Original Research

Epigallocatechin-3-gallate protects against tumor necrosis factor alpha induced inhibition of osteogenesis of mesenchymal stem cells

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Abstract

Anabolic bone accruement through osteogenic differentiation is important for the maintenance of physiological bone mass and often disrupted in various inflammatory diseases. Epigallocatechin-3-gallate, as an antioxidant and anti-inflammatory agent, has been suggested for potential therapeutic use in this context, possibly by the inhibition of bone resorption as well as the enhancement of bone formation through directly activating osteoblast differentiation. However, the reported effects of epigallocatechin-3-gallate modulating osteoblast differentiation are mixed, and the underlying molecular mechanism is still elusive. Moreover, there is limited information regarding the effects of epigallocatechin-3-gallate on osteogenic potential of mesenchymal stem cell in inflammation. Here, we examined the *in vitro* osteogenic differentiation of human mesenchymal stem cells. We found that the cell viability and osteoblast differentiation of human bone marrow-derived mesenchymal stem cells are significantly inhibited by inflammatory cytokine TNF α treatment. Epigallocatechin-3-gallate is able to enhance the cell viability and osteoblast differentiation of neversing the TNF α -induced inhibition. Notably, only low doses of epigallocatechin-3-gallate have such benefits, which potentially act through the inhibition of NF- κ B signaling that is stimulated by TNF α . These data altogether clarify the controversy on epigallocatechin-3-gallate promoting osteoblast differentiation and further provide molecular basis for the putative clinical use of epigallocatechin-3-gallate in stem cell-based bone regeneration for inflammatory bone loss diseases, such as rheumatoid arthritis and prosthetic osteolysis.

Keywords: EGCG, TNFα, osteoblast differentiation, mesenchymal stem cell, inflammatory bone disease, NF-κB

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Introduction

Patients with chronic inflammatory bone diseases, including arthritis, osteomyelitis, periodontal bone loss, and wear debris-induced osteolysis, not only suffered from the acceleration of bone resorption (osteoclast activation) but also devastated by a dramatic loss of bone formation activity (osteoblast differentiation).^{1–5} Such a dilemma remains a challenge for treating numerous inflammatory diseases that are frequently associated with severe bone loss.^{3,6,7} In complementary to the current treatment strategies involving the blockade of bone resorption,^{8–10} it is often needed for the regeneration of collagen architecture through bone neo-formation.^{3,6} The feasibility of cell-based approaches,¹¹ particular using bone marrow-derived mesenchymal stem cell (BM-MSC), thus presents an exciting option for such purposes to repair bone lesions by promoting the differentiation of autologous bone cell (including chondrocyte, osteoblast, and osteocyte). Preliminary successes of MSC therapy on animal models of inflammatory bone diseases¹²⁻¹⁶ further prompted basic research to investigate the potential mechanisms on the refinement of bone tissue repair by MSCs.

As a critical mediator of pro-inflammation, TNF α stimulates strong systemic effects as well as local responses in bone disease.¹⁷⁻¹⁹ Both arms of bone remodeling can be affected by TNF α , resulting synergistic bone loss. The activation of TNF α on bone resorption has been extensively studied in each step of osteoclast activation,²⁰⁻²³ including expansion of osteoclast precursor, amplification of RANKL/RANK-mediated osteoclast differentiation, and facilitation of osteoclast specialized function. The less clear is the effect of inflammation on bone formation. Consistent with a negative impact on bone formation, $\text{TNF}\alpha$ can induce cell death of osteogenic cells and block osteoblast differentiation *in vitro*.^{24–26} However, its molecular target is unknown. Important candidates include bone maturation marker (e.g. Runx2, alkaline phosphatase (ALP)) and Wnt signaling,^{27,28} both of which are repressed by TNF α through a variety of pathways (e.g. oxidative stress and/or its major downstream activator NF- κ B²⁹). In addition, its effect on MSC differentiation is unclear, and whether it could also inhibit bone regeneration from BM-MSC is an open question.

Epigallocatechin-3-gallate (EGCG), or catechins in green tea polyphenols, is best known as an antioxidant and antiinflammatory agent.³⁰ It, therefore, has been proposed to use in many clinic scenarios such as inhibiting cancer progression, preventing metabolic disorders, and modulating immune diseases. It is known to block a wide array of inflammatory stimuli through inhibiting intracellular NF-κB activation.³¹ Importantly, it has been reported that the association occurred between higher EGCG consumption and a decreased risk of fractures with improved bone health.32-34 The inhibitory effects of EGCG on osteoclastogenesis are relatively well established, while its action on osteoblast differentiation is still controversial, with both increasing and suppressing bone formation/osteoblast differentiation reported in the literature.³⁵⁻³⁸ In addition, there is even less information about the effect of EGCG on osteogenic differentiation of MSCs.35,39 Given the strong evidence of anti-inflammatory effect of EGCG and inhibition of bone formation by inflammatory cytokine such as TNFa, we designed and carried out the experiments to examine if EGCG would promote osteogenic differentiation in human MSCs isolated from bone marrow (BM-hMSCs) and to identify potential interplays between EGCG and TNF α in this setting. In the present study, our data support that EGCG has a protective effect on the osteogenic response inhibited by TNF α in MSCs, which is potentially mediated through an important downstream player of TNFα, namely NF-κB signaling.

Methods and materials

Isolation and culture of BM-hMSC

Bone marrow cells were placed in heparinized tubes prefilled with a-MEM and then washed twice with HBSS followed by Ficoll-Paque centrifugation (1800 \times g for 30 min at room temperature). Buffy coat was then carefully collected from the Ficoll-HBSS interface and washed again by HBSS. Viable cells were counted with a hemocytometer using Trypan blue and plated at a cell density of 50-100 cells/cm² in 175 cm² flasks or 150 mm dishes. After 24 h to remove, the adherent cells were cultured in CCM media at 37°C with 5% humidified CO₂. Vehicle controls, TNFa (1, 5, 10, and 20 ng/mL), or EGCG (5, 10, 20, and 40 μ M) were added in the media throughout the primary cultures until harvest for assays. For the blockade of NF-KB pathway, IKB-AA1 (super-repressor of NF-κB, cloned into a commercially available Rc/CMV expression vector) was transfected. The protocol was approved by the Committee on the Ethics of The Second Affiliated Hospital of Nantong University.

Osteogenic differentiation of BM-hMSC

BM-hMSCs were cultured and grown to reach complete confluence (24–72 h) and then induced with MSC osteogenic differentiation medium containing 10 mmol/L β -glycerol phosphate, 10⁻⁸ mol/L dexamethasone, and 50 mg/L ascorbic acid 2-phosphate.³⁹ Differentiation medium was changed every third day. Vehicle controls, TNF α (1, 5, 10, and 20 ng/mL), or EGCG (5, 10, 20, and 40 μ M) were added in the differentiation media throughout the differentiation processes until harvest for assays.

Characterization of BM-hMSC by flow cytometry analysis

Adherent MSCs were trypsinized and resuspend in PBS containing 5% fetal bovine serum. After washing, cells were counted and aliquoted into tubes for different staining by antibodies recognize various surface proteins, including CD29, CD45, CD105, and CD90. Isotype control antibody-stained cells were used to optimize photomultiplier tube and compensation in the analysis using BD-FACScan (Becton Dickinson, San Jose, CA). Flow cytometry data were analyzed with CellQuest software.

ALP activity assay

Cultured cells after eight days in osteogenic differentiation media were fixed in 4% paraformaldehyde for 15 min at room temperature and then washed twice with PBS followed by staining in freshly prepared 0.1% naphthol AS-MX phosphate, 56 mM 2-amino-2-methyl-1-3-propanediol, and 0.1% fast red violet LB salt. Quantitative analysis was determined by colorimetric assay of enzyme activity using the ALP kit (Sigma, St. Louis, MO, USA). Briefly, total protein lysates were extracted and then mixed with the freshly prepared colorimetric substrate para-nitrophenyl phosphate at 37°C for 30 min. The enzymatic reaction was stopped by adding NaOH (0.2 M). The optical density of the yellow product para-nitrophenol was determined by a spectrometer plate reader (Molecular Devices, CA, USA) at a wavelength of 405 nm. The ALP activity was normalized by protein amount and expressed as relative fold compared with vehicle control-treated samples.

Mineralization assays

The mineralization nodules formed 16 days after osteogenic induction of mesenchymal stem cells in dishes were determined by alizarin red-sulfate staining.³⁹ The washed cells were treated by 40 mM AR-S (pH 4.2) with gentle rocking for 30 min on room temperature. After three times of water rinse followed by a PBS washing step, the positively stained mineralization were examined by light microscopy.

Quantitative real-time polymerase chain reaction

Total RNA was isolated from cell cultures using the RNeasy kit (Qiagen, Duesseldorf, Germany) and was reverse transcribed to complementary cDNAs using Superscript II according to manufacturer's instructions (Invitrogen, NY, USA). Primers specific for *Runx2* and *Ostrix* were used. Duplicated polymerase chain reaction (PCR) reactions

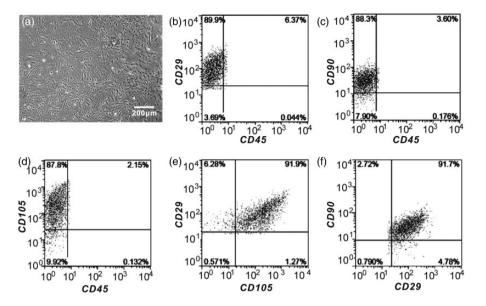


Figure 1 Cell viability of human BM-MSCs under different concentrations of TNF- α (1, 5, 10, and 20 ng/mL) and EGCG (5, 10, 20, and 40 μ M) treatment. Cell viability was measured by MTT assay and cell death was indicated by LDH release assay. Cell viability (a) and LDH release (b) of BM-MSCs normalized to that of control under TNF- α treatment for one, three, and five days, respectively. Cell viability (c) and LDH release (d) of BM-MSCs normalized to that of control under EGCG treatment for one, three, and five days, respectively. Cell viability (c) and LDH release (d) of BM-MSCs normalized to that of control under EGCG treatment for one, three, and five days, respectively. Cell viability (e) and LDH release (f) of BM-MSCs normalized to control under co-treatment of 20 ng/mL TNF- α and 5 μ M EGCG. All the experiments were independently repeated in triplicate. Data were presented by mean \pm SEM. **P* < 0.05 and #*P* < 0.01 vs. control

were carried out in ABI 7300 real-time PCR machine (Life Technologies, USA), using n=3 for each sample. SYBR Green dye-based detection method was used by using the SYBR Green PCR Master Mix assay (TaKaRa Biotechnology). A series of duplicate dilutions of cDNA from control samples were used to optimize the standard curve and validate the melting curves for each primer set.

Western blotting

For the protein analysis, cells were lysed in hypotonic buffer (10 mM HEPES, pH 7.9, 1.5 mM MgCl₂, 10 mM KCl, 0.5 mM DTT, and 0.2 mM PMSF) to separate cytoplasmic fractions from nuclear extracts. Both fractions were then measured and the same amounts of proteins were subjected to SDS-PAGE, followed by electric transfer into PVDF membrane. The antibodies used in western blot included anti-p65, anti-I κ B α , and anti-phosphorylated I κ B α (Cell Signaling). Anti-GAPDH and anti-HDAC1 (Cell Signaling) were used as loading controls for the cytoplasm and nuclear factions, respectively.

Statistics

All data in graphs are expressed as the means \pm SEM as indicated. The two-way ANOVA was used for Figure 1 and one-way ANOVA was used in Figure 2. The comparisons were determined by Bonferroni post test with **P* < 0.05 and #*P* < 0.01 (vs. the controls) considered as significant.

Results

Isolation and characterization of human bone marrow-derived MSCs

We used a conventional isolation protocol to obtain the mesenchymal stem cell derived from human bone

marrow (Figure 3(a)). The identity of the progenitor cells was also confirmed by the analysis of well-known surface markers, including positive for CD29, CD105, CD90, and negative for CD45 (Figure 3(b) to (f)).

The effects of TNF $\!\alpha$ and EGCG on cell viability of BM-hMSCs

In order to determine the effects of $TNF\alpha$ on MSCs, we tested a series of concentrations of TNFa (1, 5, 10, and 20 ng/mL) based on published reports and our own preliminary experiments. As shown by Figure 1(a) and (b), increasing concentrations of $TNF\alpha$ (particularly at 10 and 20 ng/mL) can significantly reduce cell viability and increase cell death in MSCs. These effects were found as early as one day at the highest concentration of $TNF\alpha$ (20 ng/mL) treatment and accumulated up to five days (more than 40% decrease in cell viability and around 80% increase in cell death). Similarly, we examined a series of doses of EGCG (5, 10, 20, and $40\,\mu\text{M}$) on BM-hMSCs by the same assays. Interestingly, we found an inverse dose response of EGCG, with only lower doses (5 and $10\,\mu$ M) increasing the cell viability and decreasing cell death of BM-hMSCs (Figure 1(c) and (d)). To further test the cross-talk between these two agents, we chose 20 ng/mL TNF α (the most significant damage in cell survival) and 5 µM EGCG (the most effective increase in cell viability). We found that at 5 µM EGCG concentration, the inhibitory effect of TNFa in BM-hMSCs was completely corrected in both cell viability (Figure 1(e)) and cell death assays (Figure 1(f)) during five-day treatment.

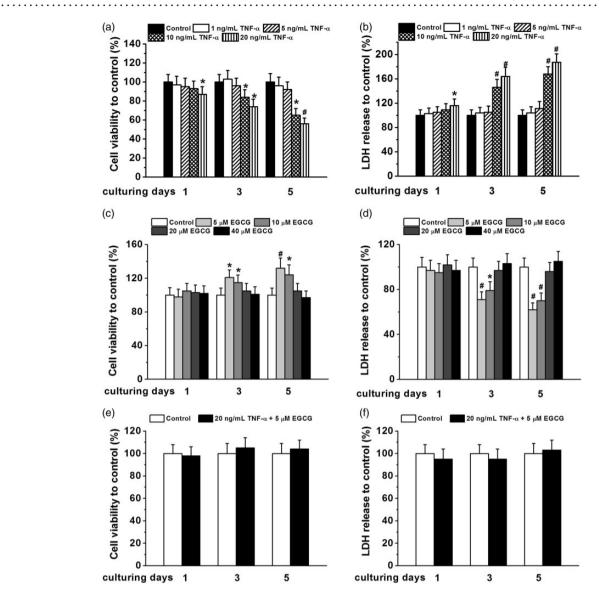


Figure 2 NF- κ B signaling pathway was involved in the protective effect of EGCG on the TNF- α -induced inhibition of osteogenic differentiation of human BM-MSCs. The changes of p65, I κ B α , and p-I κ B α protein bands in the cytoplasm (a) and p65 protein band in the nucleus (b) during the osteogenic differentiation at indicated time (day 0, 3, 7, and 16) in the cultures of human BM-MSC under 20 ng/mL TNF- α treatment. EGCG (5 μ M) was capable of inhibiting NF- κ B signaling pathway activated by 20 ng/mL TNF- α . (c and d) The p65, I κ B α , and p-I κ B α protein expressions in the cytoplasm in the cultures at day 16 during osteogenic differentiation in the experimental groups. (e and f) The p65 expression in the nucleus at day 16 during osteogenic differentiation in the experimental groups. (g and h) EGCG failed to rescue TNF- α -induced inhibition of osteogenic differentiation in BM-hMSCs when NF- κ B was blocked by its super-repressor I κ B-AA1, as indicated by no significant difference of the ALP activity, calcium content, Runx2, and Osx mRNA expression between non-EGCG and EGCG treated groups. All the experiments were independently repeated in triplicate. Data were presented by mean \pm SEM. #*P* < 0.01 vs. control

The effects of TNF α and EGCG on osteogenic differentiation of BM-hMSCs

Next, we sought to investigate the effects of TNF α and EGCG on osteogenic differentiation of BM-hMSCs. Similarly as the previous findings, both reagents showed dose-dependent effects on the osteoblast differentiation processes (Figure 4). TNF α (as low as 5 ng/mL) inhibited the ALP activity, mineralization ability, calcium content (Figure 4(a) to (c)), and gene expressions of bone formation markers (Runx2 and Osterix in Figure 4(d)) with the maximal effects at the highest dose 20 ng/mL. On the contrary, EGCG promoted osteogenic differentiation

(Figure 4(a) to (d)) only at lower doses (5 and $10 \,\mu$ M), with no effects at higher doses (20 and $40 \,\mu$ M). Finally, in the presence of a low-dose EGCG (5 μ M), TNF α could no longer block the BM-hMSC osteogenic differentiation, shown by ALP activity, mineralization nodule formation, and Runx2 and Osterix expressions (Figure 4(d) to (h)).

TNF α stimulates NF- κ B signaling in BM-hMSCs that is inhibited by EGCG

To better ascertain the importance of intracellular signaling originated from the classic inflammation mediator, we

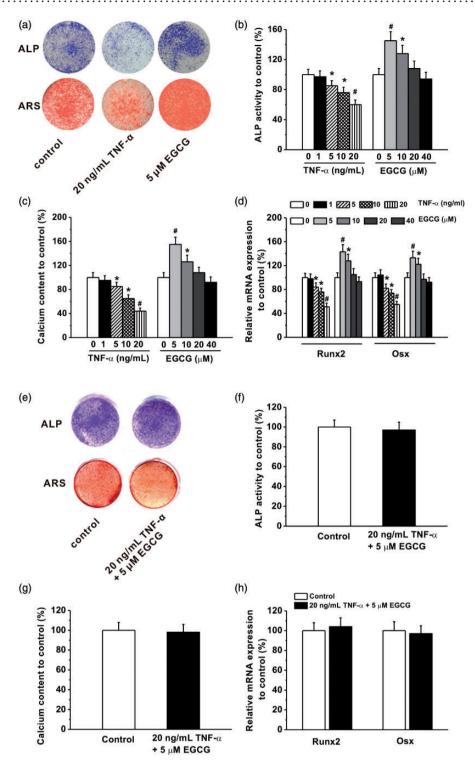


Figure 3 Phenotypic characterization of human BM-MSCs. (a) Representative image of human BM-MSCs after seeding for five days. (b to f) Flow cytometry identified BM-MSCs positive marker CD29, CD90, CD105, and negative marker CD45. (A color version of this figure is available in the online journal)

examined NF- κ B pathway in BM-hMSCs in response to TNF α treatment. During the cultures of human BM-MSC under 20 ng/mL TNF- α treatment from day 3 to day 16, we found an increased phosphorylation-associated degradation of I κ B α protein in the cells (Figure 2(a)) accompanied by a sustained nuclear translocation of p65 (Figure 2(b)) from the cytoplasm (Figure 2(a)). These data demonstrated

that TNF α activates classic NF- κ B signaling in bone marrow-derived MSCs. Importantly, EGCG blocked such activation by TNF α (again at a low dose of 5 μ M). Treating cells with EGCG, we found that the levels of I κ B α protein remained in the cytoplasm even in the presence of TNF α (20 ng/mL on day 16, Figure 2(c) and (d)), possibly due to an attenuation of I κ B α phosphorylation (Figure 2(c) and (d)).

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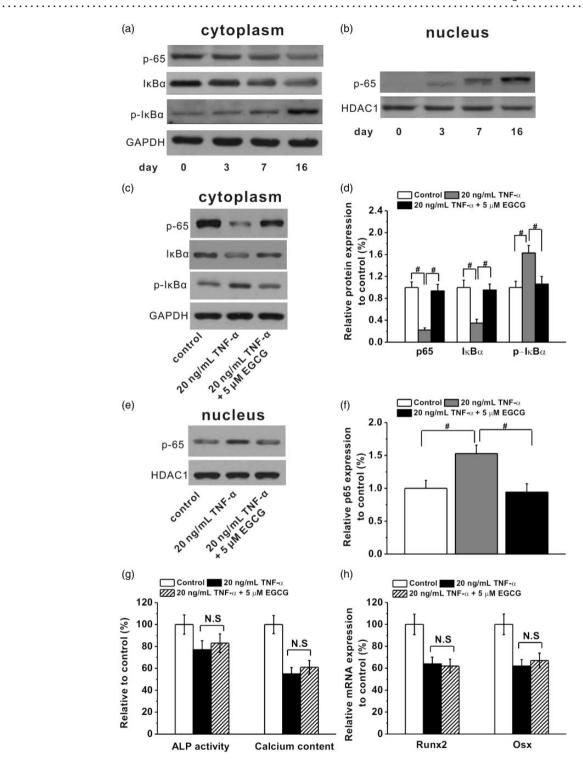


Figure 4 Effects of different concentrations of TNF- α (1, 5, 10, and 20 ng/mL) and EGCG (5, 10, 20, and 40 μ M) treatment on osteogenic differentiation of human BM-MSCs. (a) Representative photos for ALP staining in the cultures at day 8 and Alizarin red staining at day 16. Relative ALP activity (b) and relative calcium content (c) in the experimental groups, (d) Relative mRNA expression of *Runx2* and Osx in the experimental groups, quantified by RT-PCR analysis. (e) Representative photos for ALP staining in the cultures at day 16 after co-treatment of 20 ng/mL TNF- α and 5 μ M EGCG. Relative ALP activity (f), relative calcium content (g) and relative expression of *Runx2* and Osx mRNAs (h) in the cultures after co-treatment of 20 ng/mL TNF- α and 5 μ M EGCG. All the experiments were independently repeated in triplicate. Data were presented by mean \pm SEM. **P* < 0.05 and #*P* < 0.01 vs. control. (A color version of this figure is available in the online journal.)

As a result, p65 protein was still quenched in the cytoplasm (compared with TNF α treatment only, Figure 2(c) and (d)) after EGCG treatment, losing the ability to shuttle into the nucleus for transcription activation (Figure 2(e) and (f)).

Besides, when NF- κ B was blocked by NF- κ B super-repressor I κ B-AA1 in BM-hMSCs, EGCG treatment was not able to rescue the TNF α -induced inhibition of osteogenic differentiation any more, as there were no significant differences

in ALP activity, calcium content, Runx2, and Osx mRNA expressions between EGCG-treated group and EGCG-non-treated group (Figure 2(g) and (h)). This furthermore indicates the involvement of NF- κ B signaling pathway in the protective effects of EGCG against TNF α -induced impairments of osteogenesis.

Discussion

Based on prior evidences that inflammation can perturb osteoblast differentiation, we aimed to investigate the possible functions played by EGCG, a known antiinflammatory agent, in protecting bone formation inhibited by a pro-inflammatory cytokine TNF α . In particular, we focused on BM-hMSCs and found out that TNF α and EGCG have counter effects on the osteogenic differentiation process in MSCs, which might be mediated through a classic NF- κ B signaling pathway.

As others have suggested,^{26,27} we found an inhibitory effect of TNFa on the osteogenic potential of BM-MSCs, which is consistent with the notion that in the inflammatory milieu, not only bone resorption is activated but bone formation activity is also attenuated.⁷ Using highly pure progenitors of bone marrow cells (CD29⁺, CD90⁺, CD105⁺, and CD45⁻ populations) in primary cultures, our data have shown that TNF α is able to suppress ALP activity/mineralization and repress Runx2 and Osterix gene expressions in the differentiation of MSCs towards the osteoblastic lineages (Figure 4). TNF α can induce primary mesenchymal stem cell death (Figure 1), which could not be solely responsible for its inhibition on the bone differentiation since a low dose of TNFa (which didn't stimulate MSC death at 5 ng/mL) still significantly suppressed osteogenic differentiation (Figure 4, e.g. up to 20% decrease of Runx2 and Osterix gene expression). On the other hand, $TNF\alpha$ was first proposed as a crucial player in promoting postnatal bone repair, including intramembranous bone formation⁴⁰ and fracture repair,⁴¹ and later found to enhance expression of osteogenic factors like bone morphogenetic protein 2 (BMP-2) in mesenchymal cells.⁴² This discrepancy may be resulted from the differences in heterogeneous progenitor populations in MSC pools and the loss-of-function studies using TNFa knockout mice.^{40,41} In fact, TNFa-dependent osteoclast activation may be required to initiate bone remodeling that integrates bone resorption/formation cycles.^{7,21} Thus, the integrity of TNF α signaling might be indispensable for bone repair process, while the direct action of TNFa on osteoblast differentiation and/or maturation could be detrimental, as suggested by our data and many others.26,27

The recent explosion of technique advances in isolation and expansion of adult stem/progenitor cells from human bone marrow^{11,12,15} has contributed significantly in developing cell-based therapies for inflammatory bone diseases. In order to restore the three-dimensional collagen structure and facilitate the integration of the newly synthesized matrix with the resident tissues, MSCs that can differentiate into all bone cell types have become an excellent choice for bone regeneration approaches. In addition, the ease of genetically manipulated bone marrow-derived stem cells makes them attractive vehicles for gene therapy, if necessary.¹¹ Given the drastically negative effects of TNF α on the *in vitro* bone differentiation from bone marrow-hMSCs reported here, we hypothesize that a counter strategy is necessary in restraining the actions of TNF α and other pro-inflammatory cytokines.¹⁵

Based on this rationale and previous findings, EGCG could be a good candidate. First, as a dietary supplement, EGCG or green tea extracts have been known for their effective benefits on bone health. A number of human studies have reported a positive correlation between tea drinking and bone mineral density in osteoporosis patients or postmenopausal women and reduced risk of hip fractures in aged people.³²⁻³⁴ Second, EGCG has anti-inflammatory activity and antioxidative stress action.^{32,39,43} Both of which most likely can mitigate the adverse impacts of inflammation damages on bone regeneration. For instance, a recent study revealed that EGCG is able to suppress the lipopolysaccharide-induced inflammatory bone resorption in mice.⁴⁴ Also, our present study provides an *in vitro* analysis to support this acclaim in human mesenchymal stem cells. Finally, direct pre-clinical tests using green tea polyphenols have approved its ameliorating effects on the animal models of rheumatoid arthritis,45 cadmium toxicity-associated bone disorder,46 and estrogen deficiency-induced bone loss.⁴⁷ Altogether, our results are consistent with the hypothesis that EGCG and related green tea extracts could be useful in the treatment of inflammatory disease, or as supplementary approaches for other therapies such as using hormones (glucocorticoids, PTH), nutrition (vitamin D), or Wnt signaling antagonist (DKK1).^{3,7}

Compared with the well-characterized benefit of EGCG on bone metabolism, the underlying mechanism is unclear, and particularly confusing in terms of the direct action of EGCG on osteoblast differentiation. EGCG was first shown to enhance bone formation activity in a osteosarcoma Saos-2 cell line,³⁶ increase osteogenesis in a murine bone marrow mesenchymal stem cell line D1,35 and upregulate Runx2mediated differentiation of bone cells.⁴⁸ However, conflicting data exist. In an osteoblastic cell line MC3T3-L1, EGCG appeared to suppress the ALP activity and osteoblastic differentiation,³⁷ which was compatible with another prior report in which an ectopic bone formation model induced by BMP was shown to be repressed by EGCG.⁴⁹ Our data support the positive effects of EGCG on osteoblastic differentiation in general and surprisingly have identified an inverse dose-dependent action on MSC cell survival and osteogenic differentiation (Figures 1, 2, and 4). The finding that the increasing doses of EGCG diminished its otherwise beneficial effects on the bone formation suggested that cells may respond differently to the various levels of EGCG, even in the opposite directions, which may explain the contradictory results obtained previously. These interesting findings are also consistent with a recent study on alveolar bone cells in which EGCG at a low concentration increases, whereas at higher concentrations decreases osteogenic differentiation.38

The molecular basis of EGCG protecting TNF α -induced bone formation repression can be complicated, and our data pointed to an important role played by NF- κ B pathway.

Since its discovery, NF-KB signaling has been extensively investigated and known as one of the best characterized intracellular pathway in inflammation.²⁹ We have found a dramatic stimulation of classic NF-kB signaling during long-term MSCs culturing in the presence of TNFa (Figure 2), which included enhanced IkBa phosphorylation accompanied by decreased IkBa protein (presumably due to proteasome degradation of IkBa dependent on phosphorylation²⁹). Consequently, the nuclear translocation of p65 was released from the lack of IkBa sequestration (Figure 2), supposedly inducing transcription activation/ repression of its target genes. Interestingly, a low dose of EGCG (5 μ M) completely abolished the NF- κ B signaling activated by TNFa, which is consistent with previous studies on osteoclast cells, or other cell types.^{35,45} These data suggest that EGCG reverses the inhibitory effects of TNFa, at the least in part, through preventing the activation of NF-kB signaling in BM-hMSCs during osteogenic differentiation. In addition, other mechanisms could involve, such as a matrix metalloproteinase (MT1-MMP) in mesenchymal stromal cells.⁵⁰ Importantly, oxidative stress can induce inflammatory responses through NF-kB-dependent or independent pathways, which is also a widely recognized target of EGCG due to its ability to capture and quench reactive oxygen species.^{30-32,39} The potential antioxidant functions of EGCG in suppressing the inhibition effects on mesenchymal stem cells by TNFa require further investigation in the future.

Conclusion

In conclusion, our studies have shown that EGCG can prevent the inhibition by TNF α on the survival and osteogenic differentiation of human BM-MSCs. These effects are only found at low concentrations of EGCG, potentially through suppressing NF- κ B signaling stimulation by TNF α . Therefore, besides the efforts in reducing bone resorption in the inflammation-associated bone diseases, strategies such as using green tea extracts to promote bone formation while preventing the damaging of bone repair by inflammatory insult may directly benefit these bone diseases.

Author contributions: All authors participated in the design, interpretation of the studies, analysis of the data, and review of the manuscript; WL, JBF, DWX, and JZ conducted the experiments; WL and ZMC wrote the manuscript; WL and JBF contributed equally to this work.

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DECLARATION OF CONFLICTING INTERESTS

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

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