

Hedgehog Signaling and Osteogenic Differentiation in Multipotent Bone Marrow Stromal Cells Are Inhibited by Oxidative Stress

Woo-Kyun Kim,¹ Vicente Meliton,¹ Noam Bourquard,¹ Theodore J. Hahn,² and Farhad Parhami^{1*}

¹Department of Medicine, David Geffen School of Medicine at UCLA, Los Angeles, California

²VA Greater Los Angeles Healthcare System, Geriatric Research, Education, and Clinical Center, Los Angeles, California

ABSTRACT

Oxidative stress may play a major role in age-related osteoporosis in part by inhibiting osteoblast generation from osteoprogenitor cells. In the present study, we hypothesized that oxidative stress may inhibit the osteogenic differentiation of bone marrow stromal cells (MSC) in part by inhibiting the Hedgehog (Hh) signaling pathway, which is essential for bone development and maintenance and induces osteogenic differentiation of osteoprogenitor cells. To test this hypothesis, we examined the effects of oxidative stress on Sonic Hh (Shh)-induced osteogenic differentiation and signaling in M2-10B4 (M2) MSC, C3H10T1/2 embryonic fibroblasts, and mouse primary MSC. Treatment of cells with H₂O₂ inhibited Shh-induced osteogenic differentiation determined by the inhibition of Shh-induced expression of osteogenic differentiation markers alkaline phosphatase (ALP), osterix (OSX), and bone sialoprotein (BSP). Similar effects were found when oxidative stress was induced by xanthine/xanthine oxidase (XXO) or minimally oxidized LDL (MM-LDL). H₂O₂, XXO, and MM-LDL treatment inhibited Shh-induced expression of the Hh target genes Gli 1 and Patched1 as well as Gli-dependent transcriptional activity in M2 cells. H₂O₂ treatment also inhibited Hh signaling induced by the direct activation of Smoothened by purmorphamine (PM), but not by Gli1 overexpression. This suggests that oxidative stress may inhibit Hh signaling upstream of Gli activation and Gli-induced gene expression. These findings demonstrate for the first time that oxidative stress inhibits Hh signaling associated with osteogenic differentiation. Inhibition of Hh signaling-mediated osteogenic differentiation of osteoprogenitor cells may in part explain the inhibitory effects of oxidative stress on osteoblast development, differentiation, and maintenance in aging. *J. Cell. Biochem.* 111: 1199–1209, 2010. © 2010 Wiley-Liss, Inc.

KEY WORDS: OXIDATIVE STRESS; MESENCHYMAL STEM CELLS; HEDGEHOG; OSTEOGENESIS

Age-related osteoporotic bone loss is associated with decreased bone formation and increased bone resorption [Cummings and Melton, 2002]. Recent evidence suggests that oxidative stress may play a major role in this process [Almeida et al., 2007, 2009]. Aging is associated with increased oxidative stress and decreased antioxidant defenses [Garrett et al., 1990; Mody et al., 2001; Shouhed et al., 2005]. Reactive oxygen species (ROS), such as hydrogen peroxide (H₂O₂), superoxide (O₂⁻), and hydroxyl radical (OH[•]), can trigger oxidative damage in lipid, DNA, protein, carbohydrate, and proteoglycan molecules [Valiko et al., 2007]. In addition, oxidative stress has been linked to various aging disorders, including atherosclerosis, diabetes, neurodegenerative diseases, and inflammatory arthritis to name a few [Valiko et al., 2007]. It has been

suggested that oxidative stress may also contribute to age-related osteoporosis in part by decreasing bone formation through inhibition of the osteogenic differentiation of bone marrow stromal cells (MSC) [Garrett et al., 1990]. Oxidative stress induced by H₂O₂, xanthine oxidase, minimally oxidized low-density lipoprotein (MM-LDL), and lipid oxidation products inhibits osteogenic differentiation and bone formation [Mody et al., 2001; Shouhed et al., 2005; Almeida et al., 2009]. A role for MM-LDL in promoting atherosclerosis and regulating osteogenic differentiation has been suggested, with MM-LDL inducing oxidative stress, promoting osteogenic differentiation of vascular mesenchymal cells and inhibiting osteogenic differentiation of bone-derived osteoprogenitor cells [Parhami et al., 1997; Mody et al., 2001; Shouhed et al.,

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*Correspondence to: Farhad Parhami, PhD, David Geffen School of Medicine at UCLA, Center for the Health Sciences, BH-307, 10833 Le Conte Avenue, Los Angeles, CA 90095. E-mail: fparhami@mednet.ucla.edu

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2005]. In this regard, we have previously shown that antioxidants and specific osteogenic oxysterols that activate Hedgehog (Hh) signaling in osteoprogenitor cells effectively reverse the inhibitory effects of oxidative stress on osteogenic differentiation [Mody et al., 2001; Shouhed et al., 2005]. Moreover, oxidative stress induced by some forms of oxidized lipids inhibits osteoblastogenesis and increases adipogenesis by inhibiting Wnt signaling that is critical for the regulation of osteogenic and adipogenic differentiation of MSC [Almeida et al., 2009].

Hh signaling is one of the major signaling pathways that regulate osteogenesis and adipogenesis [Spinella-Jaegle et al., 2001; Suh et al., 2006]. The proteins of the Hh family are important signaling molecules that play a central role in embryonic bone development and post-embryonic bone homeostasis [McMahon et al., 2003; Kimura et al., 2008]. In vertebrates, the Hh family consists of three members: Sonic Hh (Shh), Indian Hh (Ihh), and Desert Hh (Dhh) [McMahon et al., 2003]. Hh protein binding to the transporter-like receptor Patched (Ptch) releases Ptch inhibition of Smoothed (Smo), a seven-pass transmembrane protein that transduces the Hh signal, which in turn activates members of the Gli family of DNA-binding proteins that mediate the transcription of Hh target genes in cells [McMahon et al., 2003; Beachy et al., 2004]. It has been shown that Hh signaling is involved in post-embryonic bone fracture healing and bone maintenance [Ito et al., 1999; Kimura et al., 2008]. In the initial stages of fracture repair, the expression of molecular components of Hh signaling, including Smo, Indian Hh, and Ptch, is increased [Ito et al., 1999]. Moreover, transient inhibition of Hh signaling in young mice causes severe defects in bone structure [Kimura et al., 2008]. Thus, impaired Hh signaling may be one of the mechanisms that underlie age-related osteoporosis and impaired healing of bone fractures. In the present report, we demonstrate that oxidative stress induced by H₂O₂, XXO, or MM-LDL is associated with inhibition of Hh signaling and reduced osteogenesis in pluripotent MSC that serve as the progenitors of osteogenic cells [D'ippolito et al., 1999], and whose differentiation is shifted in favor of adipogenesis at the expense of osteogenesis in aging and osteoporosis [Meunier et al., 1971; Beresford et al., 1992; Nuttall et al., 1998; Chan and Duque, 2002].

MATERIALS AND METHODS

CELL CULTURE AND REAGENTS

M2-10B4 (M2) pluripotent mouse MSC and C3H10T1/2 mouse embryonic fibroblasts were cultured as previously described [Richardson et al., 2007; Kim et al., 2009]. Cell treatment was performed in differentiation medium containing 5% fetal bovine serum (FBS), 50 µg/ml ascorbate, and 3 mM β-glycerophosphate. Primary MSC were isolated from 6 to 8 months old CD1 mice according to our previously published methods [Kha et al., 2004] and maintained in MSC basal medium and mouse mesenchymal stem cell stimulatory supplements from STEMCELL Technology (Vancouver, BC). H₂O₂, xanthine, and xanthine oxidase were purchased from Sigma-Aldrich, Co. (St. Louis, MO), purmorphamine (PM) was from Calbiochem (La Jolla, CA), and recombinant mouse Shh N-terminal peptide was from R&D Systems (Minneapolis, MN). XXO treatment was performed using xanthine (250 µM) and

xanthine oxidase (40 mU/ml) in osteogenic medium [Shouhed et al., 2005]. Polyclonal anti-mouse Patched1 antibody was purchased from Novus Biologicals (Littleton, CO), monoclonal anti-mouse Gli1 antibody was purchased from R&D Systems, and monoclonal anti-β-actin antibody was purchased from Sigma-Aldrich, and all antibodies were used according to manufacturer's instructions.

LIPOPROTEIN PREPARATION AND OXIDATION

Human LDL was isolated by density-gradient centrifugation of serum, and MM-LDL was prepared by iron oxidation of human LDL

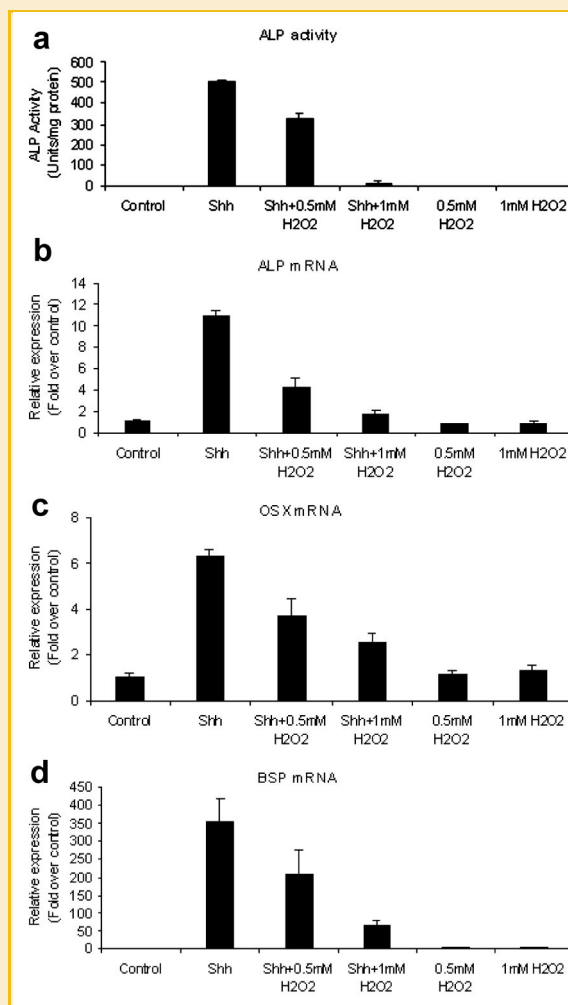


Fig. 1. Oxidative stress induced by H₂O₂ inhibits Shh-induced osteogenic differentiation of bone marrow stromal cells. a–d: M2-10B4 cells were treated at confluence with control vehicle, 200 ng/ml Shh, or 0.5–1 mM H₂O₂, alone or in combination. ALP activity was determined by a colorimetric method, and ALP and OSX mRNA expression were measured by quantitative real-time PCR 3 days after treatment. BSP mRNA expression was measured on day 6 post-treatment. Fold changes in gene expression relative to control are reported as the mean of triplicate determinations ±SD. (a: $P < 0.0001$ for Shh vs. Shh + 1 mM H₂O₂; $P < 0.01$ for Shh vs. Shh + 0.5 mM H₂O₂ or Shh + 1 mM H₂O₂; b: $P < 0.0001$ for Shh vs. Shh + 0.5 mM H₂O₂ or Shh + 1 mM H₂O₂; c, d: $P < 0.0001$ for Shh vs. Shh + 1 mM H₂O₂, $P < 0.001$ for Shh vs. Shh + 0.5 mM H₂O₂).

as previously described [Parhami et al., 1999]. The levels of lipoproteins used in the present study are reported in micrograms of protein. The lipoproteins were examined pre- and post-oxidation for lipopolysaccharide levels and found to have <30 pg of lipopolysaccharide/ml of medium.

QUANTITATIVE RT-PCR

Total RNA was extracted with an RNA isolation kit from Stratagene (La Jolla, CA) according to the manufacturer's instructions. RNA was

DNase-treated using a DNA-free kit from Ambion (Austin, TX). Three micrograms of RNA was reverse-transcribed using reverse transcriptase from Stratagene to make single-stranded cDNA. The cDNAs were then mixed with Qi SYBR Green Supermix (Bio-Rad, CA) for quantitative RT-PCR assay using a Bio-Rad I-cycler IQ quantitative thermocycler. All PCR samples were prepared in triplicate wells in a 96-well plate. After 40 cycles of PCR, melt curves were examined in order to ensure primer specificity. Fold changes in gene expression were calculated using the $\Delta\Delta C_t$ method. Primers

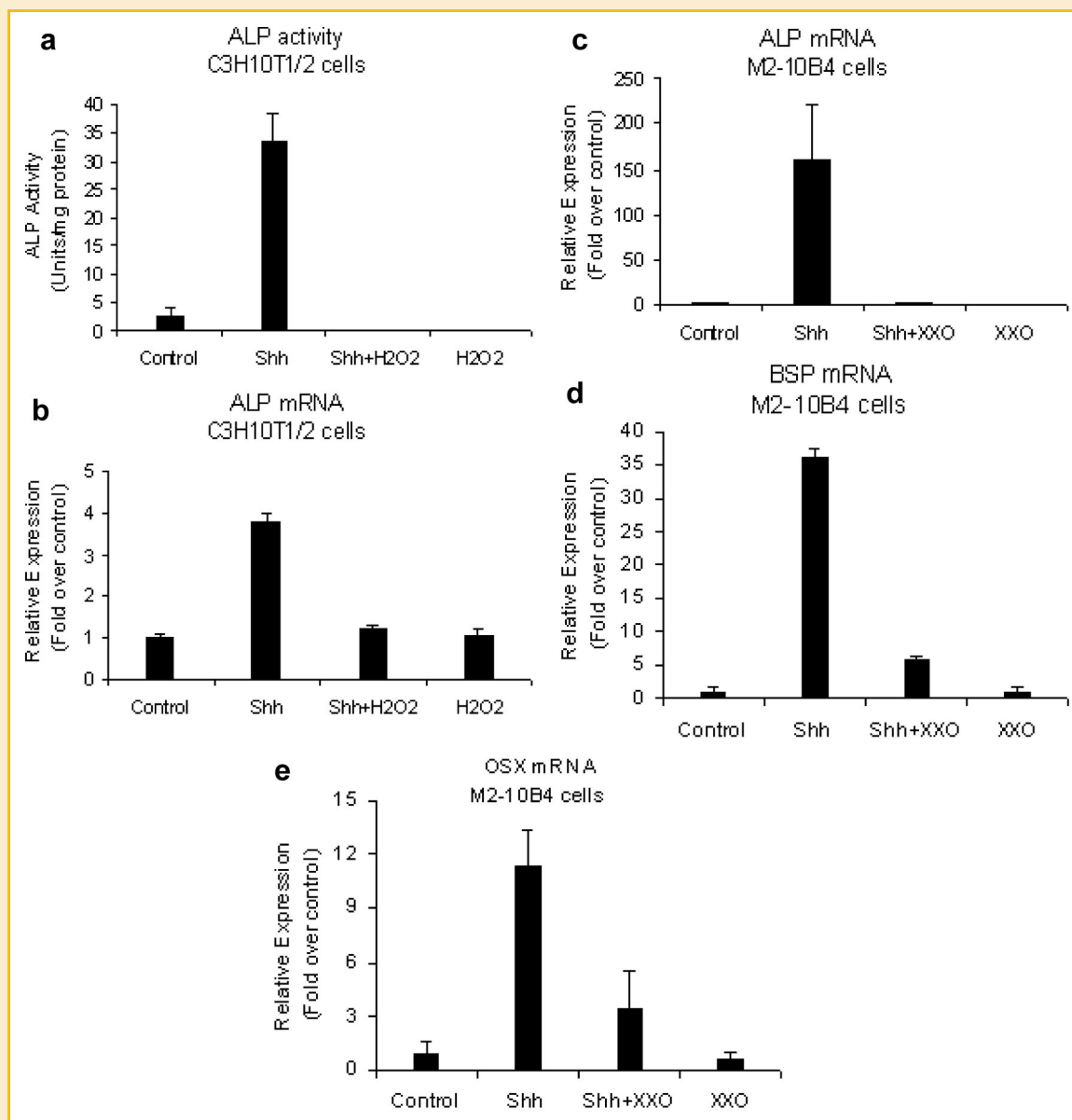


Fig. 2. Oxidative stress induced by H₂O₂ or XXO inhibits Shh-induced osteogenic differentiation in embryonic fibroblasts and bone marrow stromal cells. a: C3H10T1/2 cells at confluence were treated with control vehicle, 200 ng/ml Shh, or 0.75 mM H₂O₂ alone or in combination and ALP activity was measured after 3 days. b: C3H10T1/2 cells at confluence were treated with control vehicle, 200 ng/ml Shh, or 1 mM H₂O₂ alone or in combination for 48 h. ALP mRNA expression was measured by quantitative real-time PCR. Fold changes in gene expression relative to control are reported as the mean of triplicate determinations \pm SD. c–e: M2-10B4 cells were treated at confluence with control vehicle, 200 ng/ml recombinant mouse Shh or XXO alone or in combination for 4 days. ALP, BSP, and OSX mRNA expression were measured by quantitative real-time PCR. Fold changes in gene expression relative to control are reported as the mean of triplicate determinations \pm SD. (a,b: $P < 0.0001$ for control vs. Shh and for Shh vs. Shh + H₂O₂; c: $P < 0.0001$ for control vs. Shh and for Shh vs. Shh + XXO).

used were as follows: *GAPDH* (5'-ATGGACTGTGGTCATGAGCC-3' and 5'-ATTGTACGAATGCATCTG-3'), *Gli1* (5'-GAAGCCGAGCCGAGTATC-3' and 5'-GGTGTAGACAGAGGTTGG-3'), *Ptch1* (5'-CCATCGGCGACAAGAACC-3' and 5'-CCAGCACAGCAAAGAAA-TACC-3'), *Smoothed (Smo)* (5'-AACTATCGGTACCGTGCTGG-3' and 5'-CATCATGGGAGACAGTGTGC-3'), *c-fos* (5'-TGAGCTTC-CACCC-3' and 5'-CTCCAGGTTGCTGA-3'), *Hemoxygenase 1 (HOX1)* (5-CAGGTGATGCTGACAGAGGA-3' and 5'-GAGAGTGAGGACC-CACTGGA-3'), *Osterix (OSX)* (5'-CCCTTCTCAAGCACCAATGG-3' and 5'-AGGGTGGGTAGTCATTGTCAT-3'), *Alkaline Phosphatase (ALP)* (5'-AAACCCAGAACACAAGCATTCC-3' and 5'-TCCACCAG-CAAGAAGAAGCC-3'), and *Bone sialoprotein (BSP)* (5'-ACGCCA-CACTTCCACACTCTC-3' and 5'-TTCCTTCTCTTCTTCTTCTTCC-3').

ALKALINE PHOSPHATASE (ALP) ACTIVITY ASSAY

M2 cells at confluence were treated in differentiation medium with control vehicle or test agents. After 72 h, a colorimetric assay for ALP activity was performed on whole cell extracts as previously described [Kha et al., 2004].

MTT ASSAY

The MTT assay was used to measure the potential effects of oxidative stress inducing agents on cell proliferation and/or viability as we have previously reported [Mody et al., 2001].

TRANSIENT TRANSFECTION AND REPORTER ASSAY

M2 cells at 70% confluence were transiently transfected with a Gli response-element reporter construct (pGL3b-8xGli-Luciferase) plasmid, or pGL3b-Luciferase plasmid and pTK-Renilla-Luciferase plasmid as we have previously reported [Dwyer et al., 2007]. After 24 h cells were treated with test agents for 48 h. For Gli1 overexpression studies, M2 cells at 70% confluence were transiently transfected with a Gli1 overexpression vector (kind gift of Dr. William Matsui, Johns Hopkins School of Medicine), pGL3b-8xGli-Luciferase plasmid, or pGL3b-Luciferase plasmid and pTK-Renilla-Luciferase plasmid without Gli1 overexpression vector. Luciferase activity was measured after 48 h and normalized for transfection efficiency using the Renilla luciferase activity. Data are reported as the mean of triplicate determinations \pm SD.

WESTERN BLOTTING

Cells were lysed in lysis buffer, and protein concentrations were determined using the Bio-Rad protein assay as previously described [Kha et al., 2004]. Whole cells extracts (30 μ g) were separated on SDS-PAGE gels and transferred overnight onto nitrocellulose membrane. Blots were blocked and incubated with specific primary antibodies followed by HRP-linked secondary antibodies, and binding was revealed using the Western Lightning ECL Kit (Perkin Elmer, Boston, MA) according to the manufacturer's instructions.

STATISTICAL ANALYSIS

Statistical analyses were performed using the StatView 5 program. All *P* values were calculated using ANOVA and Fisher's projected least significant difference (PLSD) significance test. A value of *P* < 0.05 was considered significant.

RESULTS

OXIDATIVE STRESS INHIBITS SHH-INDUCED OSTEOGENESIS IN OSTEOPROGENITOR CELLS

We previously reported that oxidative stress induced by XXO or MM-LDL inhibits spontaneous in vitro osteogenic differentiation of MSC cultured in the presence of FBS, ascorbic acid, and β -glycerophosphate (i.e., osteogenic differentiation medium) [Shouhed et al., 2005]. In the present study, we examined whether oxidative stress induced by H_2O_2 inhibits Shh-induced osteogenesis in MSC by assessing its effect on markers of osteogenic differentiation, ALP activity, and ALP, OSX, and BSP mRNA expression, in M2 cells. H_2O_2 significantly reduced Shh-induced ALP activity in a dose-dependent manner (Fig. 1a). Furthermore, both concentrations of H_2O_2 inhibited Shh-induced ALP, OSX, and BSP mRNA expression measured by qRT-PCR (Fig. 1b-d), indicating that H_2O_2 -induced oxidative stress inhibits Shh-induced osteogenesis in MSC. H_2O_2 treatment also inhibited Shh-induced ALP activity as well as ALP mRNA expression in C3H10T1/2 embryonic fibroblasts (Fig. 2a,b). Furthermore, consistent with the effects of H_2O_2 , XXO, which induces oxidative stress by converting xanthine to uric acid and superoxide anion, significantly inhibited Shh-induced ALP, BSP, and OSX mRNA expression in M2 cells (Fig. 2c-e). We also confirmed that H_2O_2 produces oxidative stress in MSC by measuring its effects on the expression of key oxidative

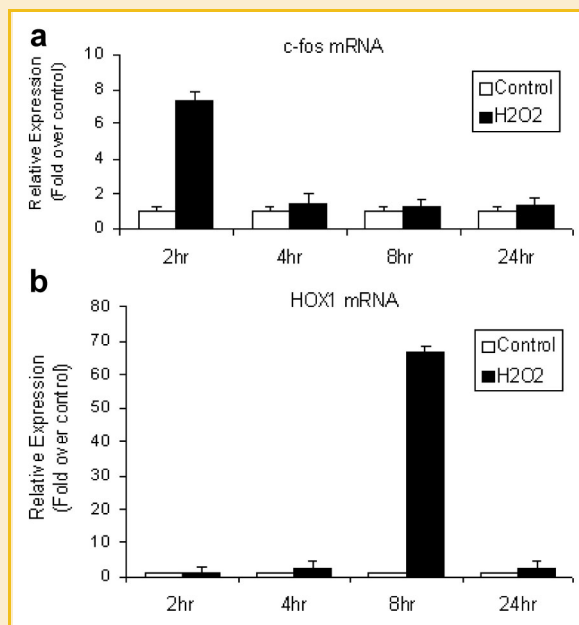


Fig. 3. Effect of H_2O_2 on oxidative stress-related gene expression in bone marrow stromal cells. M2-10B4 cells at confluence were treated with control vehicle or 1 mM H_2O_2 . *c-fos* and *HOX1* mRNA expression were measured by quantitative real-time PCR at 2, 4, 8, and 24 h post-treatment. Fold changes in gene expression relative to control cells are reported as the mean of triplicate determinations \pm SD. (a: *P* < 0.0001 for control vs. 1 mM H_2O_2 at 8 h and *P* < 0.001 for control vs. 1 mM H_2O_2 at 4 and 24 h; b: *P* < 0.0001 for control vs. 1 mM H_2O_2 at 2 h).

stress-related genes, including c-fos and HOX1 [Orozco et al., 2007] at 2, 4, 8, and 24 h after the addition of H₂O₂. The expression of c-fos mRNA was increased over sevenfold at 2 h in H₂O₂-treated cells ($P < 0.0001$) and returned to baseline levels after 4 h (Fig. 3a). HOX1 mRNA expression was also significantly increased by H₂O₂ treatment showing a robust 60-fold induction at 8 h, with a slight but significant 2.5-fold induction still present at 24 h (Fig. 3b).

To examine the potential effects of H₂O₂, XXO, and MM-LDL on proliferation and/or cell viability, we performed MTT assays anticipating that a decrease in proliferation and/or cell viability would be associated with a significant decrease in MTT values. Results showed that in M2 cells after 8, 24, 48, or 96 h, 0.5 mM H₂O₂ caused no significant effects in MTT values. Treatment of M2 cells with 1 mM H₂O₂ caused no significant effects at 8 and 24 h

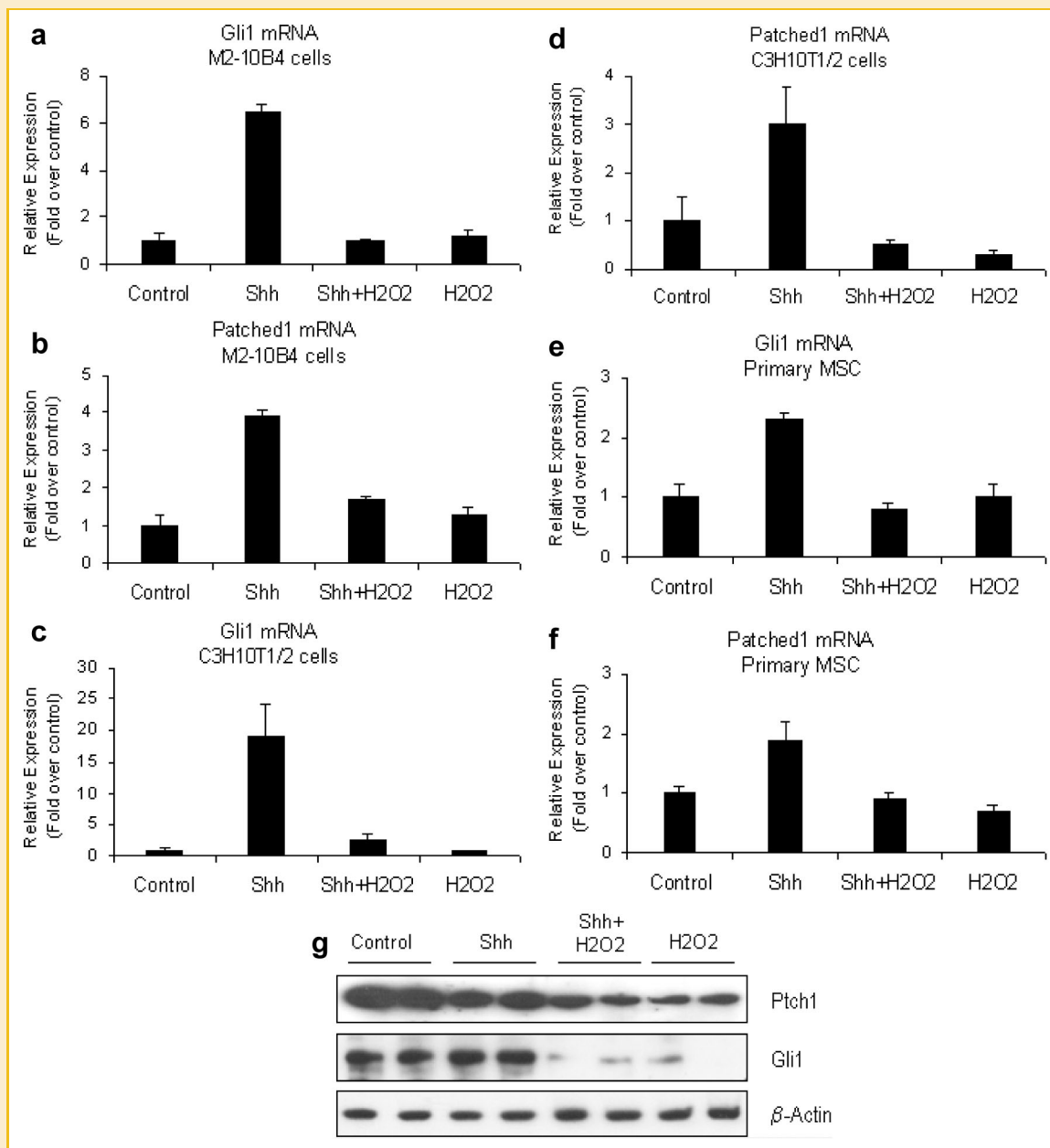


Fig. 4. Oxidative stress induced by H₂O₂ inhibits Hh signaling in osteoprogenitor cells. a,b: M2-10B4 cells at confluence were treated with control vehicle, 200 ng/ml Shh, or 1 mM H₂O₂ alone or in combination for 8 h. Gli1 and Patched1 mRNA expression were measured by quantitative real-time PCR. c,d: C3H10T1/2 cells at confluence were treated with control vehicle, 200 ng/ml Shh, or 1 mM H₂O₂ alone or in combination for 48 h. Gli1 and Patched1 mRNA expression were measured. e,f: Primary MSC at confluence were treated with control vehicle, 600 ng/ml Shh, or 0.5 mM H₂O₂ alone or in combination for 24 h. Gli1 and Patched1 mRNA expression were measured. Fold changes in gene expression relative to control cells are reported as the mean of triplicate determinations \pm SD. ($P < 0.001$ for control vs. Shh and for Shh vs. Shh + H₂O₂ in all panels shown). g: M2-10B4 cells were treated with 200 ng/ml Shh or 1 mM H₂O₂ alone or in combination for 24 h. Western blot analysis using whole cell lysates was performed using specific antibodies to Gli1, Ptch1, and β -actin for normalization. Results showing duplicate samples from each condition and representative of two separate experiments are reported.

but did cause a significant 20% decrease after 48 h ($P < 0.01$) and a significant 30% decrease after 96 h ($P < 0.01$) without any evidence of cell rounding or detachment from the monolayer up to 8 days post-treatment. XXO and MM-LDL caused no effects on MTT values up to 96 h post-treatment, and caused no evidence of cell rounding or detachment for the entire duration of the experiments.

OXIDATIVE STRESS INHIBITS HEDGEHOG SIGNALING IN BONE MARROW STROMAL CELLS

Next, we examined the effects of oxidative stress on Shh-induced Hh signaling in M2 cells. Shh increased the mRNA expression of the key Hh signaling target genes, Gli1 and Ptch1 in M2 cells after 8 h of treatment, and these effects were significantly inhibited by H₂O₂ (Fig. 4a,b). These results were confirmed in C3H10T1/2 mouse embryonic fibroblasts (Fig. 4c,d) and in primary murine MSC

where H₂O₂ treatment was also found to significantly reduce Shh-induced Gli1 and Ptch1 mRNA expression (Fig. 4e,f). Although previous reports have demonstrated the effect of Hh pathway stimulation on the subcellular distribution and localization of molecular components of the Hh pathway, including Ptch1, Smo, and Gli2, to primary cilia [Rohatgi and Scott, 2007; Milenkovic et al., 2009], we were interested in examining the potential effects of oxidative stress on absolute levels of Ptch1 and Gli1 proteins which were abundantly present at baseline in extracts from confluent cultures of M2 cells (Fig. 4g). In order to examine the effects of H₂O₂ on Gli1 and Ptch1 protein levels, M2 cells were treated with Shh or H₂O₂, separately or in combination, and whole cell extracts were analyzed after 24 h of treatment by Western blotting. In contrast to its previously reported inductive effects on their mRNA expression, Shh had almost no effect on Gli1 protein levels and a slightly inhibitory effect on Ptch1 protein levels (Fig. 4g). However, H₂O₂

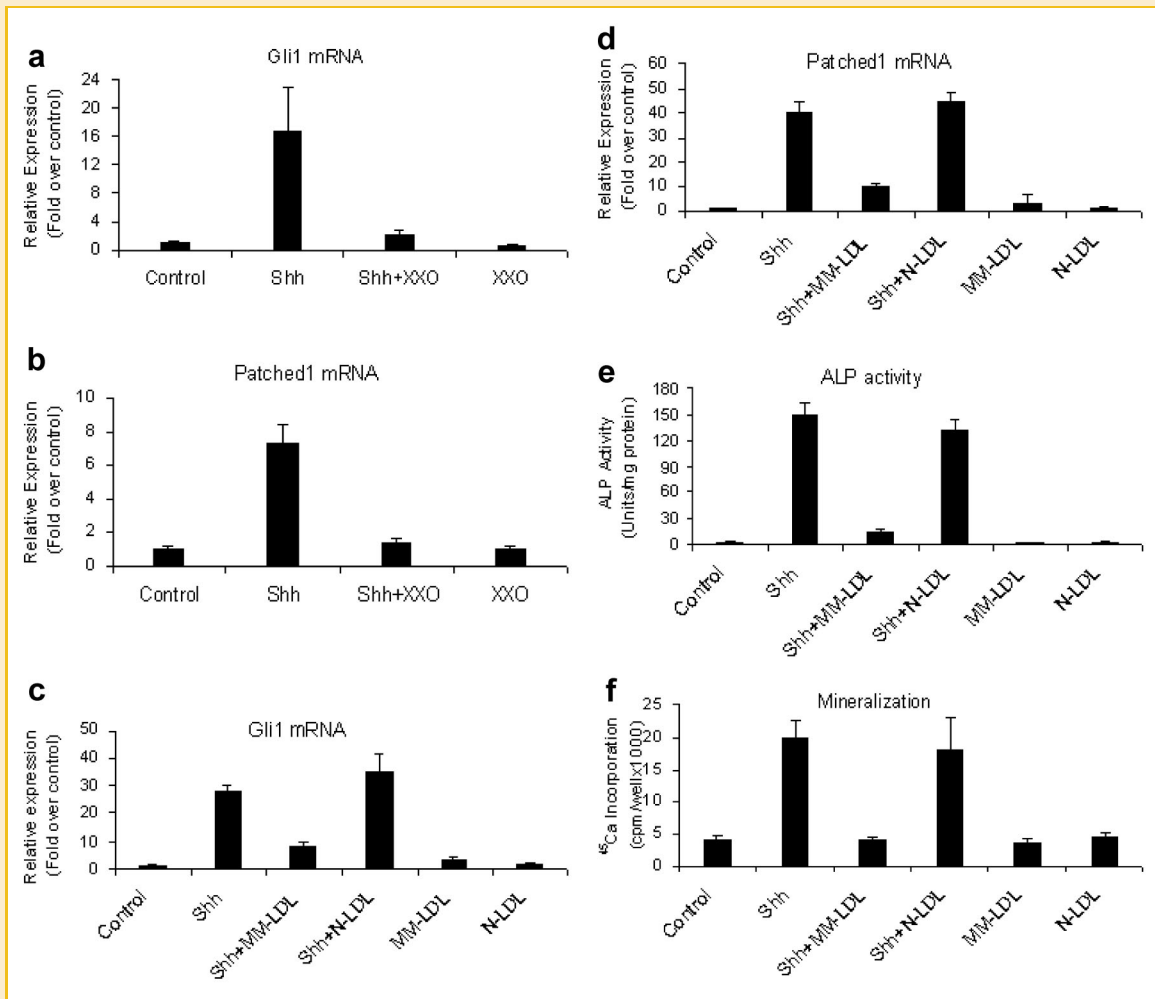


Fig. 5. Oxidative stress induced by XXO or MM-LDL inhibits Hh signaling and osteogenic differentiation of bone marrow stromal cells. a,b: M2-10B4 cells at confluence were treated with control vehicle, 200 ng/ml Shh, or XXO alone or in combination for 24 h. Gli1 and Patched1 mRNA expression were measured by quantitative real-time PCR. c–f: M2-104 cells at confluence were treated with control vehicle, 200 ng/ml Shh, MM-LDL (200 μ g/ml), or native LDL (N-LDL; 200 μ g/ml) alone or in combination. Gli1 and Patched1 mRNA expression were measured by quantitative real-time PCR at 48 h post-treatment. Fold changes in gene expression relative to control cells are reported as the mean of triplicate determinations \pm SD. ALP activity was assayed in whole cell lysates after 3 days (e), and mineralization in parallel cultures was quantified after 14 days of treatments (f). (a,b: $P < 0.001$ for control vs. Shh and for Shh vs. Shh + XXO; c–f: $P < 0.001$ for control vs. Shh and for Shh vs. Shh + MM-LDL).

robustly reduced the protein levels of both Gli1 and Ptch1 in control as well as in Shh-treated cells (Fig. 4g).

In addition, we examined whether oxidative stress induced by XXO and MM-LDL inhibits Hh signaling in M2 cells. XXO and MM-LDL treatment significantly inhibited Gli1 and Ptch1 mRNA expression (Fig. 5a–d), whereas treatment with unoxidized native LDL (N-LDL), which does not induce oxidative stress, did not inhibit the expression of these genes. We also found that MM-LDL significantly inhibited Shh-induced osteogenic differentiation of M2 cells as suggested by the inhibition of ALP activity and mineralization in these cells (Fig. 5e,f).

KINETICS OF OXIDATIVE STRESS-INDUCED INHIBITION OF HEDGEHOG SIGNALING

We examined the kinetics of H₂O₂ effects on Hh signaling in M2 cells by testing the effects of H₂O₂ when added prior to or after treating the cells with Shh. An 8 h pretreatment time point with H₂O₂ was chosen since H₂O₂ induction of oxidative stress occurred as early as 2 h after treatment as evident by the induction of c-fos, and was persistent for at least 8 h as evident by the robust induction of HOX1

at this time point (Fig. 3b). Subsequent to the 8 h pretreatment, H₂O₂ was completely removed from the cells and the cells were then treated with Shh for an additional 24 h in order to examine whether the inhibitory effects of H₂O₂ are persistent or transient and reversible upon its removal. Results showed that the effects of H₂O₂ appear to be transient and therefore the fold changes in the induction of Gli1 and Ptch1 are similar with and without H₂O₂ pretreatment (Fig. 6a,b). Next we examined whether H₂O₂ is capable of inhibiting Hh signaling if the cells have encountered Shh prior to the induction of oxidative stress by H₂O₂. Similar to co-treatment of cells with Shh and H₂O₂, treatment of M2 cells with Shh first for 16 h followed by the addition of H₂O₂ for 8 h in the presence of Shh produced a significant inhibition of Gli1 and Ptch1 mRNA expression (Fig. 6c,d) while significantly inducing HOX1 expression (Fig. 6e).

MOLECULAR MECHANISMS UNDERLYING THE INHIBITORY EFFECTS OF OXIDATIVE STRESS ON HEDGEHOG SIGNALING

In order to begin identifying the molecular mechanisms by which oxidative stress inhibits cellular responses to Hh proteins, we

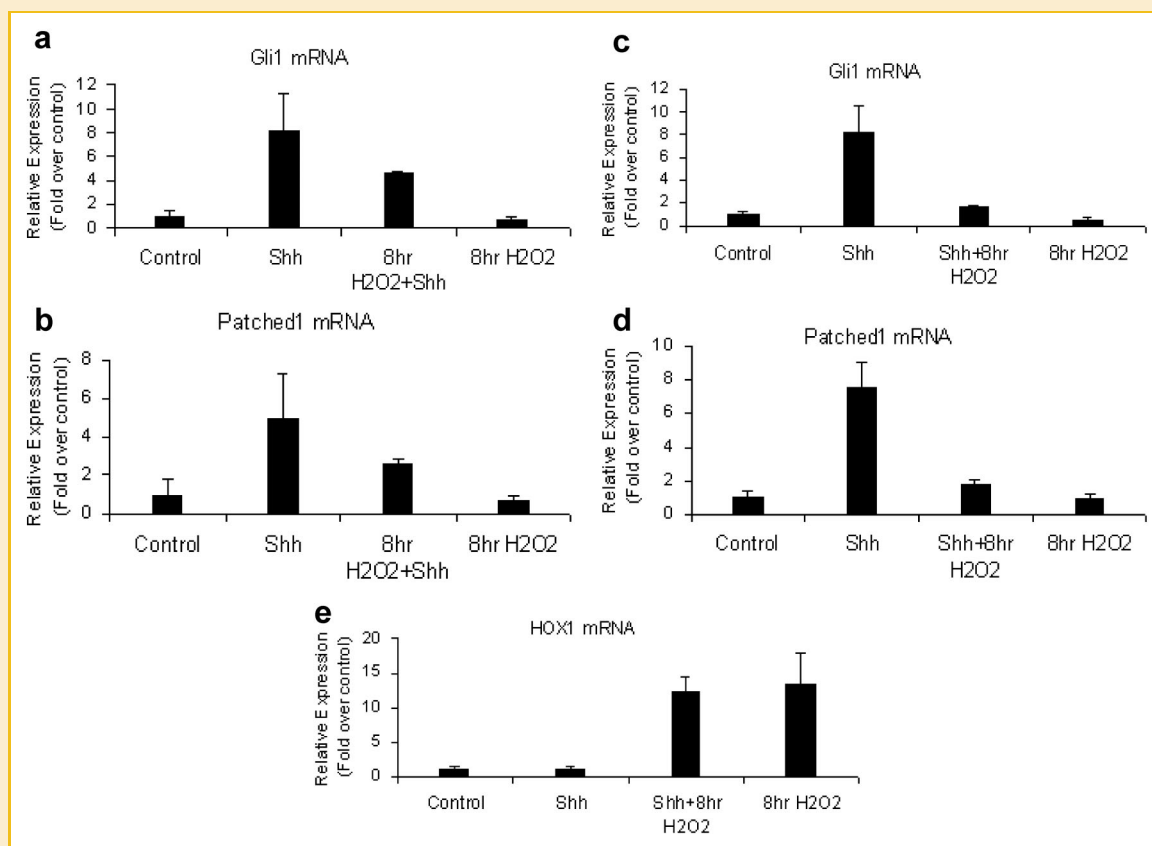


Fig. 6. Kinetics of inhibition of Hh signaling by oxidative stress in bone marrow stromal cells (a,b). M2-10B2 cells at confluence were pretreated with 1 mM H₂O₂ for 8 h. Next, H₂O₂ was removed and the cells were treated with control vehicle or 200 ng/ml Shh for 24 h. Gli1 and Patched1 mRNA expression were measured by quantitative real-time PCR. c–e: M2-10B4 cells at confluence were treated with control vehicle or 200 ng/ml Shh for 24 h. At 16 h post-treatment, 1 mM H₂O₂ was added into Shh + H₂O₂ or H₂O₂ treatment groups for the remaining 8 h. Gli1, Patched1, and HOX1 mRNA expression were measured by quantitative real-time PCR. Fold changes in gene expression relative to control cells are reported as the mean of triplicate determinations ±SD. (a,b: $P < 0.001$ for control vs. Shh and $P < 0.05$ for Shh vs. 8 h H₂O₂ + Shh; c,d: $P < 0.0001$ for control vs. Shh or Shh + 8 h H₂O₂; e: $P < 0.0001$ for control vs. Shh + 8 h H₂O₂ or 8 h H₂O₂).

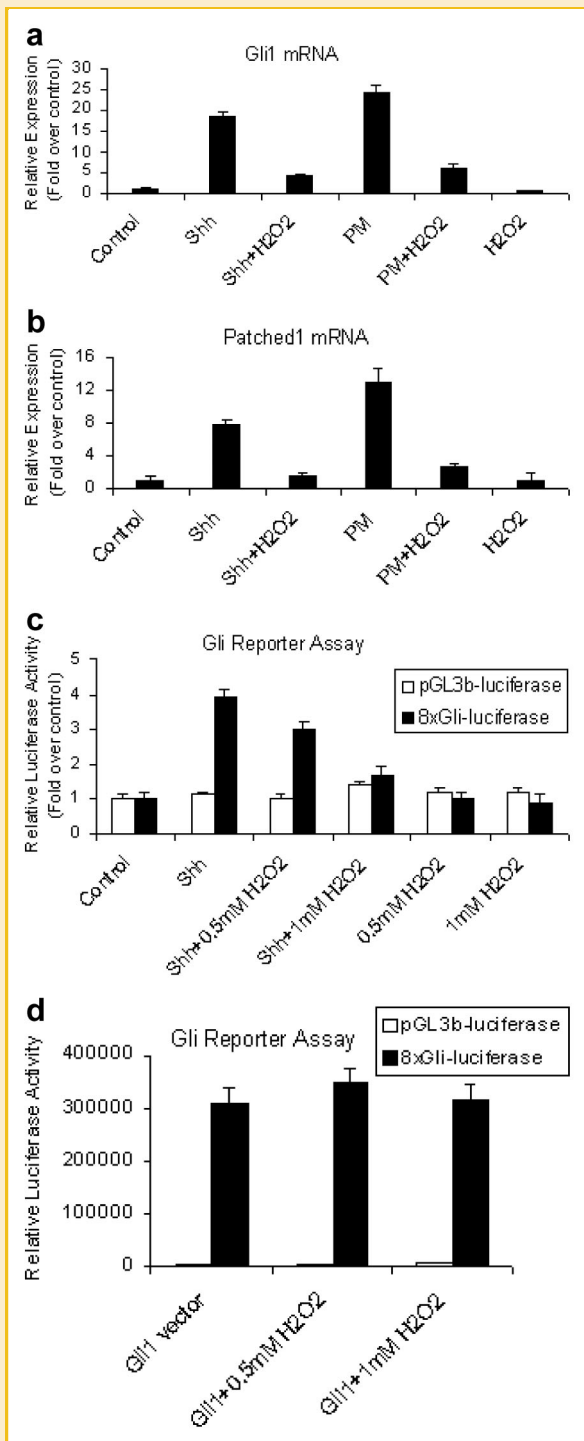
examined the effects of H₂O₂ on Hh signaling induced by the binding of Shh to Ptch versus that which is induced by Smo agonists that directly bind to Smo and bypass the Ptch regulatory step in activating Hh signaling. Direct activation of Smo was achieved by treating M2 cells with PM, which activates Hh signaling by direct binding to Smo [Sinha and Chen, 2006]. As expected, H₂O₂ treatment significantly inhibited Shh-induced Gli1 and Ptch1

mRNA expression (Fig. 7a,b). Similarly, H₂O₂ treatment also inhibited PM-induced Gli1 and Ptch1 mRNA expression (Fig. 7a,b), indicating that oxidative stress inhibits Hh signaling downstream or at the level of Smo. This is consistent with the robust H₂O₂-induced reduction in Gli1 protein levels which is in part responsible for transducing the transcription regulating signals from Smo to the nucleus. Furthermore, in preliminary unpublished studies we have found that treatment of M2 cells for 24 h with 1 mM H₂O₂ resulted in a significant 70% inhibition of Smo mRNA expression. In addition, H₂O₂ treatment significantly inhibited Shh-induced Gli reporter activity in a dose-dependent manner in M2 cells (Fig. 7c). However, H₂O₂ did not inhibit Gli reporter activity induced by the overexpression of Gli1 (Fig. 7d), suggesting that inhibition of Hh signaling occurs upstream of Gli-induced Hh target gene expression and may involve inhibition of Gli1 expression which is compensated by the overexpression of Gli1.

DISCUSSION

Several studies have suggested an association between oxidative stress and impairment of skeletal integrity, and a potentially important role for antioxidants in preventing age-related osteoporosis [Basu et al., 2001; Maggio et al., 2003; Shouhed et al., 2005]. Levels of 8-iso-PGF_{2α}, a circulating biomarker of oxidative stress that increases with age were reported to be negatively associated with bone mineral density in both women and men [Basu et al., 2001]. In mice increased oxidative stress with age was associated with reduced osteoblast number and bone formation rate as well as increased osteoblast and osteocyte apoptosis, decreased bone mineral density, and decreased bone strength [Almeida et al., 2007]. Decreased osteogenic differentiation of bone marrow progenitor cells could reduce osteoblast number and bone formation, causing or accelerating age-related osteoporosis [Chan and Duque, 2002; Chen et al., 2002]. Given the importance of Hh signaling in embryonic bone development and post-embryonic maintenance and repair of bone, we focused our attention on the potential adverse effects of oxidative stress on Hh-induced

Fig. 7. Mechanism of oxidative stress-induced inhibition of Hh signaling. a,b: M2-10B4 cells at confluence were treated with control vehicle, 200 ng/ml Shh, 1 μM PM, or 1 mM H₂O₂ alone or in combination for 48 h. Gli1 and Patched1 mRNA expression were measured by quantitative real-time PCR. Fold changes in gene expression relative to control cells are reported as the mean of triplicate determinations ±SD. c: M2-10B4 cells at 70% confluence were transiently transfected with a Gli response element reporter construct (pGL3b-8xGli-Luciferase) or pGL3b-Luciferase plasmid in combination with pTK-Renilla-Luciferase plasmid. After 24 h cells were treated with control vehicle, 200 ng/ml Shh, 0.5 or 1 mM H₂O₂ alone or in combination for 48 h. d: M2 cells at 70% confluence were transiently transfected with a Gli1 overexpression vector, a Gli response element reporter construct (pGL3b-8xGli-Luciferase) or pGL3b-Luciferase plasmid in combination with pTK-Renilla-Luciferase plasmid. Luciferase activity was measured after 48 h and normalized for transfection efficiency using the Renilla luciferase activity. Data are reported as the mean of triplicate determinations ±SD. (a,b: $P < 0.0001$ for control vs. Shh or PM, for Shh vs. Shh + H₂O₂ and for PM vs. PM + H₂O₂; c: $P < 0.0001$ for control vs. Shh and for Shh vs. Shh + 1 mM H₂O₂ and $P < 0.01$ for Shh vs. Shh + 0.5 mM H₂O₂).



osteogenesis and Hh signaling in MSC. Indeed, the studies presented here clearly demonstrate the inhibitory effects of oxidative stress on Hh-induced osteogenic differentiation of MSC and support the hypothesis that the contribution of oxidative stress to osteoporosis may be in part due to inhibition of Hh signaling-mediated osteogenesis. In addition, we speculate that oxidative stress may also inhibit Hh-mediated skeletal development, consistent with reports that showed inhibition of Hh-mediated limb patterning and neural plate closure by oxidative stress during development [Dennerly, 2007; Aoto et al., 2008]. We propose that impairment of Hh signaling by oxidative stress in adult organisms may have detrimental effects on tissue homeostasis including the skeleton. Oxidative stress may be induced by a variety of physiologically relevant factors that may interfere with Hh signaling. In this report we demonstrate that inflammatory oxidized lipoproteins, such as MM-LDL, which we have shown in the past to impair osteogenic differentiation of osteoprogenitor cells, also disrupt Hh signaling in these cells. This is of particular relevance to the previously suggested role of dyslipidemia and lipid oxidation in bone maintenance and osteoporosis [Parhami et al., 1999, 2001; Parhami, 2003; Almeida et al., 2009]. The importance of this finding, if translated *in vivo*, will be at least twofold, both in the context of embryonic development and in post-embryonic tissue homeostasis where Hh signaling plays critical roles. Inflammatory oxidized lipoproteins play fundamental roles in the development of atherosclerosis in mice and humans [Rosenfeld, 1991], and in osteopenia/osteoporosis in mice [Parhami et al., 1999, 2001]. Since Hh signaling appears to be important in both these disorders by regulating homeostasis of vessel wall and bone/cartilage [Ehlen et al., 2006; Beckers et al., 2007], respectively, we propose that understanding the molecular interactions between inflammatory oxidized lipoproteins and Hh signaling may provide new insights into the pathogenesis of cardiovascular disease and osteoporosis.

Moreover, we demonstrate here that oxidative stress induced by H₂O₂ inhibits Hh signaling induced by Hh proteins or by ligands such as PM that directly bind to and activate Smo. Together with our finding that activation of Hh signaling by Gli overexpression is not inhibited by oxidative stress, that oxidative stress robustly inhibits Gli1 protein levels, and our preliminary observation that oxidative stress appears to significantly inhibit Smo mRNA expression, we propose that oxidative stress inhibits Hh signaling in part by inhibiting the expression of Gli1 and/or by inhibiting events that occur upstream of Gli activation. Since Smo is the sole transducer of the Hh-mediated signaling and target gene expression, down-regulation of Smo expression may be the main mechanism underlying the inhibition of Hh signaling by oxidative stress. However, since regulation of Smo activity in part involves complex changes in subcellular localization of Smo protein to primary cilia [Milenkovic et al., 2009], in future studies it will be necessary to further examine the effects of oxidative stress on Smo protein localization and trafficking within subcellular compartments. Furthermore, it has been well established that Ptch causes catalytic inhibition of Smo [Taipale et al., 2000]. At present the mechanism for this catalytic inhibition is not clearly understood, although the potential role of intracellular small molecules such as specific oxysterols has been suggested [Beachy et al., 2004; Dwyer et al.,

2007; Rohatgi et al., 2007]. It is intriguing to speculate that these small molecule regulators of Ptch–Smo interaction may be targets of oxidative stress that in addition to down regulation of Smo expression may result in impairment of Hh signaling and osteogenic differentiation in cells exposed to oxidative stress.

As noted earlier, recent studies have suggested that primary cilia and the molecular machinery within them are important for the integrity of Hh signaling in mammalian cells [Rohatgi et al., 2007; Wong et al., 2009]. In preliminary studies we have found that in M2 cell oxidative stress induced by H₂O₂ significantly reduces the mRNA expression of Polaris (Parhami et al., unpublished work), a key component of primary cilia whose knockdown has been found to adversely affect primary cilia formation and Hh signaling [Huangfu and Anderson, 2005; May et al., 2005]. Since primary cilia have been reported to also mediate other cell signaling pathways, the previously reported adverse effects of oxidative stress on multiple signaling pathways is consistent with oxidative stress-induced impairment of primary cilia [Simons et al., 2005; Michaud and Yoder, 2006; Alaiwi et al., 2009; Wong et al., 2009]. Disruption of primary cilia and intraflagellar transport within cilia were found to impair normal endochondral bone formation due to disruption of multiple signaling pathways [Haycraft et al., 2007]. Since plasma membrane is highly susceptible to oxidative stress and lipid oxidation, changes in its physical properties and composition often occur with oxidation [Joseph et al., 1998; Stark, 2005; Cuschieri and Maier, 2007]. In fact, microdomains in plasma membrane that centralize signal transduction, such as lipid rafts and caveolae, are targets of oxidative stress and undergo compositional changes that modulate cellular signaling and behavior [Girotti, 2001; Cuschieri and Maier, 2007]. Since primary cilia are plasma membrane microdomains that control signal transduction, we hypothesize that oxidative stress will induce changes in physical properties of ciliary membrane, causing alterations in localization of key membrane-bound members of Hh pathway signaling, Ptch and Smo. This hypothesis remains to be tested in future studies.

In summary, our studies demonstrate for the first time that oxidative stress inhibits Hh signaling and Hh-mediated osteogenic differentiation in osteoprogenitor cells. Inhibition of Hh signaling and osteogenesis by oxidative stress may in part explain the reduced integrity of the skeletal tissue seen with aging, and together with our finding that the inhibitory effects of oxidative stress on Hh signaling appear to be transient, reemphasizes the potential benefits of antioxidants in preventing and/or reversing oxidative stress-induced osteoporosis.

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