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Human Osteoblasts Support Hematopoiesis through the Production of Granulocyte Colony-stimulating Factor

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Summary

Previous attempts at identifying the constitutive source(s) of granulocyte colony-stimulating factor (G-CSF) in human bone marrow have been unsuccessful despite the fact that normal bone marrow supports abundant myelopoiesis in vivo. We hypothesized that the intimate physical association between bone and hematopoietic cells facilitates interactions between osteoblasts and hematopoietic stem cells. Here we provide the first direct evidence that human osteoblasts participate in hematopoiesis by constitutively producing G-CSF and present the protein in a membrane-associated fashion to human hematopoietic progenitors. These results suggest a direct and central role for osteoblasts in normal myelopoiesis.

rematopoietic stem cell differentiation after birth is largely restricted to the bone marrow cavity. Whether osteoblasts and bone matrix provide essential adhesive, stimulatory, or regulatory signals to stem cells is not known. Overall, the unique physiological conditions that bone tissue provides for hematopoietic cells is essentially a mystery. Mesenchymal cells of fibroblast origin, dispersed throughout the marrow cavity, secrete cytokines such as GM-CSF, IL-6, and c-kit ligand to support basal hematopoiesis, and respond to inflammatory monokines by secreting large quantities of GM-CSF and G-CSF (1-7). This cytokine secretion profile is not, however, specific to bone marrow and is shared by fibroblasts from many tissues. In addition, in the absence of inflammation human fibroblasts do not produce G-CSF, and the source of basal G-CSF production that supports normal granulopoiesis is unknown (6, 7).

The development of the bone marrow cavity is a coordinated process in which blood precursors migrate and colonize spaces carved out of embryonic bone and cartilage (8). Thus, an intimate physical association between bone cells and blood cells is established early in life. Since (a) osteoblasts are found on endosteal marrow surfaces; (b) primitive hematopoietic stem cells are closely approximated with the endosteal surfaces rather than randomly distributed throughout the marrow cavity; and (c) many stromal cell lines share several phenotypic characteristics with osteoblasts (9–14), we hypothesized that osteoblasts might directly support stem cell survival and/or differentiation. In particular we asked whether osteoblasts might be a heretofore undetermined source of constitutively produced G-CSF in the bone marrow microenvironment.

Materials and Methods

Human Osteoblasts. Human osteoblasts were obtained using a modification of methods described by Robey and Termine (15). Normal human trabecular bone was obtained from patients undergoing orthopedic surgery in accordance with the University of Michigan's Policies for Human Subjects. Bone cleaned of loosely adherent tissue was ground to produce a uniform particle size (size ≤1 mm²) (BioComp Minimill; W. Lorenz, Jacksonville, FL) and incubated in 1 mg/ml bacterial collagenase (Type P; Boehringer Mannheim Biologicals, Indianapolis, IN). The explants were placed into culture until confluent monolayers were produced in a 1:1 (vol/vol) mixture of F12/DMEM medium (Biofluids Inc., Rockville, MD) with low Ca⁺² and 10% FCS. Thereafter, cultures were maintained in calcium-replete DMEM/F12 (1:1 vol/vol) medium containing 10% heat inactivated FCS, antibiotics, 10 mM β -glycerophosphate, and 10 mg/ml L-ascorbate. To verify that the cells were osteoblasts, several histochemical assays were performed including in vitro mineralization, and expression of high levels of alkaline phosphatase (16).

RNA Preparation. Total cellular RNA was recovered from osteoblasts as reported in (7). RNA quantity and integrity were checked by gel electrophoresis with ethidium bromide and absorbance at A₂₆₀/A₂₈₀.

Oligonucleotide Primers. Sense and antisense primers were prepared by the oligonucleotide synthesis core at the University of Michigan. The primers used in these investigations were reported previously (7) with the exceptions of those for osteocalcin (nucleotides 1046–1066 sense) GGCAGCGAGGTAGTGAAGAG and (nucleotides 1364–1384, antisense) GATGTGGTCAGCCA-ACTCGT.

Reverse Transcriptase Polymerase Chain Reaction (RT-PCR). RT-PCR was performed for 35 cycles as previously reported (7). PCR products were electrophoresed in 3% agarose and visualized with ethidium bromide. As positive cytokine controls, RNA was isolated from either peripheral blood lymphocytes stimulated with PHA (3.0 μ g/ml) and PMA (3 ng/ml) or IL-1 α (25 U/ml) stimulated human bone marrow stromal cells. To control for DNA contamination, reverse transcriptase was omitted from the reverse transcriptase reaction.

ÉLISAs. G-CSF ELISAs were performed using the doubleantibody sandwich method (R&D Systems Inc., Minneapolis, MN). Based upon parallel assays of known diluted standards, the sensitivities of the assays for G-CSF in unconcentrated medium is 25-2,000 pg/ml. Aliquots of conditioned medium were concentrated 4-10-fold by centrifugation at 1,000 g in a 25° C fixed angle JA-17 rotor (Beckman Instruments Inc., Fullerton, CA) in Centricon-10 concentrators (Amicon Division of W. R. Grace and Co., Beverly, MA) until the desired volume was reached. Used in conjunction with concentrated medium, the G-CSF ELISA was able to detect concentrations as low as 4 pg/ml.

Immunohistochemistry. Indirect immunohistochemistry phasecontrast microscopy was performed for G-CSF on 2% paraformaldehyde-fixed primary human osteoblasts grown in 96-well tissue culture plates. Paraformaldehyde-fixed primary human osteoblasts were incubated with 10 μ g/ml of a murine monoclonal anti-human G-CSF in PBS (M7; kind gift of Dr. Michael Widmer, Immunex Corporation, Seattle, WA) or an isotype control (FOPC-21, Sigma Chemical Co., St. Louis, MO) at 4°C for two hours followed by a rabbit antimurine FITC-conjugated serum at a 1/64 dilution (Sigma Chemical Co.). Nonspecific binding was blocked with 10% normal rabbit serum.

Isolation of $CD34^+$ Cells. Bone marrow cells obtained from healthy adult volunteers were diluted 1:4 (vol/vol) in IMDM and separated by density separation on Ficoll-Hypaque (specific gravity 1.077) to recover mononuclear cells. After two rounds of plastic adherence at 37°C for 1 h each in IMDM medium with 20% FCS (to remove monocytes, platelets, and megakaryocytes), the nonadherent cells were recovered. CD34⁺ hematopoietic progenitor populations were isolated using an avidin-biotin immunoaffinity process (CellPro Inc., Bothell, WA).

Coculture of NFS-60, CD34⁺, and Osteoblasts. CD34⁺ or NFS-60 cells were seeded directly onto osteocalcin⁺, c-kit ligand⁻ confluent osteoblast monolayers in 96-well tissue culture plates at a final density of 10⁴ cells/well and incubated for 3–14 d. Either vehicle, 10 μ g/well of an affinity purified IgG fraction of neutralizing goat anti-human G-CSF serum (R&D Systems), or 10 μ g/well of normal goat IgG serum were added to cultures daily. Absolute cell numbers were determined by manual hemocytometer counting in PBS containing 0.4% trypan blue (Sigma Chemical Co.).

Results and Discussion

We first determined whether unstimulated primary human osteoblasts are a normal source of G-CSF. Primary osteoblasts were obtained using the methods of Robey and Termine (15), where osteoblasts emerge from collagenase-digested human bone during the second to third week of culture. We found that these cells express several functional characteristics of osteoblasts including the expression of mRNA for matrix Gla protein (not shown) and the osteoblast-specific protein, osteocalcin (bone Gla protein) (16–18) (Fig. 1), mineralization of their extracellular matrix, and the expression of alkaline phosphatase examined by histochemical methods (data not shown) (16).

Primary human osteoblast cultures were examined for their



Figure 1. RT-PCR detection of cytokine mRNA by (A) primary human osteoblasts and (B) stromal cells. RT-PCR was performed using primers for IL-1 α , TNF, hymphotoxin (LT), IL-6, G-CSF, GM-CSF, IL-3, c-kit ligand, and osteocalcin. Negative controls omitted reverse transcriptase from the reverse transcription reaction. RNA from IL-1 β and TNF- α stimulated stromal cells served as positive controls for G-CSF and c-kit ligand.

expression of mRNA for several cytokines using RT-PCR. As shown in Fig. 1, osteoblasts constitutively express mRNA for several cytokines including TNF- α , IL-6, GM-CSF, and G-CSF, but failed to express mRNA for IL-1 α , IL-3, and *c-kit* ligand. As *c-kit* ligand is constitutively produced by stromal fibroblasts, the absence of detectable mRNA for this hematopoietic growth factor suggests minimal fibroblast contamination in the primary cultures (Fig. 1 and 4, 6, 7). To further assure that these results are representative of normal osteoblast physiology, we induced the differentiation of osteoprogenitors using a serum deprivation method (19). A similar cytokine profile was observed for these osteocalcin⁺, *c-kit* ligand⁻ cells. In addition, we noted a comparable pat-



Figure 2. Immunohistochemical detection of G-CSF on osteoblast cell surfaces. 2% paraformaldehyde-fixed primary human osteoblasts (A) were incubated with (C) 10 μ g/ml of murine monoclonal anti-human G-CSF in PBS or (B) an isotype control at 4°C for 2 h followed by a rabbit antimurine FITC-conjugated serum at a 1:64 dilution (Sigma Chemical Co.). Nonspecific binding was blocked with 10% normal rabbit serum.

tern of cytokine mRNAs expressed by the human osteosarcoma cell lines, MG-63 and SaOS-2 (data not shown).

The RT-PCR results suggest that osteoblasts constitutively express G-CSF mRNA. We next asked whether the G-CSF message is translated and how the protein is presented to hematopoietic progenitors. To determine whether the G-CSF message is translated, we performed immunohistochemical staining for G-CSF on paraformaldehyde-fixed primary human osteoblasts using a monoclonal anti-human G-CSF antibody. As shown in Fig. 2, G-CSF is localized on the cell membrane of the osteoblasts and not in the interstitial extracellular matrix. To quantify G-CSF production from unstimulated osteoblasts expressing G-CSF mRNA, salt extracts of extracellular matrix (20, 21) and concentrated, conditioned medium were analyzed for G-CSF by ELISA (7). As positive controls for G-CSF production, osteoblasts or stromal cells were exposed to lipopolysaccharide or IL-1 α , respectively (22-24). LPS stimulation of osteoblasts resulted in the production of 21 pg/ml/24 h/10⁴ cells of soluble G-CSF. No G-CSF could be detected in the medium or extracellular matrix of unstimulated osteoblasts, suggesting that under basal conditions G-CSF may not be released as a soluble protein.

To determine whether the G-CSF detected by immunohistochemical methods has functional activity, we evaluated the proliferation of the G-CSF-dependent cell line NFS-60 in coculture with osteocalcin⁺, c-kit ligand⁻ human explant osteoblasts (25). A neutralizing antibody to human G-CSF was used to directly test whether osteoblast-derived G-CSF has biological activity. As shown in Fig. 3, NFS-60 cells proliferate in the presence of osteoblast monolayers but not in medium alone. Inclusion of a neutralizing anti-G-CSF antibody in the cultures caused a 41% reduction in the proliferation of the NFS-60 cells. The proliferation of NFS-60 cells was not affected by a control antiserum nor were significant differences in proliferation of the NFS-60 cells observed when the cells were grown in osteoblast-conditioned medium (50% vol/vol) alone, with or without the neutralizing anti-human G-CSF antibody.

These results obtained with the NFS-60 cell line suggest that osteoblasts might stimulate the proliferation of human



Figure 3. NFS-60 cell proliferation on osteoblast monolayers. NFS-60 cells were seeded onto confluent human osteoblasts at a final density of 10⁴ cells/well in 96-well tissue culture plates for 3 d. Where indicated vehicle (no AB), 10 μ g/well of an affinity purified IgG fraction of neutralizing goat anti-human G-CSF serum (Anti-G-CSF) or 10 μ g/well of normal goat IgG serum (control AB) were added daily. Absolute NFS-60 cell numbers were determined by manual hemocytometer counting and reported as mean \pm SD (n = 4). (Asterisk) Significant difference from antibody or vehicle control, p < 0.01.



Figure 4. Human CD34⁺ cells obtained by affinity purification of nonadherent low density mononuclear bone marrow cells (CellPro Inc.) were seeded directly into 96-well tissue culture plates or onto confluent osteocalcin⁺ c-kit ligand⁻ human osteoblast monolayers at a final density of 10⁴ cells/well and incubated for 14 d. The Results of one of two experiments are presented. Where indicated vehicle (no AB), 10 μ g/well of either neutralizing goat anti-human G-CSF serum or goat IgG serum (control AB) were added daily (R&D Systems). Cell counts were determined by manual hemocytometer counting and are reported as mean ± SD (n = 3). The hematopoietic cells recovered after 2 wk of coculture maintained an immature (23 ± 6% myeloblasts/promyelocytes; 51.7 ± 5.5% myelocytes/metamyelocytes; 21 ± 5% bands/polymorphonuclear neutrophils; and 4.3 ± 0.6% monocytes, treatment with anti-G-CSF reduced the recovery of band/polymorphonuclear neutrophils to 8.7 ± 10%). (Asterisk) Significant difference from antibody or vehicle control, p < 0.01.

hematopoietic progenitors by producing G-CSF. To examine this possibility, CD34⁺ hematopoietic progenitor cells were isolated and seeded into confluent osteocalcin+, c-kit ligandosteoblast monolayers. At 14 d the hematopoietic cells were recovered and counted. The significance of G-CSF in this system was tested by the addition of an antiserum to human G-CSF. An eightfold increase in cell recovery was observed for hematopoietic cells grown in coculture with osteoblasts over a 2-wk period (Fig. 4). 55.4% of the osteoblast-induced increase in cell recovery was directly blocked by coincubation of neutralizing anti-G-CSF antibody in the culture. It is interesting that 44.5% of the growth activity produced by osteoblasts was not due to G-CSF, either alone or in combination with other osteoblast-derived cytokines. Furthermore, in Wright-Geimsa cytospin preparations many of the hematopoietic cells recovered after 2 wk of coculture maintained an immature morphology $(23 \pm 6\% \text{ myeloblasts}/$ promyelocytes; 51.7 ± 5.5% myelocytes/metamyelocytes; $21 \pm 5\%$ bands/polymorphonuclear neutrophils; $4.3 \pm 0.6\%$ monocytes). These data suggest that osteoblasts might produce additional factors that support the survival and limited proliferation of primitive myeloid cells as well.

The biologic data using CD34⁺ and the NFS-60 cell line in conjunction with the mRNA, and immunohistochemistry results strongly suggest that human osteoblasts produce G-CSF. Osteoblast-derived G-CSF appears to be accessible to target cells when it is associated with osteoblast's cell membrane. The possible mechanisms by which growth factors become cell associated or retained by components of the extracellular matrix and presented to target cells have not yet been fully elucidated. Cell associated forms of M-CSF, c-kit ligand, IL-1 α , and matrix-associated forms of IL-3 and GM-CSF have been described (20, 21, 25-30). The implications of the experiments evaluating G-CSF production with bioassays vs. ELISA of conditioned osteoblast medium suggest that either (a) immunoreactive G-CSF is diffusible only over extremely short distances before inactivation; (b) the neutralizing G-CSF antibody does not have access to the microenvironment between osteoblasts and stem cells; (c) directed cell-cell transfer of G-CSF occurs between the osteoblasts and hematopoietic progenitors, as has been reported for IL-2 (31); or (d) G-CSF or a closely related molecule is presented to hematopoietic cells as a cell membrane associated protein or bound in an active form to extracellular matrix molecules. Of interest, at least three forms of G-CSF have been identified on immunoblots, including 18-, 28-, and 32-kD forms (32). To date it appears that the major secreted species is the 18-kD form, whereas the biological nature of the higher molecular mass proteins are unknown but possibly represent cellassociated or inactive protein precursors (32). We found no evidence for G-CSF detectable in either the LPS-treated or untreated osteoblast matrix fraction; until more extensive investigation of osteoblast's extracellular matrix is performed however, these data do not rule out such a possibility.

It is interesting that the antibody to G-CSF neutralized only 55% of the CD34⁺ cell's proliferative activity produced by the osteoblasts (Fig. 4). Thus, more than one factor produced by osteoblasts may support the proliferation of hematopoietic stem cells. At present we do not know the identity of the molecule(s) which are responsible for this activity. In preliminary experiments neutralizing antibody to IL-6 and GM-CSF failed to affect proliferation of the NFS-60 cell line. It is possible that unique osteoblast-derived cytokines are responsible for proliferative effects on NFS-60 and CD34⁺ cells. These observations suggest that stem cell self-renewal is influenced by either osteoblast-derived cytokines or adhesive interactions between the two cell types or possibly both.

In sum, strong evidence indicates that human osteoblasts produce cytokine signals that regulate hematopoiesis. Hematopoietic myeloid stem cell differentiation occurs in direct proximity to osteoblasts within the bone marrow cavity. We speculate that this intimate physical association established early in life facilitates interactions between bone and hematopoietic cells. We determined that human osteoblasts are a constitutive source of the hematopoietic growth factor G-CSF. If osteoblasts are a major source of the colony stimulating cytokines in bone marrow, then understanding of the regulation of osteoblast-elaborated cytokines and the functional consequences of alterations in osteoblast metabolism are critical to our understanding of normal hematopoiesis. The authors wish to thank Drs. L. G. Biesicker, L. Xiao, L. B. Taichman, and J. Caldwell for their helpful discussion and control RNA; D. Gianolla for technical assistance; and Drs. L. Mathews, S. W. Weiss, and E. L. Poy for their invaluable help in obtaining human bone samples.

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