Human Full-Length Osteoprotegerin Induces the Proliferation of Rodent Vascular Smooth Muscle Cells both in vitro and in vivo

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Abstract

Background/Aims: Since elevated plasma levels of osteoprotegerin (OPG) represent a risk factor for death and heart failure in patients affected by diabetes mellitus and coronary artery disease, this study aimed to elucidate potential roles of OPG in the pathogenesis of atherosclerosis. Methods and Results: Recombinant human full-length OPG, used at concentrations comparable to the elevated levels found in the serum of diabetic patients, significantly increased the proliferation rate of rodent vascular smooth muscle cells (VSMC). To mimic the moderate chronic elevation of OPG observed in diabetic patients, low doses (1 μg/mouse) of full-length human OPG were injected intraperitoneally every 3 weeks in diabetic apolipoprotein E (apoE)-null mice. The group of animals treated for 12 weeks with recombinant OPG showed a small increase in the total aortic plaque area at necropsy in comparison to vehicle-treated animals. Importantly, while no differences in the amount of interstitial collagen or the degree of macrophage infiltration were observed between OPG-treated and vehicle-treated apoE-null diabetic animals, a significant increase in the number of α-actin-positive smooth muscle cells was observed in the plaques of OPG-treated mice. Conclusions: Our data suggest that OPG promotes VSMC proliferation and might be directly involved in pathogenetic aspects of atherosclerosis.

Key Words

Osteoprotegerin · Vascular smooth muscle cells · Apolipoprotein E-null mice · Cell proliferation · Atherosclerosis

Introduction

Osteoprotegerin (OPG), a soluble member of the tumor necrosis factor (TNF) receptor superfamily, was originally characterized for its ability to suppress osteoclastogenesis by binding to and inhibiting receptor activator of NF-κB ligand (RANKL), a member of the TNF superfamily of cytokines [1, 2]. Subsequently, it has been shown that OPG can also interact with another member of the TNF superfamily, TNF-related apoptosis-inducing ligand (TRAIL), which shares approximately 25% amino acid homology with RANKL [3] and participates together with RANKL and OPG in the control of osteoclastogenesis [4–6]. Interestingly, a recent study has demonstrated that the affinity of OPG for TRAIL is slightly lower than but comparable to that for RANKL [7].
Several studies have clearly demonstrated that the serum levels of OPG are elevated in diabetic and nondiabetic patients affected by cardiovascular diseases, and increased levels of OPG represent a risk factor for cardiovascular mortality, especially in diabetic patients [8–21]. However, in spite of the reported findings, the physiological role of elevated serum levels of OPG in vascular biology is not well understood. In this context, a possible pathogenetic link between elevated levels of OPG and inflammation has been suggested by recent in vitro studies of our group [22] and that of Mangan et al. [23] demonstrating that exposure to recombinant OPG promotes leukocyte adhesion to endothelial cells. These findings are particularly noteworthy since atherosclerosis, which constitutes the single most important contributor to the growing burden of cardiovascular disease, is regarded as a form of chronic low-grade inflammatory process which can ultimately lead to the development of complex lesions, or plaques, that protrude into the arterial lumen [24]. Moreover, it has recently been demonstrated that OPG might be involved in the pathogenesis of pulmonary hypertension by promoting the growth of human vascular smooth muscle cells (VSMC) obtained from pulmonary artery [25].

The aim of this study was to investigate whether OPG is involved in pathogenetic aspects of atherosclerosis. For this purpose, the effect of increasing concentrations of human recombinant OPG, comparable to those found in the plasma of diabetic patients [14], was investigated in rodent VSMC. In addition, recombinant full-length OPG was administered in vivo in an experimental model of atherosclerosis represented by apolipoprotein E (apoE)-null diabetic mice. Upon induction of diabetes mellitus by streptozotocin injection, atherosclerotic lesions developed over time in these animals and resemble in appearance and distribution those observed in humans [26].

**Materials and Methods**

**Cell Cultures**

Primary rat aortic VSMC were purified as previously described and used between passages 3 and 6 [27]. Expression of lineage-specific markers was monitored by immunofluorescence analysis with anti-α-smooth muscle actin (α-SMA) antibody (Dako, Copenhagen, Denmark). For cell cycle analysis, rodent VSMC were seeded at a low cell density and made quiescent by using serum-reduced (0.1% FBS) medium before incubation with 5-bromodeoxyuridine (BrdU; Sigma, St. Louis, Mo., USA) at 37 °C for 1 h. Anti-BrdU antibody was bound to BrdU incorporated into neosynthesized DNA, and the complex was detected by fluorescein isothiocyanate-conjugated secondary antibody. Cells were stained with propidium iodide and analyzed by flow cytometry. To avoid nonspecific fluorescence from dead cells, live cells were gated tightly using forward and side scatter, as described elsewhere [28].

**Animals and Experimental Protocol**

Animal care and treatment conformed to institutional guidelines in compliance with national and international laws and policies (EEC Council Directive 86/609, OJL 358, 12 December 1987). Six-week-old apoE-null mice (n = 20) were rendered diabetic by 5 daily intraperitoneal injections of streptozotocin (Sigma Chemical Co., St. Louis, Mo., USA) at a dose of 55 mg/kg body weight. The animals were maintained on regular mouse chow. Diabetic animals were further randomized to receive an intraperitoneal injection of either human recombinant OPG (R&D Systems, Minneapolis, Minn., USA; 1 µg/mouse in a total of 200 µl of HEPES-buffered saline) or an equivalent volume of vehicle, every 3 weeks. After 3 months, the animals were anesthetized by an intraperitoneal injection of pentobarbital sodium (60 mg/kg body weight) and sacrificed for blood tests and histological examination. In each group of animals, serum glucose, glycosylated hemoglobin (HbA1c), systolic blood pressure, triglycerides and total and high-density lipoprotein (HDL) cholesterol were determined as previously described [26, 29]. Human OPG levels were measured in mouse serum samples using a sandwich-type enzyme-linked immunosorbent assay (ELISA) kit (Alexis Biochemicals, Lausen, Switzerland), which does not cross-react with mouse OPG, according to the manufacturer’s instructions. Mouse RANKL and OPG serum levels were measured in mouse serum samples using ELISA kits from R&D Systems. The results were read at an optical density of 450 nm using an Anthos 2010 ELISA reader (Anthos Labtec Instruments GmbH, Wals/Salzburg, Austria). Measurements were performed in duplicate.

**Evaluation of Atherosclerotic Plaques**

To analyze the atherosclerotic lesions, 2 complementary approaches were used: an en face whole approach and histological section evaluations. The en face approach was used to obtain information about the distribution and extent of atherosclerosis in the aorta, whereas microscopic histological analysis was used to evaluate the lesion composition and complexity, as previously described [26].

In brief, for the en face approach the entire aorta was opened longitudinally and stained with Sudan IV-Herxheimer’s solution (Sigma), and lesion area measurements were performed by calculating the proportion of aortic intimal surface area occupied by the red stain in the arch, descending thoracic and abdominal aorta, with the use of a video-based image analysis program (MCID, Imaging Research, St. Catharines, Ont., Canada). All aortic segments were then embedded in paraffin, and 4-µm-thick cross-sectional serial sections were obtained. For quantification of collagen content, aortic cross-sections were stained with either Mason trichrome or Sirius red, as previously described [26].

**Immunohistochemistry**

Aortic lesions were characterized for cell composition by immunohistochemical analysis of cross-sectional sections. For this purpose, after neutralization of endogenous peroxidase, paraffin-embedded sections of aorta were incubated with the following primary antibodies: α-SMA (smooth muscle cell marker) and...
proliferating cell nuclear antigen (PCNA; both from Dako; diluted 1:200). Biotinylated immunoglobulins (diluted 1:200) were then applied as a secondary antibody, followed by horseradish peroxidase-conjugated streptavidin (Dako; diluted 1:625). The staining was visualized by reaction with 3,3′-diaminobenzidine tetrahydrochloride (Sigma). Macrophage detection was performed using the primary antibody for F4/80 (Serotec, Oxford, UK; diluted 1:200), followed by secondary antirat immunoglobulins (Vector Laboratories, Burlingame, Calif., USA; diluted 1:200) and a catalyzed signal amplification mouse amplification kit (Dako), following the manufacturer’s instructions. Syndecan-1 detection was performed using monoclonal rat antiamouse CD138 (BD Pharmingen, San Jose, Calif., USA; diluted 1:500), followed by secondary biotinylated rabbit antirat antibody (Vector Laboratories; diluted 1:200) and an ABC kit (Vector Laboratories), following the manufacturer’s instructions.

After counterstaining with hematoxylin, all the sections were examined by light microscopy and digitized using a high-resolution camera. Atherosclerotic lesions were manually traced on the computer, taking care to exclude normal-appearing media and to include only the intimal/subintimal atherosclerotic lesions, and quantification of collagen staining and of the specific immunostaining within the plaques was assessed using the MCID image analysis system. All immunohistochemical quantifications were performed by 2 independent blinded observers and are shown as the percentage of stained areas or of positive cells.

Real-Time Quantitative Reverse Transcription-Polymerase Chain Reaction

Three micrograms of total RNA extracted from each aorta were used to synthesize cDNA with the Superscript First Strand synthesis system for reverse transcription-polymerase chain reaction (RT-PCR; Gibco BRL, Grand Island, N.Y., USA). Angiopoietin-2 and vascular cell adhesion molecule-1 (VCAM-1) gene expression was analyzed by real-time quantitative RT-PCR using the Taqman system (ABI Prism 7900 HT, Perkin-Elmer). Gene expression of the target sequences was normalized in relation to the expression of an endogenous control, 18S ribosomal RNA (18S rRNA Taqman Control Reagent kit, ABI Prism 7900 HT, Perkin-Elmer). Primers and Taqman probes for angiopoietin 2 and VCAM-1 and the endogenous reference 18S rRNA were designed with the help of Primer Express (ABI Prism 7900 HT, Perkin-Elmer, PE Biosystems). Expression of the angiopoietin 2 cDNA, the forward primer was 5′-GTCCAACCTACAGATTCACCTTACAG-3′ and the reverse primer was 5′-TTGTCCGAATCTTTGTGCTAA-3′. The probe specific to angiopoietin 2 was FAM-5′-CAGCCAACAGGAAAG-3′-MGB. For the VCAM-1 cDNA, the forward primer was 5′-AAGTCTGTGATGTGGCCTACA-3′ and the reverse primer was 5′-TCAGTCTTTAGGTACCTTGATATGC-3′. The probe specific to VCAM-1 was FAM-5′-CCGCATCCTGCACAGC-3′-TAMRA. Each sample was tested in triplicate. Results were expressed relative to control aorta values.

Statistical Analysis

The mean, median, minimum and maximum values were calculated for each group of data. Box plots were used to show the medians and 25th–75th percentiles. Data were analyzed by ANOVA and the Mann-Whitney rank sum test. Comparison of group means was performed by the Bonferroni method. Correlation coefficients were calculated by Spearman’s method. Statistical significance was defined as p < 0.05.

Results

Recombinant Full-Length OPG Promotes Proliferation of Primary Rodent VSMC

Since it has been previously demonstrated that OPG levels are significantly elevated in the plasma or serum of diabetic patients [14–21], in the first group of experiments we investigated whether exposure to recombinant full-length human OPG had any effect on cell cycle progression of rodent VSMC, analyzed by BrdU incorporation (fig. 1). As shown in figure 1, full-length human OPG at concentrations similar to those found in human serum significantly (p < 0.05) promoted the proliferation of VSMC [14].

In vivo Injection of Low Concentrations of Full-Length Recombinant Human OPG Induces a Small but Significant Increase in Atherosclerotic Plaques in apoE-Null Diabetic Mice

After the induction of diabetes mellitus (3 weeks after streptozotocin injection), apoE-null mice were randomized to be injected intraperitoneally either with recombinant human OPG or vehicle. Using an ELISA kit specific for human OPG, the presence of human OPG was detectable, although at low levels, in the serum of OPG-injected animals 24 h after injection but not in the serum of animals injected with vehicle (fig. 2). Levels of human OPG similar to those found after 24 h were detectable in animal sera also 1 week after intraperitoneal injection (data not shown). It should be noted that repeated intraperitoneal injections of human recombinant OPG (every 3 weeks for 3 months) were well tolerated by the apoE-null mice. After 3 months, the diabetic animals were analyzed for various parameters and then sacrificed for pathological examinations. The body weight and systolic blood pressure as well as the levels of blood glucose, HbA1c, total cholesterol, HDL cholesterol and triglycerides were not significantly different in diabetic animals injected or not with human recombinant OPG (table 1). In addition, to ascertain whether the injection of human recombinant OPG might modulate the levels of endogenous mouse OPG and RANKL, the levels of these molecules were quantitatively evaluated by ELISA at the end of the experimental period. As shown in table 1, no significant
differences in serum levels of OPG and RANKL were observed between mice injected with recombinant human OPG or vehicle. Unfortunately, it was not possible to measure the levels of TRAIL, the second ligand for OPG [6, 7], due to the lack of commercially available ELISA kits for mouse TRAIL.

En face dissection (fig. 3a) and cross-sectional serial sections (fig. 3b) showed the presence of plaque areas, characterized by asymmetrically thickened intima, in the entire aorta of apoE-null diabetic mice (fig. 3a, b). Of interest, OPG-treated animals exhibited a significantly (p < 0.05) increased total plaque area compared to the mice injected with vehicle (fig. 3a). Such an increase could not be ascribed to an aspecific, allergic or toxic effect consequent to the injection of a human protein in the apoE-null animals since the administration of recombinant human TRAIL induced the opposite effect, decreasing the total plaque area [29].

**The in vivo Injection of Recombinant Human OPG Modulates the Histological Composition of Atherosclerotic Plaques in apoE-Null Diabetic Mice**

We next investigated whether, besides increasing the extent of atherosclerotic lesions, OPG treatment also affected the histological composition of the plaques. For this purpose, cross-sectional serial sections of the aortic
Atherosclerotic lesions of apoE-null diabetic mice were quantitatively analyzed by immunohistochemistry for the presence of collagen and the cell composition of the plaques (fig. 4, 5). Of note, no significant differences in the collagen content or in the percentage of infiltrating macrophages were observed between OPG- and vehicle-injected mice (fig. 4a, b). Similarly, the levels of syndecan-1, which has been proposed to act as a ligand for OPG in multiple myeloma [30], were not changed in the aortas of OPG-injected mice with respect to vehicle-treated animals (data not shown).

On the other hand, the percentage of VSMC (α-SMA-positive cells) was significantly (p < 0.05) increased in the aortic lesions of OPG-injected apoE-null animals.
compared to diabetic animals injected with vehicle (fig. 5a, b). To elucidate whether the increased content of VSMC in the plaques of OPG-treated animals was accompanied by an increase in cell proliferation, we analyzed the percentage of PCNA-positive cells, which accurately reflects the degree of cell proliferation, within the plaques. As shown in figure 5a and b, OPG administration induced a significant (p < 0.01) increase in PCNA-positive cells. Interestingly, the percentage of PCNA-positive cells evaluated in all animals showed a significant positive correlation with the percentage of the α-SMA-positive area evaluated in the plaques of each mouse (R = 0.76, p < 0.05), suggesting that VSMC were the major cell type involved in cell proliferation within the plaques.

**Treatment with Recombinant Human OPG Increases the mRNA Expression of Angiopoietin 2 in the Aorta of apoE-Null Animals**

Taking into consideration the fact that a recent in vitro study demonstrated that angiopoietin 2 plays a key role in promoting inflammation of the vessel wall [31], we next investigated whether the in vivo administration of human OPG was able to modulate angiopoietin 2 expression in the aortic wall of apoE-null mice. The constitutive steady-state mRNA levels of angiopoietin 2 were quantitatively evaluated by real-time RT-PCR in RNA extracted from aortic samples of OPG- and vehicle-treated mice (fig. 6). The mRNA levels of angiopoietin 2 were significantly (p < 0.05) higher in the aortas of OPG-treated animals than in the aortas obtained from vehicle-treated
animals (fig. 6). On the other hand, the inflammatory cell adhesion molecule VCAM-1 [32] did not show significant modifications in OPG-treated versus vehicle-treated mice (fig. 6).

**Discussion**

Several studies have demonstrated that increased amounts of serum OPG correlate with increased cardiovascular risk in both diabetic and nondiabetic patients [8–21]. Possible interpretations of this positive relationship include OPG playing an active role in disease progression, or OPG serving as a compensatory response to minimize disease progression, or OPG representing a noncompensatory (neutral) response to disease. The interpretation that OPG serum elevation represents a compensatory mechanism mainly relies on studies performed on OPG knockout mice showing a protective role of OPG against vascular calcifications [33–35]. However, in each of these model systems, the prevention of vascular calcification by OPG was associated with strong suppression of bone turnover. Moreover, both recombinant OPG [34] and bisphosphonate [36] treatments of rats suppressed vascular calcification in a warfarin model of vascular calcification, in association with suppression of bone turn-

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**Fig. 5.** Recombinant human OPG increased the content of VSMC and their proliferative activity in the atherosclerotic plaques of apoE-null diabetic mice. Serial sections from aortas of apoE-null diabetic mice injected intraperitoneally with either vehicle (n = 10) or recombinant OPG (n = 10) were analyzed to quantitatively evaluate the presence/percentage of VSMC (using α-SMA marker) and the content of PCNA-positive cells in the aortic intima. Staining results, representatively shown in a (original magnification ×20), were quantitatively determined in each mouse with the MCID video-based image analysis program and expressed as the percentage of stained area or of positive cells. In b, horizontal bars are medians, upper and lower edges of the boxes are 75th and 25th percentiles and vertical lines are 10th and 90th percentiles.
in the vascular microenvironment. In spite of this cautionary consideration, it is noteworthy that human OPG could be detected in the general circulation of mice injected intraperitoneally with human OPG for at least 1 week after injection. We were able to demonstrate that repeated injections in apoE-null mice of relatively low concentrations of human OPG resulted in a significant increase in the total plaque area. Taking into account the fact that we previously demonstrated in a series of in vitro and in vivo experiments that soluble recombinant TRAIL shows anti-inflammatory [40–42] and antiatherosclerotic [29] activity, one mechanism by which OPG might accelerate the development of atherosclerotic lesions is through the inhibition of soluble circulating TRAIL. Consistent with this interpretation, it has previously been shown that soluble TRAIL levels have a tendency to decrease in the sera of patients affected by cardiovascular diseases [43, 44]. Moreover, the demonstration that OPG significantly increases the mRNA expression of angiopoietin 2 at the aortic level is in line with the results of a recent in vitro study [23]. Since angiopoietin 2 has been shown to play a key role in the early steps of atherosclerosis [31], the ability of OPG to increase the expression of angiopoietin 2 might represent an important additional pathogenetic mechanism to explain the ability of OPG to promote atherosclerosis. On the other hand, we also noticed a slight trend for an increase in systolic blood pressure in the OPG-treated mice (table 1), which might account for the changes observed in atherosclerotic plaques, but we were unable to perform hemodynamic assessments that may have provided additional information.

It is also noteworthy that the plaques in OPG-treated mice showed a significant increase in the number of VSMC. In line with our finding that OPG increases the number of proliferating VSMC in aortic plaques of apoE\(^{-/-}\) knockout mice, a study performed on OPG/apoE double (OPG\(^{-/-}\)apoE\(^{-/-}\)) knockout mice [35] demonstrated that the atherosclerotic lesions observed in these mice were characterized by a severe reduction in the number of smooth muscle cells, which rendered these plaques unstable. Moreover, in keeping with our finding that recombinant full-length OPG promotes rodent VSMC proliferation in vitro, a recent study has demonstrated that recombinant OPG is able to significantly increase the proliferation of cultured human pulmonary VSMC [25].

Our present findings should also be considered in the light of a recent study by Morony et al. [45], who demonstrated that subcutaneous injection of pharmacological concentrations of human recombinant OPG (OPG-Fc) decreased the degree of atherosclerotic calcified lesions
in diet-fed atherogenic ldlr−/− mice, without affecting the total burden of atherosclerotic lesions. The most likely explanation to reconcile the findings of Morony et al. [45] with our present findings is that Morony et al. [45] used human OPG-Fc rather than native human full-length OPG. In OPG-Fc, the signal peptide, heparin-binding domain and death domain homologous regions are removed [46], and we and others have previously shown that the heparin-binding domain of OPG is essential to mediate important proinflammatory effects of OPG [7, 22, 23]. Moreover, it has been shown that surface proteoglycans, which mediate interactions with the heparin-binding domain of OPG [7, 22], play a key role in promoting VSMC migration and proliferation [47].

Besides contributing to vessel wall inflammation and lipoprotein retention, VSMC play a key role in the formation of the fibrous cap that provides stability to plaques, preserving them from rupture, and therefore the benefit/risk of therapeutic inhibition of VSMC proliferation in atherosclerosis is unclear [24]. In this context, the effect of OPG on plaque formation seems to be twofold; on the one hand, it increases the extent of the total plaque area, but on the other hand, it may contribute to plaque stability by promoting the proliferation of VSMC. Anyhow, although one must be cautious in extrapolating these experimental data to the clinical context, an important implication of our data is that denosumab, a fully human monoclonal antibody which blocks osteoclastogenesis by inhibiting RANKL [46], might be preferable to recombinant OPG for the treatment of pathologies associated with high bone turnover, such as osteoporosis.

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References


OPG Promotes VSMC Proliferation

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261


