Short Communication

Gene Expression of Osteoprotegerin Ligand, Osteoprotegerin, and Receptor Activator of NF-κB in Giant Cell Tumor of Bone

Possible Involvement in Tumor Cell-Induced Osteoclast-Like Cell Formation

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Giant cell tumor of bone (GCT) is a rare primary osteolytic tumor of bone that is characterized by massive tissue destruction at the epiphysis of long bones. There is no evidence that tumor cells themselves are capable of bone destruction; instead, it appears that the tumor cells of GCT act by promoting osteoclastogenesis and, as a consequence, osteoclastic bone resorption. However, the mechanism by which this is achieved is not understood. Here we attempted to determine whether osteoprotegerin ligand (OPGL), the factor that is necessary and essential for osteoclastogenesis, is involved in tumor cell-recruited osteoclast-like giant cell formation in GCT. Using fluorescence in situ hybridization, we sought to determine mRNA expression of OPGL, its receptor RANK, and its decoy receptor OPG in three major cell types of GCT. We demonstrated that OPG mRNA was expressed in all three cell types of GCT, OPGL transcripts were mainly detected in spindle-shaped stromal-like tumor cells, whereas RANK was expressed only in macrophage-like mononuclear cells and multinuclear osteoclast-like giant cells. By semiquantitative RT-PCR, we also showed that the level of OPGL mRNA in GCT is much higher than that in normal bone and osteogenic osteosarcoma. In contrast, a similar level of OPG transcripts was detected in these three kinds of tissues, and RANK mRNA was detectable only in GCT tissues. We have further examined the regulation of gene expression of OPGL and OPG in tumor cells in response to osteotropic hormones. Administration of 1,25(OH)2D3 and dexamethasone resulted in maximum up-regulation of OPGL level and down-regulation of OPG level in cultured GCT stromal-like tumor cells and the mouse bone marrow-derived ST-2 stromal cell line. Furthermore, we have shown that tumor cells of GCT induce differentiation of RANK-expressing myeloid RAW264.7 cells into osteoclast-like cells in the presence of 1,25(OH)2D3 and dexamethasone. Our findings suggest that OPGL is involved in the tumor cell-induced osteoclast-like cell formation in GCT. The ratio of OPGL/OPG by tumor cells may contribute to the degree of osteoclastogenesis and bone resorption.

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Two novel tumor necrosis factor (TNF) superfamilies, osteoprotegerin (OPG) and osteoprotegerin ligand (OPGL), have recently been identified as members of a ligand-receptor system that directly regulates osteoclast differentiation and bone resorption.1–4 OPGL, also known as osteoclast differentiation factor (ODF),3 TNF-related activation-induced cytokine (TRANCE),3,5,6 and receptor activator of NF-κB ligand (RANKL)5 belong to the membrane-associated TNF-ligand family. It has been shown, using an in vitro culture system, that OPGL can both induce osteoclastogenesis and activate mature osteoclasts.3,4 The expression of OPGL in osteoblast/stromal...
cells parallels the formation of osteoclasts in cocultures with bone marrow or spleen cell populations. The recombinant OPGL can replace the requirements for stroma cells in the in vitro model of osteoclastogenesis.4 Mice with a disrupted opgl gene show severe osteopenosis and a defect in tooth eruption and completely lack osteoclasts as a result of an inability of osteoblasts to support osteoclastogenesis.7 It has been assumed that OPGL acts as an osteoclastogenesis-inducing factor linked to interaction between stromal cells and osteoclast progenitors. The cell surface receptor that interacts with OPGL has recently been shown to be the ligand for the TNFR-related protein receptor activator of NF-κB (RANK).8 Transgenic mice expressing a soluble RANK-Fc fusion protein display osteopenosis, a defect of osteoclast activity, whereas polyclonal antibody against the RANK extracellular domain promotes osteoclastogenesis in bone marrow cultures.8 On the other hand, the decoy receptor of OPGL, OPG, has been shown to neutralize and interrupt stromal cell-derived OPGL signals, resulting in the reduction of osteoclastogenesis.1,2,13 OPG, also known as osteoclastogenesis inhibitory factor (OCIF),2 is a soluble member of the TNF receptor family. OPG inhibits not only formation of osteoclast-like cells in murine cultures in vitro2 but also bone resorption in vitro and in vivo.1,2 OPG knock-out mice exhibited severe osteopenia due to accelerated bone resorption.1 In short, it is conceivable that OPGL and OPG are key extracellular regulators of osteoclastogenesis and bone resorption. RANK is receptor necessary for the activation of OPGL.

Giant cell tumor of bone (GCT), a rare primary osteolytic tumor of bone, is characterized by massive bone destruction at the epiphysis of long bones.9 Previous studies have shown that the spindle-shaped stromal-like mononuclear cells of GCT are the most likely candidate cells for the tumor’s neoplastic component.9-12 However, there is no evidence that tumor cells themselves are capable of bone resorption.13 Instead, tumor cells of GCT act by recruiting multinucleated osteoclast-like giant cells and hence promoting tumor-induced osteolysis.9-12 Moreover, tumor cells of GCT are related to the induction of osteoclastogenesis and bone resorption.16-18 It is noteworthy that although individual cytokines play a role in the recruitment of osteoclasts and in osteoclastic bone resorption, none of them are capable of directly inducing the macrophage-like cells to form osteoclasts. In fact, cells from osteogenic tumors such as osteosarcomas can express a similar cytokine profile,19-21 as do cells from GCT, but induction of osteoclast formation is observed to a lesser degree. Thus the mechanism by which tumor cells of GCT generate osteoclast-like giant cells from recruited macrophage-like cells is unclear.

To determine whether OPGL is involved in the interaction of stromal-like tumor cells and macrophage-like mononuclear cells, which results in the generation of osteoclast-like giant cells in GCT, we have investigated the expression of OPGL, OPG, and RANK in all three cell types of GCT at the mRNA level and examined the regulation of gene expression in response to osteotropic hormones. A coculture system of stromal-like tumor cells and murine osteoclast precursor cell line RAW264.7 cells6 has been used to evaluate the ability of tumor cells to further induce osteoclastogenesis.

Materials and Methods

Materials

Human GCT from five different cases and normal cancellous bone containing no bone marrow cells were collected fresh from patients after operations at Sir Charles Gairdner Hospital (Nedlands, WA, Australia). The human osteoblast-like osteosarcoma cell line U2OS was purchased from the American Type Culture Collection (Rockville, MD). The mouse stromal cell line ST-2 was provided by Prof. G. C. Nicholson at the Geelong Hospital (Geelong, VIC, Australia). The mouse myeloid cell line RAW264.7 was provided by Dr. Ian Cassady at the University of Queensland (Queensland, Australia). Dulbecco’s minimum essential medium (MEM), α-MEM, and fetal bovine serum (FBS) were purchased from TRACE (Sydney, Australia). 1α,25-Dihydroxyvitamin D₃ (1,25(OH)₂D₃) was obtained from Calbiochem-Novabiochem (Alexandria, Australia); dexamethasone was from Sigma (St. Louis, MO); recombinant human RANKL (sRANKL) was from Pepro Tech (NJ, USA); RETROscript was from Ambion (Austin, TX); the DIG RNA Labeling kit and the Fluorescent Antibody Enhancer Set were from Boehringer Mannheim (Sydney, Australia); the diagnostics tartrate resistant acid phosphatase (TRAP) kit was from Sigma; rabbit anti-rat/mouse calcinonin receptor (CTR) polyclonal antibody was provided by Dr. P. M. Sexton at the Department of Pharmacology (University of Melbourne, Victoria, Australia); Texas Red-X-conjugated swine anti-rabbit Ig secondary antibody was purchased from Molecular Probes (Eugene, OR). All other chemicals used were of the highest grade available.

Culture of Giant Cell Tumor of Bone

As described previously,4 tumor tissues were freshly chopped in DMEM containing 10% FBS, 100 U/ml penicillin, and 100 μg/ml streptomycin. The resultant cell suspension together with small pieces of tissue was transferred to 25-cm² flasks for culture at 37°C in a humidified atmosphere of 5% CO₂ and 95% air. The pieces of tissue contained many multinuclear giant cells and mononuclear cells that had migrated out of tissue fragments across the culture flask surface and contributed to the cell population in culture. Half of the culture medium was changed every 3 days with fresh DMEM containing 10% FBS, 100 U/ml penicillin, and 100 μg/ml streptomyc-
cin. On reaching confluence, primary cultures were subcultured, and each passaged culture was stored in liquid nitrogen. To examine the gene expression of OPGL and OPG in tumor cells of GCT, GCT stromal-like tumor cells obtained after the ninth passage were cultured by the addition of $10^{-8}$ mol/L 1,25(OH)$_2$D$_3$ in the absence or presence of $10^{-7}$ mol/L dexamethasone. After 7 days of incubation, cells from each group were prepared for RNA extraction.

**Coculture System with Stromal-Like Tumor Cells**

Culture of ST-2 Cells and Osteoblastic U2OS Cells

Both ST-2 and U2OS cells were cultured in α-MEM containing 10% FBS, 2 mmol/L l-glutamine, 100 U/ml penicillin, and 100 μg/ml streptomycin, in a humidified atmosphere of 5% CO$_2$ and 95% air at 37°C. When the effect of 1,25(OH)$_2$D$_3$ and dexamethasone was tested, dose dependence was studied by the addition of $10^{-8}$ mol/L 1,25(OH)$_2$D$_3$ in the presence or absence of dexamethasone at doses from $10^{-9}$ to $10^{-6}$ mol/L. All cultures were harvested and used for total RNA extraction.

**Coculture System with Stromal-Like Tumor Cells and RAW$_{264.7}$. Cells**

GCT stromal-like tumor cells at the ninth passage obtained from the liquid nitrogen store were seeded into culture chambers (flask; Nunc) and treated with $10^{-8}$ mol/L 1,25(OH)$_2$D$_3$ and dexamethasone. To the stromal-like tumor cells reaching 50% confluence, 500 RAW$_{264.7}$ cells were added to each chamber. The cocultures were continuously treated with both 1,25(OH)$_2$D$_3$ and dexamethasone. In separate experiments, RAW$_{264.7}$ Cells were cultured in medium alone or in medium with either 30 ng/ml sRANKL, or a combination of $10^{-8}$ mol/L 1,25(OH)$_2$D$_3$ and $10^{-7}$ mol/L dexamethasone. All cultures were fed every 3 days with fresh medium. After 10 days, all cultures were fixed and proceeded to either TRAP histochemistry, CTR immunohistochemistry, or bone resorption pit assay (Zheng et al) to confirm the identity of osteoclasts. TRAP-positive multinuclear cells with more than three nuclei were scored and data were statistically analyzed by Student’s t-tests.

**Calcitonin Receptor Immunofluorescence Confocal Microscopy**

Calcitonin receptor in cocultures of stromal-like cells and RAW$_{264.7}$ cells was detected by the method described by Quinn et al. In brief, cocultures prepared as described above were first incubated with 5% sheep serum in 0.5% BSA/PBS for 10 minutes. The cells were then incubated in PBS/BSA containing rabbit anti-rat/mouse CTR antibody (diluted 1:50) preincubated with MGP-CTR antigen. After a 1-hour incubation at room temperature, the cells were rinsed in PBS, and then PBS/BSA containing Texas Red-X-conjugated swine anti-rabbit Ig antibody (diluted 1:100) was added. After another 1-hour incubation in the dark, the antibodies were rinsed away by washing in PBS. Cells were examined under a confocal laser scanning microscope (Bio-Red 10000) for the immunofluorescence of calcitonin receptor.

**RNA Extraction and RT-PCR**

By the use of RNAzol B (Tel-Test) protocols, total cellular RNA was isolated from solid tumors. Cultured cells derived from human GCT, normal bone, U2OS cells, ST-2 cells, RAW$_{264.7}$ cells, and the coculture system and then reverse-transcribed into cDNA, using 100 units of Moloney-Murine Leukemia Virus (M-MLV) reverse transcriptase (Ambion) according to the manufacturer’s instructions. Primers used for the detection of OPGL, OPG, and RANK are listed in Table 1. Housekeeping genes glyceraldehyde 3-phosphate dehydrogenase (GAPDH) and acidic ribosomal phosphoprotein (36B4) were used as internal controls in the examinations of human and mouse gene expressions, respectively.

**Table 1. Primers for OPG, OPGL, and RANK**

<table>
<thead>
<tr>
<th>Region</th>
<th>Sense primer</th>
<th>Antisense primer</th>
<th>Product size (bp)</th>
</tr>
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<tbody>
<tr>
<td>Human OPG</td>
<td>5’ GCTAAGCTCCTTCCAG 3’</td>
<td>5’ TGATTGGACCTGGTTACC 3’</td>
<td>324</td>
</tr>
<tr>
<td>Mouse OPG</td>
<td>5’ AAAGCACCCTGTAAGAAAAGC 3’</td>
<td>5’ CCCTTTATCCCTCCTACACT 3’</td>
<td>257</td>
</tr>
<tr>
<td>Human OPG</td>
<td>5’ GCCAGGGGAGGATGGTAG 3’</td>
<td>5’ TTAGCTGGAAGTTTTGCC 3’</td>
<td>486</td>
</tr>
<tr>
<td>Mouse OPGL</td>
<td>5’ AAGCTTGGATCATCCACAGAATACG 3’</td>
<td>5’ AAGCTTCAGTCTATGTCGAACTT 3’</td>
<td>726</td>
</tr>
<tr>
<td>Human RANK</td>
<td>5’ TTAGCAGCGTGTTTGACGGC 3’</td>
<td>5’ ACCTAGACCGTATGCTGTCG 3’</td>
<td>497</td>
</tr>
<tr>
<td>Mouse RANK</td>
<td>5’ AAGATGGTTCCAGAAGAGC 3’</td>
<td>5’ CATAGAGTCAGTTCGCTGGCA 3’</td>
<td>350</td>
</tr>
<tr>
<td>Human GAPDH</td>
<td>5’ GGACTCGACGGTTTGTG 3’</td>
<td>5’ GTGATGGAGAATTCATTGATT 3’</td>
<td>206</td>
</tr>
<tr>
<td>Mouse 36B4</td>
<td>5’ TCATTGTGGGACGACA 3’</td>
<td>5’ TCTCCGACTTCCCTTT 3’</td>
<td>832</td>
</tr>
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was at 72°C for 10 minutes. The PCR products were electrophoresed on a 1.5% agarose gel, stained with ethidium bromide, and absorbance measured by densitometry.

**Fluorescence in Situ Hybridization**

The cDNA fragments of human OPGL, OPG, and RANK, which were 486, 324, and 497 bp, respectively, were generated by RT-PCR of total RNA from GCT solid tumor and then separately inserted into pCR2.1 with the Original TA Cloning Kit (Invitrogen). Recombinant plasmids with correct orientation were then purified and transcribed into digoxigenin-labeled antisense riboprobes with T7 RNA polymerase, using a DIG RNA labeling kit (Boehringer Mannheim). All of the clones were sequenced for the confirmation of authentic genes. In situ hybridization was performed as previously described.23,24 The final concentration of each probe in hybridization solution was 0.3 ng/μL, and RNase treatment (100 μg/ml) before hybridization was used as the negative control. Detection of hybridization products was performed with a fluorescent antibody enhancer set (Boehringer Mannheim). We counterstained slides with 2 μg/ml of propidium iodide for 30 minutes at room temperature to view nuclei or with TRAP histochemistry to confirm the characteristics of both osteoclast-like giant cells and macrophage-like cells. Signals were detected by confocal microscopy (BioRed, 1000).

**Results**

**Expressions of OPG, OPGL, and RANK Genes in GCT**

To allow estimation of the level of OPG, OPGL, and RANK gene transcripts in solid tumor and cultures of GCT, cycle-dependent PCR reactions were performed to generate amplification curves for each gene, using specific primers (Figure 1A). A total of 28 cycles was selected for semiquantitation of OPGL and OPG gene transcripts, and 33 cycles for RANK and 26 cycles for GAPDH gene expression. As shown in Figure 1B, all GCT solid tumors expressed OPGL, OPG, and RANK mRNAs, whereas both normal bone and U2OS cells contained much less OPGL gene transcript and no RANK gene transcript. It appears that the ratio of OPGL/OPG in GCT was higher than that in normal bone and osteosarcoma. No correlation of OPGL, OPG, and RANK expression with the Enneking Clinical Stage25 of GCT was found at presentation. Two cases of GCT at stage III (Figure 1B, lanes 2 and 3) expressed almost the same level of these gene transcripts as others in GCT at stage I or II (Figure 1B, lanes 1, 4, and 5).

**Localization of OPGL, OPG, and RANK Gene Transcripts in GCT**

To further examine the localization of OPGL, OPG, and RANK gene transcripts in GCT at the cellular level, fluorescence in situ hybridization (FISH) was performed by using digoxigenin-labeled specific riboprobes (Figure 2). Signals for each gene transcript were distributed differently in the cytoplasm of various cell types. OPG signals were detected in spindle-shaped stromal-like tumor cells, macrophage-like mononuclear cells, and multinuclear osteoclast-like giant cells. However, OPGL signals were mainly in stromal-like tumor cells, whereas RANK signals were mainly in macrophage-like mononuclear cells and multinuclear osteoclast-like giant cells. It is noteworthy that the RANK-positive macrophage-like mononuclear cells were also positive for TRAP when double staining of RANK and TRAP was used (data not shown). Treatment with RNase resulted in the absence of signals in all cells, indicating the specificity of the probes for target mRNA sequence (Figure 2).

**Regulation of OPGL and OPG mRNAs by 1,25(OH)2D3 and Dexamethasone in Mesenchymal Stromal Cells**

Various osteotropic hormones have been shown to be the regulators of osteoclastogenesis.3,25,26 Because GCTs are considered to arise from mesenchymal stromal cells that have the capacity to recruit and harbor osteoclasts,9–16 we attempted to determine whether 1,25(OH)2D3 and dexamethasone were capable of regulating OPGL and OPG expression in the stromal-like tumor cells of GCT and the mouse bone marrow-derived
the cells were incubated with 100 g giant cells (g). Original magnification, 400. C: The signal disappeared when the cells were incubated with 100 µg/ml of RNase before hybridization of OPG. Original magnification, 400. D and E: The OPGL gene transcript was found in stromal-like tumor cells (t) but not in multinuclear osteoclast-like giant cells (g). Original magnification, 400. F: The signal disappeared when the cells were incubated with 100 µg/ml of RNase before hybridization of OPGL. Original magnification, 400. G and H: The RANK gene transcript was detected in round macrophage-like cells (m) and multinuclear osteoclast-like giant cells (g), but not in stromal-like tumor cells (t). Original magnification, 400. I: The signal disappeared when the cells were incubated with 100 µg/ml of RNase before hybridization of RANK. Original magnification, 400.

Figure 2. Localization of OPG, OPGL, and RANK gene transcripts in GCT by FISH. Hybridization signals are in green (FITC), and nuclei staining is in red (PI). A and B: The OPG gene transcript was detected in multinuclear osteoclast-like giant cells (g) and spindle-shaped stromal-like tumor cells (t). It seems that the round mononuclear cells display weak positivity of the OPG mRNA signal. Original magnification, 400. C: The signal disappeared when the cells were incubated with 100 µg/ml of RNase before hybridization of OPG. Original magnification, 400. D and E: The OPGL gene transcript was found in stromal-like tumor cells (t) but not in multinuclear osteoclast-like giant cells (g). Original magnification, 500. F: The signal disappeared when the cells were incubated with 100 µg/ml of RNase before hybridization of OPGL. Original magnification, 500. G and H: The RANK gene transcript was detected in round macrophage-like cells (m) and multinuclear osteoclast-like giant cells (g), but not in stromal-like tumor cells (t). Original magnification, 400. I: The signal disappeared when the cells were incubated with 100 µg/ml of RNase before hybridization of RANK. Original magnification, 400.

mesenchymal stromal cell line ST-2 cells. Cells were treated with 1,25(OH)2D3 at 10−8 mol/L and dexamethasone at 10−7 mol/L for a period of 7 days. As shown in Figure 3A, 1,25(OH)2D3 significantly increased OPGL mRNA levels while reducing OPG levels in ST-2 cells. Dexamethasone alone had no effect on OPGL and OPG gene expression (data not shown) but enhanced those effects of 1,25(OH)2D3 on both gene expression in a dose-dependent manner (Figure 3A, bottom). On the other hand, the effect of 1,25(OH)2D3 and dexametha-

Figure 3. Regulation of gene expression of OPGL and OPG by 1,25(OH)2D3 and dexamethasone. Housekeeping genes GAPDH and 36B4 were used as internal controls for tumor cells of GCT and mouse ST-2 cells, respectively. A: Gene expression of OPGL and OPG in ST-2 stromal like cells. Top: 1,25(OH)2D3 at 10−8 mol/L and dexamethasone at 10−7 mol/L increased the level of OPGL mRNA but decreased the level of OPG gene transcript. Bottom: dose effects of dexamethasone on 1,25(OH)2D3-regulated OPGL and OPGL gene expression. Note that dexamethasone increased the level of OPGL but decreased the level of OPG in a dose-dependent manner. B: Gene expression of OPGL and OPG in cultured stromal-like tumor cells of GCT. Top: 1,25(OH)2D3 at 10−8 mol/L and dexamethasone at 10−7 mol/L increased the level of OPGL mRNA but had no obvious effect on OPG gene expression in case 2 GCT primary culture (ninth passage). Bottom: dexamethasone increased OPGL levels and decreased OPG levels in a dose-dependent manner when combined with 1,25(OH)2D3 in case 4 GCT primary culture (ninth passage).

Discussion

Osteoclasts, the bone-resorbing cells, are derived from hematopoietic cells of monocyte/macrophage lineage.27 The cell-to-cell interaction, the so-called juxtacrine28 between stromal cells/osteoblasts and osteoclast progenitor cells, is essential for the formation of osteoclasts. OPGL was identified as a membrane-bound TNF ligand family protein necessary for the interaction between stromal cells and osteoclast progenitor cells during osteoclastogenesis.3,5,6 Unlike any other cytokines, the recombinant soluble OPGL is able to replace the stromal cells/osteoblasts in the induction of osteoclast formation.1–7 Thus OPGL is considered to be the “master”
cytokine that is necessary and sufficient for the induction of osteoclastogenesis.\textsuperscript{6} GCT is characterized by abundant multinuclear osteoclast-like giant cells scattered among mononuclear cells.\textsuperscript{15} Although the histogenesis of GCT is not fully elucidated, it is generally believed that it is the stromal-like tumor cells that have the ability to recruit circulating monocytes to become multinuclear osteoclast-like giant cells in GCT.\textsuperscript{9–14} We reported here the novel findings that OPGL, the osteoclastogenesis-inducing factor, was abundantly expressed in stromal-like tumor cells of GCT, and that RANK, the receptor for OPGL, was expressed in macrophage-like cells and osteoclast-like multinuclear giant cells. On the other hand, OPG, the decoy receptor for OPGL, was also ubiquitously expressed in the stromal-like tumor cells, indicating that a negative feedback loop may exist in which the tumor cells of GCT themselves may modulate the presentation of OPGL molecules on their surface, which in turn can be inhibited by OPG. Thus the ratio of OPGL and OPG gene expression in tumor cells may determine local osteoclastogenesis and osteoclastic bone resorption in the lesion. If the level of OPGL gene expression exceeds that of OPG in the microenvironment, osteoclast formation may be effectively induced. Conversely, if OPG gene expression is at higher levels than OPGL, osteoclast formation may be suppressed. In this study, we have shown a higher ratio of OPGL to OPG mRNA in all cases of GCT than that in normal bone and osteoblast-like osteosarcoma cell lines. This suggests that the production of OPGL may be of great importance for the tumor cell-induced formation of osteoclast-like giant cells in GCT. Bearing in mind that there is still a limitation of using mRNA to assess the expression of OPGL and OPG, further study should be conducted to elucidate their ratio at the protein level.

Previous studies have shown that the combination of 1,25(OH)\textsubscript{2}D\textsubscript{3} and dexamethasone significantly stimulates osteoclast-like cell formation in cocultures of mouse spleen cells or in mouse bone marrow cultures.\textsuperscript{29–31} In general accord with these findings, our results demonstrated that the combination of these two agents resulted in maximum up-regulation of OPGL gene expression and down-regulation of OPG expression in the mouse mesenchymal stromal cells tested. It is noteworthy that OPGL mRNA was also remarkably increased in primary cultured tumor cells of GCT when treated with 1,25(OH)\textsubscript{2}D\textsubscript{3} and dexamethasone. Although tumor cells of GCT did not show a consistent decrease in OPG mRNA levels in the presence of both agents, it is reasonable to presume that these tumor cells, which express increased OPGL molecules on their membrane in the presence of 1,25(OH)\textsubscript{2}D\textsubscript{3} and dexamethasone, would induce the differentiation of osteoclast progenitors into osteoclasts. In our present study, we have showed that the murine myeloid RAW\textsubscript{264.7} cells differentiated into multinucleated osteoclast-like cells when cocultured with stromal-like tumor cells of GCT in the presence of 1,25(OH)\textsubscript{2}D\textsubscript{3} and dexamethasone. The osteoclast-like cells formed in the coculture satisfied the major criteria of osteoclasts, including multinucleation and the presence of TRAP activity and calcitonin receptor. Given that the induction of osteoclast formation is required by the addition of 1,25(OH)\textsubscript{2}D\textsubscript{3} and dexamethasone, which are the agents for up-regulation of OPGL expression, it is possible that tumor cell-induced osteoclast-like giant cell formation is mediated through the OPGL molecule expressed in stromal-like tumor cells.

In summary, we have shown for the first time that stromal-like tumor cells of GCT express OPGL, whereas...
macrophage-like cells and multinuclear osteoclast-like giant cells express the receptor for OPG, the RANK. OPG, the decoy receptor for OPG, was also found in GCT. However, the ratio of OPG to OPG mRNA was much higher in GCT than that in normal bone and osteosarcoma cells. Osteotropic agents 1,25(OH)2D3 and dexamethasone up-regulate gene expression of OPG in tumor cells of GCT. RAW264.7 cells, which express high levels of RANK mRNA, can differentiate into osteoclast-like cells when cocultured with stromal-like tumor cells in the presence of 1,25(OH)2D3 and dexamethasone. Our findings suggest that OPG may be essential for the tumor cell to induce the formation of osteoclast-like giant cells in GCT. It appears that the ratio of OPG/OPG produced by stromal-like tumor cells may contribute to the degree of osteoclast-like giant cell formation and bone destruction in GCT.

References


