibitory effects on osteoblast differentiation,¹³ oxidized LDL was found to have no negative effect on the formation of osteoclasts (OCLs) *in vitro*.¹⁷ OCLs are multinucleated bone resorbing cells formed from hematopoietic precursors via membrane fusion. Absence of native LDL suppresses OCL formation, and this was attributed to a requirement for cholesterol during the fusion process.¹⁷ This could be due in part to the very low rate of *de novo* sterol synthesis in the OCL,¹⁸ thus necessitating exogenous sources of cholesterol.

In the present study, we examined the role of exogenous cholesterol availability on OCL formation and survival in vitro. Unlike mesenchyme-derived cells (i.e. osteoblasts), OCLs do not respond to rapid cholesterol removal by inducing hydroxymethylglutaryl coenzyme-A reductase (HMGCR), suggesting the lack of this feedback regulation in the cholesterol biosynthesis pathway. Moreover, cholesterol removal by HDL, apolipoprotein A1 (ApoA1) or methyl- β -cyclodextrin (MBCD) triggers induction of apoptosis and OCL death, whereas similar treatments of osteoblasts were without lasting effect. Consistently, as a result of MBCD-induced cholesterol efflux, we found suppression of antiapoptotic signaling pathways involving Akt and mammalian target of rapamycin (mTOR)/ribosomal protein S6 (S6) kinase (S6K) in OCLs. On the other hand, LDL-mediated cholesterol delivery significantly enhances OCL survival. Furthermore, OCLs derived from the LDL receptor (LDLR)-/- mice exhibit abnormal morphology and shortened lifespan *versus* OCL from LDLR+/+ mice. These findings suggest that OCLs depend on lipoproteins to control their cellular cholesterol levels and, in turn, their lifespan.

Results

Cholesterol efflux from OCLs

For promoting cholesterol efflux, we used the pharmacological agent MBCD and the physiological cholesterol acceptors HDL and ApoAl. The ability of these agents to stimulate cholesterol efflux from labeled OCLs was validated by the transfer of ³H-cholesterol released from the OCL into the medium after incubation with acceptors, as described in Materials and Methods. We found efficient ³H-labeled cholesterol transfer from OCLs to MBCD (20 mM) and HDL (400 μ g/ml) at rates comparable to those for mesenchymal cells (osteoblastic and fibroblastic cells, data not shown). MBCD promoted rapid cholesterol efflux with 80% transferred from OCL (versus 3% in controls) within 1 h, (P<0.001). Transfer to MBCD at 15 mM elicited a similar response. Cholesterol transfer to HDL and ApoAI was much slower, at 70 and 36% versus 12% in controls (P<0.001) in 24 h, respectively. These profound differences in the kinetics of cholesterol efflux have been reported also in other cell systems such as in fibroblasts, in which MBCD was shown to accept cholesterol at up to 70-fold greater rates (V_{max}) than HDL.19

Cholesterol removal via MBCD induces OCL apoptosis

The very low rate of *de novo* sterol synthesis in OCLs¹⁸ suggests the possibility that these terminally differentiated cells could depend on external sources to control cellular cholesterol levels. This would be consistent with the epidemiological link between lipoproteins and osteoporosis. To test this hypothesis, we examined the effects of rapid cholesterol removal, via MBCD, on OCL life span. OCLs were treated with MBCD (5-20 mM) or left untreated for up to 2 h, after which the media were replaced with media containing lipoproteindeficient FBS (LPDS) to limit cholesterol uptake from the medium. Interestingly, MBCD treatment caused extensive dose- and time-dependent induction of OCL apoptosis, which we assessed microscopically in the large, multinucleated OCL population (Figure 1). At 5h after initial treatment, OCL contained numerous vacuoles (Figure 2b). Classical apoptotic signs were later observed between 6 and 12h of treatment. This included membrane blebbing (Figure 2e), nuclear condensation (Figure 1a,b and 2e,e',f,f'), caspase-3 activation (Figure 1e), actin cytoskeleton disruption (Figure 1c,d and 2d) and formation of apoptotic bodies (Figure 2f). We and others have previously found that nuclear condensation/ pyknotic body formation correlates with positive TUNEL staining in this model system^{20,21}. The apoptotic response seemed remarkable since mesenchymal and HepG2 cells did not undergo apoptosis in response to similar MBCD treatment (data not shown). Similar 2 h treatment was previously shown



Figure 1 Removal of cholesterol with MBCD induces OCL apoptosis. OCLs were treated with MBCD at several doses and for different times, as noted below. After treatment, cells were further cultivated in medium containing 10% LPDS to monitor survival. Time- and dose-dependent studies showed maximal induction of apoptosis with equivalent effects at 15 or 20 mM, respectively, after a 9–12 h post-treatment period. Data were collected within these ranges based on morphological examination by microscopy. (**a**, **c**) TRAP-positive OCLs were examined for (**a**) nuclear and (**c**) actin cytoskeleton integrity 12 h after 90 min MBCD treatments at indicated concentrations. (**b**, **d**) OCLs were treated with 20 mM MBCD for indicated times, after which cultures continued 9 h in the absence of MBCD. TRAP-positive OCLs were assessed for (**b**) nuclear and (**d**) actin cytoskeletal integrity. (**e**) Dose-dependent effects of MBCD on caspase 3 activity immediately following 2 h treatments. The bars represent mean \pm S.E.**P*<0.05, ****P*<0.001 (Co versus treatment)

to be nontoxic to fibroblasts, although continuous 8 h exposure could induce some cell death.¹⁹ Although there was no evidence of cell death in MB1.8, NIH/3T3 or HepG2 cells, initial and transient morphological rounding was observed (data not shown). This response and the actin disruption seen in OCLs suggest that the cytoskeleton was commonly affected following cholesterol removal. Indeed, disruption of the ring-like f-actin pattern in OCLs (up to 67% greater than in controls, Figure 1c, d) was more prominent than the induction of nuclear condensation (greater by up to 31%, see Figure 1a, b) when quantified 12 h after MBCD treatment.

The earliest response to MBCD measured in the OCL was caspase activation, which was evident at 60 min (data not shown) and maximally increased after 120 min (Figure 1e) of continuous treatment. Morphological apoptosis was not apparent at either initial time point. A similar magnitude of caspase activity was observed 9h after MBCD treatment



Figure 2 Morphological features of OCL apoptosis after cholesterol removal. Purified OCLs were untreated (**a**, **b**) or treated with 20 mM MBCD (1 h followed by 6 h (**ce**, **e**') or 9 h (**f**, **f**') without treatment). OCLs were stained for actin cytoskeletal (**b**, **d**) and nuclear (**e**', **f**') integrity, as described in Materials and Methods. Morphology (**a**, **c**) and actin staining (**b**, **d**) of control (**a**, **b**) *versus* MBCD-treated OCLs (**c**, **d**). (**e**) Phase-contrast and (**e**') nuclear integrity of OCL in an early stage of apoptosis. Arrowheads indicate surface blebs containing pyknotic nuclei. (**f**) TRAP staining and (**f**') nuclear integrity of OCL in a later stage of apoptosis. TRAP-positive apoptotic bodies are indicated with arrowheads and contain pyknotic nuclei. Scale bars indicate 340 μm (**a**-**d**) and 100 μm (**e**, **e**', **f**, **f**')

(data not shown) at a time point when morphological apoptosis was apparent. However, when FBS was substituted for LPDS in the media used to maintain the OCLs after MBCD treatment, apoptosis was suppressed by one-third at this later time point, as assessed by quantifying multinucleated OCLs with pyknotic nuclei (data not shown). This suggested that although the apoptotic program was immediately activated by MBCD, the response was asynchronous within the OCL population and one-third of the cells that had not already committed to the apoptotic program were rescued by the subsequent restoration of cholesterol levels via the lipoproteins found in normal FBS.

By comparison to osteoblasts, fibroblasts and hepatocytes, OCLs showed high sensitivity to rapid cholesterol removal. Interestingly, other cells of the macrophage/monocyte lineage (J774 and Raw264.7 cells) shared this sensitivity and responded to MBCD treatment by undergoing apoptosis, albeit at a somewhat later time point (data not shown). Whereas OCL apoptosis was seen at 5–10 h, both J774 and Raw264.7 cells appeared normal during this time. However by 24 h, extensive apoptosis was observed. Thus several cell types from the macrophage/monocyte lineage did undergo apoptosis in response to rapid cholesterol removal, the most profound response being seen in OCLs.

HDL induces OCL apoptosis, while LDL extends survival

To further test the role of external cholesterol on OCL lifespan, we examined the physiologically relevant effects of HDL and LDL particles. HDL-mediated removal of OCL cholesterol

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Figure 3 Removal of cholesterol induces OCL apoptosis, while addition increases survival. (**a**, **b**) Dose-dependent effects of (**a**) HDL (48 h) or (**b**) LDL (72 h) on nuclear integrity. After treatments, OCLs were stained for both TRAP and DNA as in Materials and Methods. OCLs with pyknotic nuclei were quantified and are expressed as a percent of the total population. Dose-dependent effects of (**a**) HDL and (**b**) LDL (n=3 for both analyses). The bars represent means \pm S.E. ***P<0.001 (Co versus treatment)

induced dose-dependent apoptosis featuring the same classical apoptotic signs as observed with MBCD treatment (Figure 2), by up to five-fold (Figure 3a). As expected, the timeframe was much longer than that for MBCD (48 *versus* 1 h), consistent with the slower kinetics of cholesterol removal by HDL. In addition, note that some subclasses of HDL are not full acceptors but are also capable of donating cholesterol. Similarly, cholesterol removal using ApoAI particles lead to a significant, dose-dependent induction of OCL apoptosis by up to 40% (data not shown). To test if HDL, like MBCD, could induce apoptosis in other cells, we also examined its effects on macrophages, osteoblasts, fibroblasts and hepatocytes. However, unlike the OCL, all cell types remained viable after extended (72 h) HDL treatment (data not shown). Thus among the cell types tested, HDL-induced apoptosis was limited to

OCLs, which seem to exhibit the highest sensitivity to cholesterol withdrawal.

Apoptotic death in response to cholesterol removal by HDL suggested that lipoproteins can control OCL lifespan and that the cholesterol donor LDL might extend the lifespan of OCL *in vitro*. In general, purified OCLs maintained in the presence of M-CSF and RANKL remain mostly intact over a period of 48 h followed by a steep decline in their survival rate with substantial spontaneous apoptosis seen at 72 h. Consistent with our hypothesis, LDL exposure of OCLs for 72 h significantly reduced the number of apoptotic OCLs by 50% (Figure 3b) as measured by quantifying OCL with pyknotic nuclei. Thus, HDL and LDL have the potential to modulate OCL lifespan: the cholesterol donor, LDL, extends lifespan.

Cholesterol removal impairs cytokine survival signaling to Akt, mTOR and S6 kinase

It was recently shown that cholesterol removal via MBCD could negatively impact RANKL signaling to Akt, although it had no effect on signaling to NF-kB.²² We very recently found that RANKL and M-CSF signaling also activate the mTOR and S6 kinase (S6K) pathway and that this is critical for OCL survival.23 Whereas RANKL signaling to Akt elicits only a small stimulation, signaling to the mTOR/S6K pathway is robust. We therefore examined the impact of MBCD treatment on cytokine-induced activation of several targets in this pathway. As expected, M-CSF profoundly stimulated phosphorylation of Akt (Thr 308) ranging from 7- to 38-fold at 30 min, depending on the experiment (Figure 4a). However, MBCD pretreatment of OCLs for 45 min reduced signaling to Akt by MCSF and RANKL to 50%. Meanwhile, RANKL stimulation was minimal, as we have previously reported.²³ In most replicates, RANKL did not stimulate Akt, and there was no response to pretreatment with MBCD (Figure 4a). Akt signaling participates in regulation of S6K activity via TSC2 and mTOR, which regulate activation via phosphorylation of Thr 389 in S6K.²⁵ We next measured the phosphorylation of S6K (Thr 389), and its activity on the downstream target, S6 (Figure 4b). M-CSF stimulated phosphorylation of S6K and S6 by 4.2- and 7.6-fold, respectively. As we have previously reported, RANKL, which did not substantially activate Akt, consistently induced S6K and S6 phosphorylation by two- and 4.2-fold, respectively. Interestingly, pretreatment with MBCD strongly suppressed the stimulatory effects of both M-CSF and RANKL on S6K phosphorylation up to 88%, as measured at 30 min, and reduced S6 phosphorylation by about 60%. In order to examine effects of MBCD treatment on mTOR more closely, we also assessed the phosphorylation of 4E-BP1, a translational regulator that can act as a direct substrate of mTOR. Whereas M-CSF and RANKL stimulated phosphorylation of 4E-BP1 (Ser 65) in control OCLs by up to 9.8- and 3.5-fold (Figure 4b), respectively, pretreatment with MBCD reduced these effects by an average of 70%. In certain replicates, MBCD pretreatment completely abrogated RANKL-induced 4E-BP1 phosphorylation. Together, these data demonstrate that MBCD-mediated removal of cholesterol has a profound impact on survival cytokine-induced



Figure 4 Cholesterol removal impairs cytokine signaling to mTOR and S6 kinase. (a) Purified OCL were pretreated with or without methyl- β -cyclodextrin (MBCD, 15 mM) in medium containing 0.5% FBS for 45 min, and stimulated with M-CSF (100 ng/ml) or sRANKL (200 ng/ml) for 30 min, as indicated. Cell lysates were immunoblotted with phosphospecific antibodies for Akt, S6 and 4E-BP1 and S6K1. Blots were stripped and reprobed with control antibodies for Akt and actin. Relative band intensities (*versus* control) are indicated beneath each lane, and are normalized to Akt (a) or actin (b) expression levels. One representative experiment out of three repeats with similar results is shown

signaling to the Akt and especially the mTOR/S6K pathway. Negative effects on these kinases and their respective substrates are consistent with the capacity of MBCD to induce apoptosis in the OCL.

Cholesterol efflux upregulates hydroxymethylglutaryl coenzyme A reductase (HMGCR) in mesenchymal and liver cells, but not OCLs

Induction of OCL apoptosis by MBCD suggested the possibility that the normal responses to cholesterol removal (i.e. induction of cholesterol synthesis machinery) might be absent in these cells. To test the degree to which the cholesterol biosynthetic machinery was induced in OCLs by cholesterol removal, we examined feedback induction of HMGCR, a critical enzyme of this pathway. HMGCR mRNA expression is ordinarily induced when cellular cholesterol levels decline, via sterol regulatory elements within its promoter.^{26,27} MBCD, combined with lovastatin, has previously been shown to increase HMGCR mRNA by threefold.²⁸ We have previously observed that OCLs synthesize very little sterol¹⁸ and undergo apoptosis in response to lovastatin treatment.²⁰ Lovastatin was therefore omitted in our experiments to avoid confounding effects of apoptosis. Cells were treated with MBCD for 1 h, shown to remove 80% of

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cellular cholesterol, and then maintained in LPDS for a period of 4 h, which is a time point well before profound apoptosis in OCL can be observed. In osteoblastic, fibroblastic and hepatic cells, HMGCR mRNA expression was, as expected, induced 2–4-fold after treatment with MBCD (Figure 5a). Meanwhile, although OCLs could rapidly and extensively transfer cholesterol to MBCD, no significant HMGCR mRNA induction (1.1fold) was observed (Figure 5a). Furthermore in the uninduced state (i.e. no MBCD treatment), basal levels of HMGCR mRNA expression were comparatively low. This and the lack of response to MBCD treatment indicate absence of feedback regulation of cholesterol synthesis in OCLs, but not in other cell types tested.

We also examined caveolin expression in OCLs in comparison to other cells. Caveolin interacts with cholesterol in caveolae membranes and appears to play a major function in transporting cholesterol between caveolae and the endoplasmic reticulum and Golgi apparatus. The caveolin promoter contains two sterol regulatory elements, which control expression in response to changes in cholesterol levels.²⁹ Interestingly, in OCLs as well as J774 and Raw 264.7 cells, all derived from the macrophage/monocyte lineage, caveolin-1 protein was expressed at extremely low levels (Figure 5b). This was in contrast to mesenchymal MB1.8 osteoblasts and NIH3T3 fibroblasts, where a robust caveolin-1 signal was detected. Caveolin-2 was abundantly expressed in the J774 macrophages, and a faint signal was detected in RAW 264.7 macrophages and in OCLs (Figure 5b). However, it is notable that caveolin-2 lacks the three cysteine palmitoylation motifs found in caveolin-1 that are essential for cholesterol binding and for proper transport of caveolin to the plasma membrane.³⁰ Absence of caveolin expression in HepG2 cells (Figure 5b) has been previously described.³¹ For the OCL this suggests that, in addition to a lack of substantial



Figure 5 Cholesterol efflux induces feedback regulation of cholesterol synthesis in mesenchymal cells, but not in OCL. Cells have been treated with/ without 15 mM MBCD for 1 h, thereafter maintained in LPDS for 4 h followed by RNA extraction. (a) Induction of HMGCR mRNA in untreated and MBCD-treated NIH/3T3, MB1.8, HepG2 cells and purified OCLs, measured by Northern analysis. (b) Expression of caveolin-1 and -2 protein in untreated HepG2, NIH/ 3T3, MB1.8, Raw 264.7, J774 and OCLs was assessed by Western blot analysis

cholesterol synthesis and feedback regulation of HMGCR, these cells may also have limited ability to transport cholesterol intracellularly through caveolin-1. The absence of these internal sterol regulatory mechanisms might explain the apoptotic response of OCLs to cholesterol withdrawal.

OCL formation in the LDL receptor -/- mouse: impaired spreading/fusion and survival

To extend beyond pharmacological observations described above, we used a genetic approach involving the LDL receptor -/- mouse (hereafter -/-) to examine OCL formation and survival. It is notable that both +/+ and -/-OCLs were generated using mouse MB1.8 osteoblast-like cells, derived from a +/+ background, to drive differentiation in response to 1,25(OH)₂ vitamin D₃, as described in Materials and Methods. OCL formation using the bone marrow of -/versus C57BL6/J background mice (hereafter +/+) revealed that OCLs from -/- mice were significantly smaller (Figure 6a, b). This was assessed in OCL (-/-)/osteoblast (+/+)cocultures by staining for the OCL marker enzyme, tartrateresistant acid phosphatase (TRAP). In -/- males, a 5.8-fold reduction in the proportion of large TRAP-positive OCLs $(>340 \,\mu\text{m}$ in diameter) was observed, while in females there was a 2.3-fold reduction on day 9 of culture (Figure 6c). There was no enhancement of OCL size between days 7 and 11 of differentiation (data not shown), suggesting that formation of OCL (>340 μ m) was suppressed rather than delayed. As expected, the reduction in large OCLs in -/- cultures was accompanied by a significant increase in the number of smaller multinucleated TRAP-positive OCLs (<340 µm, containing at least five nuclei, arrows in Figure 6b) and a general shift towards smaller sized OCLs (Figure 6d). OCLs from -/cultures mostly fell into the smallest grouping ($< 10\,000\,\mu\text{m}^2$), while within the +/+ cultures, most OCLs ranged from $10\,000-50\,000\,\mu m^2$.

Interestingly, small -/- OCLs contained numerous nuclei and demonstrated irregular morphologies, as visualized by TRAP staining (Figure 6e, g). Phalloidin staining revealed that the f-actin rings in the small -/- OCLs, although often limited to small areas within the cell, frequently spread beyond the limits of TRAP staining (Figure 6f). Nonetheless, the f-actin cytoskeleton was intact and arranged in a ring-like pattern. During the OCL purification process, we noticed that these small, irregular shaped -/- OCLs were less tightly attached to the extracellular matrix. Indeed, while large OCLs were typically resistant to collagenase treatment, most of the small -/- OCLs were detached. We quantified the number of collagenase-sensitive OCLs by transferring them into new culture vessels and counting after reattachment. By this measure, we observed a significant 1.6-fold increase in collagenase-sensitive -/- versus +/+ OCLs (data not shown). Interestingly, after replating, the -/- OCLs adopted the more typical round morphology generally seen in +/+OCLs.

We also quantified the mean area of OCLs from female mice. Consistent with the reduction in the number of large OCLs (Figure 6c), the mean size of -/- OCLs was reduced 2.4-fold *versus* +/+ OCLs (Figure 7f). To mimic the effects of



Figure 6 Reduced spreading and/or fusion of LDLR-/- OCLs. OCL formation with bone marrow cells from background (C57BL6J) or LDLR-/- mice. Morphologies of TRAP-stained (**a**) +/+ and (**b**) -/- OCLs, and (**c**) percent of large (> 340 μ m diameter) male and female OCLs. Arrows in (**b**) indicate some of the small, multinucleate OCLs. (**d**) Size distribution of +/+ and -/- female OCLs. Size groupings are indicated (inset and beneath X-axis). (**e**) TRAP, (**f**) actin and (**g**) nuclear staining of small, multinucleated -/- OCLs. Arrows in (**f**) indicate the 'actin rings', the white lines in (**f**) and (**g**) mark the outlines of TRAP-positive OCLs as seen in panel e. Scale bars represent 340 μ m (**a**,**b**) and 50 μ m (**e**-**g**), error bars in (**c**) represent means ± S.E. ****P* < 0.001 (+/+ *versus* -/-), $\diamond \diamond \diamond P < 0.001$ (-/- male *versus* -/- female)

LDLR deletion, we also examined OCL formation using LPDScontaining medium. The morphology of TRAP-positive +/+OCLs in the presence of LPDS (Figure 7b) resembled that of -/- OCLs in normal FBS (Figure 7c), showing a four-fold reduction in the number of large OCLs (Figure 7e) and a threefold reduction in the mean OCL size (Figure 7f). Interestingly, the use of LPDS during differentiation further reduced the number of large OCLs in -/- cultures by about two-fold versus that seen with FBS (Figure 7e). Indeed, in LPDS there was no difference in the mean area of +/+ and -/- OCLs (Figure 7f). Overall, the findings demonstrate that cholesterol uptake during differentiation via the LDLR has a major influence on OCL morphology, and also suggests that OCLs have other ways than via the LDLR to import extracellular cholesterol. In support with this hypothesis, we incubated the OCLs with a cholesterol-cyclodextrin complex for 30 min daily. Such cholesterol-saturated complexes have been shown to effectively and rapidly enrich cells with cholesterol,³² thereby bypassing the absence of the LDLR. In the

osteoblast-OCL coculture system, pulsatile treatment with cholesterol–cyclodextrin complexes significantly increased the formation of large -/- OCLs ($>340\,\mu\text{m}$) up to four-fold (Figure 7c-d, g). This treatment also had a positive effect on OCL formation in +/+ cultures, although the effect was not always statistically significant.

In separate analyses, we examined the survival of -/- OCL after purification. For these analyses, both -/- and +/+ OCL were maintained for 1 and 2 days in FBS supplemented with the differentiation/survival cytokines M-CSF and RANKL. In three of four experiments, the rate of spontaneous apoptosis was significantly increased by greater than 60% for -/- OCLs versus +/+ OCLs. Spontaneous apoptosis was observed after both 24 and 52 h of culture (Figure 8a–c) and did not require cholesterol removal by MBCD or HDL. These data are consistent with the observation that LDL extends OCL lifespan and that the inability of cholesterol uptake via the LDLR leads to induction of apoptosis, albeit milder than that seen with pharmacological cholesterol removal (Figure 1).

Discussion

Osteoclasts are terminally differentiated, multinucleated cells formed by the fusion of mononuclear progenitors of the monocyte/macrophage family. OCL activation results in the degradation of the calcified collagenous bone matrix. The delicate balance between bone resorption by OCLs and deposition of new bone by osteoblasts determines bone mass in the adult undergoing consistent adaptive bone remodeling. Disparity of these coupled activities in favor of bone resorption results in the loss of bone and deterioration of its trabecular architecture leading to osteoporosis. Interestingly, osteoporosis often coexists with atherosclerotic calcification in patients, associated with formation of bone mineral in vascular walls and simultaneous calcium loss from bone.1,2,6,7 Growing evidence suggests that abnormalities in lipid metabolism could be a common factor underlying both diseases, 10,16,33–35 and that plasma lipids might play a role in maintaining bone mass.^{13,36-38} A better understanding of the coupling of osteoporosis and atherosclerosis is of critical importance as indicated by some studies whereby treatment of one disease may have beneficial effects on the other. 39-44

In the present study, we investigated the effects of exogenous cholesterol on OCL formation and survival. We show here for the first time that lipoproteins influence survival of purified OCL, and that the addition of cholesterol via native LDL could double the lifespan of purified OCL *in vitro*. More importantly, failure to gather cholesterol seems to have a negative impact on OCL survival, since purified LDLR-/-OCLs showed significantly higher levels of apoptosis than +/+OCLs. Conversely cholesterol deficiency, caused by the use of LPDS in the coculture system, significantly suppressed the formation of TRAP-positive, multinucleated OCLs. This is in concordance with earlier reports suggesting a role for LDL, isoPGE2 and oxidized LDL in OCL fusion and differentiation via effects on bone marrow hematopoietic precursors.^{17,45}

Furthermore, we find that cholesterol withdrawal has detrimental effects on OCL survival. The correlation between



Figure 7 Effects of cholesterol withdrawal and repletion on OCL formation. (**a**–**d**) TRAP staining of OCLs from (**a**,**b**) +/+ and (**c**,**d**) -/- mice. +/+ OCLs were formed in the presence of (**a**) FBS or (**b**) LPDS, while -/- OCLs were formed in the presence of (**c**) FBS or (**d**) FBS with daily pulsatile cholesterol treatments (MBCD-cholesterol inclusion complexes), as described in Materials and Methods. (**e**) Percent of large ($>340 \mu$ m diameter) and (**f**) mean area of +/+ and -/- OCLs differentiated in FBS *versus* LPDS-containing medium. (**g**) Change in large -/- OCLs when formed in the absence (FBS) or presence (FBS + chol) of MBCD-cholesterol inclusion complexes. Scale bar represents 340 μ m (**a**–**d**); error bars in (**e**–**f**) represent means \pm S.E. ***P < 0.001 (+/+ versus -/-), $\diamond \diamond P$ < 0.001, $\diamond \diamond P$ < 0.01 (-/- FBS versus -/- LPDS); (**g**) ***P < 0.001 (-/- FBS versus -/- FBS plus MBCD-cholesterol inclusions)



Figure 8 Reduced survival of LDLR -/- OCLs. OCLs derived from +/+ and -/- mice were purified and maintained for 1–2 days in the presence of M-CSF (15 ng/ml) and sRANKL (25 ng/ml). Nonapoptotic and apoptotic OCLs were quantified as TRAP-positive, intact cells or a ring of apoptotic bodies (arrows), respectively, as described in Figure 3. (a) -/- and (b) +/+ OCLs 52 h after purification. (c) Percentage of apoptotic OCLs 24 and 52 h after purification. Scale bar represents 340 μ m (a, b), error bars (c) represent means \pm S.E. **P < 0.01 (+/+ versus -/-)

cholesterol efflux and OCL apoptosis is reflected by the differences in kinetics of cholesterol removal, which substantially vary for the pharmacological agent MBCD, and the physiological cholesterol acceptors HDL and ApoAI. These

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differences in cholesterol efflux (MBCD > > HDL > ApoAI) are consistent with reports in other cell systems.¹⁹ Although the rate of cholesterol release to phospholipid-containing acceptors like HDL can be cell type dependent,⁴⁶ we observed comparable cholesterol efflux rates to HDL in OCLs as well as MB1.8 osteoblasts and NIH3T3 fibroblasts over 24 h (five-, versus seven-fold, respectively). As a consequence of promoting cholesterol efflux in OCLs, apoptosis was induced up to five-fold, as verified by cell morphology, disruption of the actin cytoskeleton, nuclear condensation and caspase-3 activation. Two cell lines of the macrophage/monocyte lineage (J774, Raw264.7) did also show an apoptotic response to MBCD, but interestingly, among the cell types tested, HDLinduced apoptosis was strictly limited to OCLs. Thus, in our hands the OCL was outstanding in its susceptibility to apoptosis due to cholesterol removal.

It is notable that we have found no previous study showing that native HDL could induce cell death in any cell type. On the contrary, several reports show that HDL protects endothelial cells by blocking the mitochondrial pathway of apoptosis, thereby preventing caspase-9 and -3 activation, and apoptotic alterations of the plasma membrane.^{47–49} However, glucose

modification and/or oxidation of HDL resulted in deterioration of the mitochondrial function and apoptosis of endothelial cells.⁵⁰ In the present study, no such prior modification of HDL was necessary to induce OCL apoptosis.

To further investigate the role of cholesterol uptake in OCL formation and survival, we used the LDLR knockout mouse (-/-) as a model.⁵¹ These mice are extensively used to investigate lipoprotein metabolism and atherosclerosis. Preliminarily data indicate that the LDLR-/- mice exhibit a bone phenotype with 8% lower BMD when placed on a high fat diet.⁵² Based on earlier work,^{16,36} it was hypothesized that the observed changes in BMD were due to accumulation of oxidized lipids in bone leading to an inhibition of osteoblastic differentiation. However, additional evidence suggests that lipid oxidation products and hyperlipidemia might also enhance osteoclastic potential and function,45 thus making the OCL likely to contribute to the lower BMD. We found that multinucleated -/- OCLs were significantly smaller and more susceptible to spontaneous induction of apoptosis when compared to +/+ OCLs. There was a significant 2.3-fold and 5.8-fold difference in the average OCL area between -/and +/+ female and male mice, respectively. These apparent gender-based differences were consistently observed in independent experiments. However, the reduction of OCL area in -/- was accompanied by a significant increase in the total number of multinucleated OCL (data not shown). Specifically, there were more multinucleated OCLs, and a greater proportion of these multinucleated OCLs were of a smaller measured size. This indicates that spreading and, perhaps to a lesser extent, fusion of these cells might be impaired. In addition, OCL formation and size in both LDLR + / + and - / - cultures were equally suppressed when LPDS was used during differentiation, providing further evidence that reduced receptor-mediated cholesterol uptake is a primary driver for the phenotype observed. Interestingly, by limiting exogenous sources of cholesterol (LPDS versus FBS), formation of large OCL (>340 um diameter) was further diminished two-fold in LDLR-/- mice. This suggests that, yet insufficient, OCL have means of cholesterol uptake other than via LDLR.

More importantly, we provide proof for the role of exogenous cholesterol during the OCL formation process by using cholesterol–cyclodextrin inclusion complexes. β -Cyclodextrin form stable inclusion complexes with cholesterol by incorporating it into their hydrophobic cavity, ^{53–55} and (receptor independently) enrich cells with cholesterol–cyclodextrin inclusion complexes significantly enhanced the formation of OCLs (>340 um) by 2–4-fold. This observation verified that the morphological phenotype of LDLR –/– OCL was driven by the loss of functional LDLR and could be rescued by providing cholesterol in a receptor-independent mechanism.

Our experiments do raise the question of why in particular the OCL is so sensitive to cholesterol efflux in comparison to the other cells that we examined. We tested the expression and regulation of HMGCR as a major, rate-limiting enzyme in the cholesterol biosynthetic pathway. Treatments that stimulate cholesterol efflux have been shown to increase the expression of this enzyme at the level of mRNA and protein in various cell types.^{28,59–62} Surprisingly, we found that HMGCR mRNA level was not responsive to MBCD treatment in OCL,

although 2-4-fold induction was seen in cell types resistant to MBCD-induced apoptosis (MB1.8, NIH/3T3 and HepG2 cells). Moreover, baseline HMGCR mRNA expression was found to be low in OCLs. Indeed, we previously found a very low level of cholesterol synthesis in OCL as <0.1% of radiolabeled mevalonate is incorporated into nonsaponifiable lipids (sterols).¹⁸ Based on these two observations and the fact that HMGCR is only one of several cholesterol biosynthetic genes regulated via sterol response elements in their promoters, this suggests that more than one enzyme of this biosynthetic pathway is downregulated. Even more so, we describe here extremely low, if any, expression of caveolins (caveolin 1 and -2) in the OCLs. Absence of caveolin-1 in OCLs has been reported very recently in another study.²¹ Caveolins, which form a family of membrane-anchored proteins, have been implied to function in cellular cholesterol homeostasis, particularly intracellular transport of cholesterol.⁶³ Altogether our data point towards an impaired feedback regulation of cholesterol synthesis and - transport in these terminally differentiated cells.

OCL survival primarily depends not only on antiapoptotic signaling mediated by M-CSF but also other cytokines like RANKL and TNF α .^{64,65} A very recent study demonstrated that MBCD treatment reduced survival and activity of in vitro generated OCL.²¹ This correlated with impairment of RANKLinduced signaling. In the present study, we observed a partial reduction of M-CSF signaling to antiapoptotic Akt, as well as more substantial inhibition of signaling farther downstream of M-CSF and RANKL to the mTOR/S6K pathway, by cholesterol withdrawal. Signaling to this pathway was greatly reduced (up to 80%), especially as related to mTOR activation of S6K and phosphorylation of 4E-BP1. We have previously shown that the mTOR inhibitor, rapamycin, induces OCL apoptosis and suppresses OCL formation.²² Thus, the suppressive effects of cholesterol withdrawal on Akt/mTOR/S6K signaling might provide a contributing mechanism by which OCL survival and formation are reduced.

In summary, our data suggest that OCL formation and survival is highly dependent on exogenous sources of cholesterol. Addition of native LDL cholesterol increases OCL viability by suppressing spontaneous apoptosis, while cholesterol removal strongly induces apoptosis in OCL and, to a lesser degree, macrophages. This was accompanied by a profound inhibition on the cytokine-induced signaling to Akt/mTOR/S6K survival pathways. We further suggest that the susceptibility of OCLs to cholesterol removal might be due to impaired feedback regulation of cholesterol synthesis involving HMGCR. OCL from the hyperlipidemic LDLR-/- mouse showed reduced size and lifespan in vitro, indicating a major role for cholesterol delivery via the LDLR in OCL formation and survival. The present findings suggest a regulatory role of lipoproteins in OCL biology, which might play a role in the epidemiological link between osteoporosis and atherosclerosis.

Materials and Methods

Cell culture

Osteoclast-like cells (OCL) were generated in six-well plates by coculturing mouse bone marrow cells with mouse MB1. 8 cells⁶⁶ in differentiation

medium (MEM- α supplemented with 10% FBS and 10 nM 1,25(OH)₂-Vitamin D₃) in a humidified atmosphere with 10% CO₂. MB1.8 cells were removed by collagenase treatment,⁶⁷ and purified OCL were further cultured in the presence of 15 ng/ml M-CSF (R&D, Minneapolis, MN, USA) and 25 ng/ml soluble RANK ligand (sRANKL) (R&D) in lipoprotein-deficient medium containing 92% less total cholesterol than FBS (MEM- α supplemented with 10% LPDS (Intracel, MD, USA), and 10 nM 1,25(OH)₂-Vitamin D₃. For signal transduction experiments, purified OCL were cultured in 0.5% FBS without cytokines with/without treatment for 1–2 h. NIH/3T3 fibroblasts, J774 macrophages, Raw 264.7 macrophages and HepG2 hepatocytes were purchased from ATCC.

For modulating cholesterol levels in OCL, the following reagents were used at indicated concentrations: LDL (38–190 μ g/ml; Calbiochem, San Diego, CA, USA), HDL (200–800 μ g/ml; Calbiochem), ApoA1 (20–40 μ g/ml), (MBCD (5–20 mM: Sigma, St. Louis, MO, USA).

Preparation of Steroid–MBCD inclusion complexes

A stock solution of 0.4 mg/ml cholesterol and 10% cyclodextrin was prepared by combining 10 ml of 10% cyclodextrin with 200 μ l of cholesterol (20 mg/ml in 2:1 methanol/chloroform) at 40°C by vortexing. Each solution was freeze-dried for storage, and filtered through a 0.2 μ m filter before use. Complexes were added to the cultures for 30 min at 37°C each day, starting with day 5 when first signs of OCL fusion were apparent and until differentiation was complete. After each treatment, media were aspirated, cultures were washed with PBS and regular differentiation medium was added.

Histochemistry

Osteoclast-like cells were washed with PBS, fixed in 3.7% formaldehyde for 15 min and stained as previously described.⁶⁸ Briefly, cells were stained for TRAP in 50 mM sodium acetate (pH 5.0), 30 mM sodium tartrate, 0.1% Triton X-100 with 0.3 mg/ml fast red violet LB and 0.1 mg/ml naphthol AS-MX phosphate (both from Sigma). Actin and nuclei were stained with 5 U/ml TRITC or FITC-labeled phalloidin together with 50 μ g/ml Hoechst 33342 dye (both from Molecular Probes, Eugene, OR, USA) and mounted using antifade (Molecular Probes).

OCL survival and actin structure

OCL were triple-stained as described above. Total OCL and OCL with pyknotic nuclei and/or a disrupted actin cytoskeleton were counted. Each treatment was performed in triplicate wells and at least 35 representative observation fields were assessed per well under 10 \times magnification. Data were analyzed by analysis of variance (ANOVA).

Caspase-3 activity assay

Fluorescent caspase-3 activity assay kit was from Biovision Inc. (Mountain View, CA, USA). Cells were washed with Hank's buffered salt solution (HBSS), and lysed and scraped in the lysis buffer supplied by the manufacturer. Lysates were cleared by centrifugation and supernatants were assayed according to instructions provided with the kit. In all, 20–40 μ g of total protein was used per assay, and activities were determined after 2 h incubation at 37°C. Values are in relative fluorescence units (RFUs) per μ g protein.

Cholesterol efflux assay

MB1.8, NIH-3T3 and purified OCLs in six-well plates were incubated with growth medium containing 10% LPDS and 1 $\mu\text{Ci/ml}$ of cholesterol (50 Ci/

mmol) (Perkin-Elmer; Boston, MA, USA) for 24 h. Following labeling, cells were washed and incubated for an additional 6 h in fresh growth medium to allow for equilibration of ³H-cholesterol with intracellular cholesterol. Owing to the short lifespan of purified OCLs in culture, the equilibration step was shortened to 6 h for these and all comparator cell lines. In two control experiments using NIH/3T3 fibroblasts, we found that the shorter equilibration period resulted in increased efflux to HDL by 10–25% after 24 h, while efflux to MBCD (1 h) was slightly increased up to 7% (data not shown). Efflux was assessed with three different cholesterol acceptors ApoAI, HDL and MBCD. After the efflux period of 1–24 h, media were collected and cells were dissolved in 0.1 M NaOH. Aliquots of both the supernatants and dissolved cells were subjected to liquid scintillation spectrometry. Cholesterol efflux is expressed as the amount of ³H-cholesterol in the medium as a percentage of the total label in both fractions.

Preparation of protein lysates and Western blotting

After treatments, cells were placed on ice and washed twice with ice-cold β -glycerophosphate-HEPES-buffered solution (β -HBS): 50 mM HEPES (pH 7.6), β -glycerophosphate (50 mM), EGTA (1 mM), NaCl (150 mM), and then lysed in β -HBS containing Triton X-100 (0.2%), microcystin LR (1 μ M), Na₃VO₄ (1 mM), dithiothreitol (1 mM), phenylmethylsulfonyl fluoride (1 mM) and a protease inhibitor mixture (Sigma) as described.⁶⁹ Protein concentrations were determined using a Bradford reagent kit (Bio-Rad, Hercules, CA, USA).

Equal amounts of protein lysates were loaded and separated on 12% or 4–15% gradient Tris-HCl gels (Bio-Rad). Proteins were transferred onto PVDF membranes, which were blocked with 5% nonfat milk/5% BSA in TBST (10 mM Tris-HCl pH 8.0, 150 mM NaCl, 0.1%Tween-20). Primary antibodies were incubated overnight at 4°C with gentle agitation. Anticaveolin-caveolin-1 and -2 antibody (BD Transduction Laboratories, San Diego, CA, USA), anti-phospho-p70 S6 Kinase (Thr389) antibody, phospho-S6 ribosomal protein (Ser235/236) antibody, phospho-4E-BP1 (Ser65) antibody, phospho-Akt (Thr 308) antibody (all from Cell Signaling Technology, Beverly, MA, USA) were diluted 1 : 1000–5000 in blocking solution. Detection was performed with alkaline phosphatase coupled secondary antibodies (1 : 10 000, Santa Cruz) and ECF-substrate (Amersham, Piscataway, NJ, USA) using a Storm 860 system (Molecular Dynamics, Sunnyvale, CA, USA).

RNA extraction and Northern blotting

Total RNA was isolated with Trizol (Invitrogen), electrophoresed, and probed as described previously.²⁴As hybridization probes, we used a pHRed102 *Hind*-III restriction fragment encoding human HMG CoA-reductase (ATCC 57042). To ensure even loading, we hybridized the same blot with a β -actin cDNA fragment which was amplified by RT-PCR with β -actin specific primer (Clontech) from human HL-60 mRNA (Clontech). Northern blots were exposed and quantitated using a Molecular Dynamics Storm 860 system (Sunnyvale, CA, USA), normalized to β -actin and expressed as fold induction compared to control.

OCL formation in a hyperlipidemic mouse model: the LDLR-/- mouse

Six-week-old homozygous LDLR-null (LDLR-/-) mice and background strain C57BL6/J mice were obtained from the Jackson Laboratory (Bar Harbor, ME, USA). OCL formation was performed as described above. In

some experiments, a lipoprotein-deficient serum was used to replace FBS during OCL differentiation (10% LPDS instead of 10% FBS), as stated in figure legends. After 7–11 days, cocultures were fixed in 3.5% paraformaldehyde and triple-stained as described above. The percentage of large OCL (> 340 μ m in diameter) was determined by counting the total number of OCL that contain at least five nuclei in at least 40 observation fields per well under 10 \times magnification. Analyses were performed in triplicate, yielding at least 120 samples per group. In one set of experiments, the OCL area was quantified for 350 individual OCLs per group using the BioQuant IV image analysis system (BioQuant, Nashville, TN, USA), as noted in legends.

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