TGF-β signaling blockade inhibits PTHrP secretion by breast cancer cells and bone metastases development

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Breast cancer frequently metastasizes to the skeleton, and the associated bone destruction is mediated by the osteoclast. Growth factors, including transforming growth factor-β (TGF-β), released from bone matrix by the action of osteoclasts, may foster metastatic growth. Because TGF-β inhibits growth of epithelial cells, and carcinoma cells are often defective in TGF-β responses, any role of TGF-β in metastasis is likely to be mediated by effects on the surrounding normal tissue. However, we present evidence that TGF-β promotes breast cancer metastasis by acting directly on the tumor cells. Expression of a dominant-negative mutant (TβRIIAc) of the TGF-β type II receptor rendered the human breast cancer cell line MDA-MB-231 unresponsive to TGF-β. In a murine model of bone metastases, expression of TβRIIAc by MDA-MB-231 resulted in less bone destruction, less tumor with fewer associated osteoclasts, and prolonged survival compared with controls. Reversal of the dominant-negative signaling blockade by expression of a constitutively active TGF-β type I receptor in the breast cancer cells increased tumor production of parathyroid hormone–related protein (PTHrP), enhanced osteolytic bone metastasis, and decreased survival. Transfection of MDA-MB-231 cells that expressed the dominant-negative TβRIIAc with the cDNA for PTHrP resulted in constitutive tumor PTHrP production and accelerated bone metastases. These data demonstrate an important role for TGF-β in the development of breast cancer metastasis to bone, via the TGF-β receptor-mediated signaling pathway in tumor cells, and suggest that the bone destruction is mediated by PTHrP.


Introduction

In 1889, Paget (1) proposed the “seed and soil” hypothesis to explain the predilection with which breast cancer grows in bone. Despite these observations, the mechanisms underlying the affinity with which breast cancer grows in bone are not completely understood. Breast cancer metastasizes to bone in greater than 80% of patients with advanced disease and causes local osteolysis (2). The associated pain, pathological fracture, hypercalcemia, and nerve compression syndromes are consequences of the bone destruction. These morbid complications can be devastating, because patients with breast cancer and bone metastases may survive for many years.

A unique characteristic of the skeleton is the storage within bone matrix of immobilized growth factors such as transforming growth factor (TGF)-β, insulin-like growth factor (IGF)-1 and -2, fibroblast growth factor (FGF)-1 and -2, and platelet-derived growth factors (3). The most abundant repository for TGF-β is the bone matrix, and it is released locally in the microenvironment as a consequence of osteoclastic bone resorption (4). Thus, tumor cells with the capacity to stimulate osteoclastic bone resorption may enrich the bone microenvironment with growth factors that may alter behavior of tumor cells. Data in support of this include histological examination of breast cancer metastatic to bone, which reveals tumor cells adjacent to bone-resorbing osteoclasts (5). Bisphosphonates, potent inhibitors of osteoclastic bone resorption, decrease the morbidity associated with breast cancer bone metastases (6, 7). These findings indicate that the bone destruction by breast cancer is mediated by tumor stimulation of osteoclastic bone resorption.

Parathyroid hormone-related protein (PTHrP) is a tumor product (8–10) that stimulates osteoclastic bone resorption and renal tubular reabsorption of calcium by binding to a common PTH/PTHrP receptor (11, 12). The majority of patients with solid tumors and hypercalcemia have increased plasma PTHrP concentrations (13). PTHrP may have a more common role in malignancy as a mediator of osteolytic bone metastasis in breast cancer, even in the absence of hypercalcemia. Women with PTHrP-positive primary breast tumors are more likely to develop bone metastases (14). Human breast cancer cells express PTHrP more often in bone (15) than in the primary (16) or soft tissue sites, and neutralizing antibodies to PTHrP inhibit the development of osteolytic metastases by human breast cancer cells in vivo (17). The reasons for increased expression of PTHrP in bone are unknown, but bone-derived growth factors may be responsible.

Because TGF-β is one of the most abundant growth...
factors in bone matrix (3), is released in active form during osteoclastic bone resorption (4), and increases PTHrP expression by cancer cells in vitro (18–21), we postulated that TGF-β was responsible for enhancing PTHrP production in the bone microenvironment and the subsequent bone destruction. To test this hypothesis, the human breast cancer cell line MDA-MB-231 (22) was transfected with a cDNA encoding a TGF-β type II receptor lacking most of the cytoplasmic domain (TβRIIΔcyt), which acts as a dominant–negative to block the biologic effects of TGF-β (23). Expression of this mutant receptor in human breast cancer cells inhibited TGF-β-stimulated PTHrP production and blocked the growth inhibitory effects of TGF-β in vitro. Blockade of TGF-β responsiveness in breast cancer cells resulted in decreased osteolysis, less tumor burden in bone, and enhanced survival in mice bearing tumors that expressed the dominant–negative receptor. When TGF-β responsiveness of the breast cancer cells expressing the dominant–negative type II receptor was restored by expression of a constitutively active TGF-β type I receptor subunit, growth of tumor in bone and osteolysis were markedly enhanced, and survival was decreased. Because PTHrP production was increased fivefold in this tumor line, the data suggested that the effects of TGF-β to enhance bone destruction were mediated by PTHrP. To confirm this notion, a cDNA-encoding human PTHrP was transfected into the breast cancer cells expressing the dominant–negative type II receptor. The resulting increase in constitutive PTHrP production in vitro was associated with accelerated bone metastases in vivo.

**Methods**

**Cells.** MDA-MB-231 cells were cultured in DMEM (Life Technologies Inc., Rockville, Maryland, USA) containing 10% FCS (HyClone Laboratories, Logan, Utah, USA), 1% penicillin/streptomycin, and nonessential amino acids (GIBCO BRL, Gaithersburg, Maryland, USA). To test the effect of bone-derived growth factors on PTHrP secretion by MDA-MB-231 cells, 10^6 cells/ml were plated onto 48-well plates. When near confluence, cells were washed with PBS, and 250 μl of serum-free DMEM containing the following growth factors was added to each well: TGF-β1 (0, 1, 5, or 10 ng/ml), FGF-1 (0, 4, 20, or 40 ng/ml), PDGF (0, 1, 5, 10, or 40 ng/ml), IGF-1 (0, 1, 5, 10, or 40 ng/ml), and IGF-2 (0, 1, 5, 10, or 40 ng/ml). TGF-β1, FGF-1, FGF-2, PDGF, IGF-1, and IGF-2 were purchased from R&D Systems Inc. (Minneapolis, Minnesota, USA). Conditioned media were collected after 48 h, stored at –70°C for PTHrP measurement, and cell number was counted for each well to correct the PTHrP concentration of the conditioned media. Triplicate measurements were performed for each treatment.

To measure the effects of TGF-β on the growth rate of MDA-MB-231 cells and respective clones, 10^6 cells/ml were plated onto each of two 24-well plates. One plate was treated with TGF-β1 (20 ng/ml) and the other with vehicle. Cell number was counted daily for 8 days; each measurement was performed in triplicate.

**Stable transfection of MDA-MB-231 cells with cDNA for the truncated type II TGF-β receptor (TβRIIΔcyt).** TβRIIΔcyt cDNA (23) was subcloned from pMEP4 into the pcDNA3 expression vector (Invitrogen Corp., Carlsbad, California, USA) as a HindIII–BamHI fragment. The pcDNA3/TβRIIΔcyt DNA or the empty vector, pcDNA3, was transfected into MDA-MB-231 cells by calcium phosphate precipitation. Single clones were isolated by limiting dilution in the presence of the selective marker, G418 (Sigma Chemical Co., St. Louis, Missouri, USA). Clones were screened by measuring the amount of secreted PTHrP in serum-free 48-h conditioned media in the presence or absence of TGF-β (5 ng/ml). Clones in which PTHrP secretion did not increase in response to TGF-β were selected for further study.

**Figure 1**

(a) Effects of bone growth factors on PTHrP secretion from MDA-MB-231 cells in vitro. MDA-MB-231 cells were plated onto 48-well plates and grown to near confluence. Cells were washed and treated with serum-free media containing the respective growth factors for 48 h. PTHrP concentrations in conditioned media were corrected for cell number. Only the results for the highest concentration of each growth factor are shown. Inset: Dose response for PTHrP secretion by MDA-MB-231 cells treated with TGF-β. Values represent the mean ± SEM (n = 3 per group). (b) Effect of TGF-β on PTHrP secretion by MDA-MB-231, MDA/pcDNA3, and MDA/TβRIIΔcyt clones. Respective cells were plated onto 48-well plates and treated as described in a. Values represent the mean ± SEM (n = 3 per group). P = parental MDA-MB-231; EV = empty vector pcDNA3 clone; 1, 2, and 3 are respective MDA/TβRIIΔcyt clones. (c and d) Osteolytic lesion area from radiographs of two separate experiments comparing clones 3 and 2 (c) or clones 1 and 2 (d) with controls of MDA-MB-231 (P) or pcDNA3 vector (EV). Values represent mean ± SEM (n = 4 per group). *P < 0.05, **P < 0.01, ***P < 0.001 vs. controls. PTHrP, parathyroid hormone–related protein; TGF-β, transforming growth factor–β.
TβRI(T204D) cDNA insert (24, 25) was subcloned into pcDNA3.1zeo (Invitrogen Corp.) as a HindIII–BamHI fragment. The pcDNA3.1zeo/TβRI(T204D) DNA or the empty vector, pcDNA3.1zeo, was transfected into the MDA-MB-231 clonal line expressing the truncated type II TGF-β receptor, MDA/TβRIIΔcyt, by calcium phosphate precipitation. Single clones were isolated by limiting dilution in the presence of the selective markers, G418 and zeocin (Invitrogen Corp.), and screened as described. Clones with increased PTHrP production, basally and in response to TGF-β, were selected for further study.

Stable transfection of MDA/TβRIIΔcyt with cDNA for PTHrP. Human preproPTHrP cDNA encoding the 1–141 isoform was subcloned from pCMVIE-AK1-DHFR (26) into pcDNA3.1zeo (Invitrogen Corp.) as a HindIII–BamHI fragment. The pcDNA3.1zeo/PTHrP DNA or the empty vector, pcDNA3.1zeo, was transfected into the MDA-MB-231 clonal line expressing the truncated type II TGF-β receptor, MDA/TβRIIΔcyt, by calcium phosphate precipitation. Single clones were isolated by limiting dilution in the presence of both selective markers, G418 and zeocin (Invitrogen Corp.), and screened as described. Clones with increased PTHrP production, in the basal state, and which did not respond to TGF-β, were selected for further study.

Cross-linking and immunoprecipitation. MDA-MB-231 clones expressing either TβRIIΔcyt (MDA/TβRIIΔcyt) or the empty vector (MDA/pDNA3) were plated onto two 10-cm petri dishes (106 cells per dish). Cells were grown to confluence and incubated for 3 h at 4°C with 125I-labeled TGF-β1 (160 pM) (Du Pont Nen Research Products, Boston, Massachusetts, USA) alone or with cold TGF-β1 (5 nM) as a competitor (23). After washing away the unbound TGF-β1, ligand-receptor was cross-linked with disuccinimidyl suberate and solubilized with 10 mM Tris buffer (pH 7.4) containing 1% Triton X-100, 1 mM EDTA, and protease inhibitors. The receptor complexes were immunoprecipitated with 2.5 μg/ml mouse anti-HA monoclonal antibody, 12CA5 (Boehringer Mannheim Biochemicals, Indianapolis, Indiana, USA), followed by adsorption to protein G-Sepharose (Pharmacia Biotech Inc., Piscataway, New Jersey, USA). This mixture was washed, and bound protein was eluted by heating samples in SDS-PAGE sample buffer containing 100 mM dithiothreitol. Samples were run on 12.5% SDS-PAGE gels under reducing conditions, and gels were fixed, dried, and exposed to film at ~70°C.

Animals

Bone metastasis. In the pilot experiments (Fig. 1, c and d), mice were inoculated with tumor-cell suspensions of MDA/TβRIIΔcyt (independent clones: 1, 2, and 3), MDA/pDNA3, or parental MDA-MB-231 cells into the left cardiac ventricle (n = 4 per group) on day 0. At sacrifice, on day 28, radiographs were obtained and analyzed as described later in this paper. Because MDA/TβRIIΔcyt, clone 2, was the least responsive to TGF-β in vitro (Fig. 1d), more detailed experiments were performed comparing this clone with the controls. In the first set of experiments, mice were inoculated with tumor-cell suspensions of MDA/TβRIIΔcyt (clone 2), MDA/pDNA3, or parental MDA-MB-231 cells into the left cardiac ventricle (n = 13 per group) on day 0 after baseline radiographs, body weights, and blood for Ca2+ and plasma PTHrP concentrations were obtained. Radiographs were taken on day 21 and at sacrifice on day 24 to monitor progression of osteolytic metastases. Ca2+ and body weight were measured weekly after tumor inoculation until sacrifice, at which time most control mice were cachectic and paraplegic. Blood was collected for Ca2+ and PTHrP measurement, and all bones and soft tissues were obtained and analyzed as described later in this paper.

Experimental protocols

Bone metastasis. In the pilot experiments (Fig. 1, c and d), mice were inoculated with tumor-cell suspensions of MDA/TβRIIΔcyt (independent clones: 1, 2, and 3), MDA/pDNA3, or parental MDA-MB-231 cells into the left cardiac ventricle (n = 4 per group) on day 0. At sacrifice, on day 28, radiographs were obtained and analyzed as described later in this paper. Because MDA/TβRIIΔcyt, clone 2, was the least responsive to TGF-β in vitro (Fig. 1d), more detailed experiments were performed comparing this clone with the controls. In the first set of experiments, mice were inoculated with tumor-cell suspensions of MDA/TβRIIΔcyt (clone 2), MDA/pDNA3, or parental MDA-MB-231 cells into the left cardiac ventricle (n = 13 per group) on day 0 after baseline radiographs, body weights, and blood for Ca2+ and plasma PTHrP concentrations were obtained. Radiographs were taken on day 21 and at sacrifice on day 24 to monitor progression of osteolytic metastases. Ca2+ and body weight were measured weekly after tumor inoculation until sacrifice, at which time most control mice were cachectic and paraplegic. Blood was collected for Ca2+ and PTHrP measurement, and all bones and soft tissues were obtained and analyzed as described later in this paper. Autopsy was performed on all mice, and those with tumor in the chest were excluded from analysis, because this indicated that the tumor inoculum did not properly enter the left cardiac ventricle. A separate experiment was similarly performed to assess sur-
vival. In this experiment, mice were sacrificed when they became moribund.

In the second set of experiments, female mice were inoculated with tumor-cell suspensions of clonal MDA-MB-231 lines TαRIIΔcαyt + TβRI(T204D) or TαRIIΔcαyt + pcDNA3.1zeo into the left cardiac ventricle on day 0. Radiographs, body weight, and blood for Ca²⁺ and PTHrP were obtained at baseline and sacrifice as in the first set of experiments. Ca²⁺, body weight, and radiographs were monitored weekly for 4 weeks, at which time the mice were sacrificed. Tissue processing, autopsy, data analysis, and a separate survival experiment were performed as in the first experiment.

In the third set of experiments, female mice were inoculated with tumor-cell suspensions of clonal MDA-MB-231 lines TαRIIΔcαyt + PTHrP (two different clones) or TαRIIΔcαyt + pcDNA3.1zeo into the left cardiac ventricle on day 0. The same parameters were measured as in the previous experiments. Two separate experiments were performed using two different TαRIIΔcαyt + PTHrP clones. Four TαRIIΔcαyt + PTHrP clones were studied in total.

Local tumor growth. To investigate whether expression of the dominant-negative type II TGF-β receptor subunit in breast cancer cells altered tumor growth in sites other than bone, tumor-cell suspensions (10⁷/100 μl PBS) of either MDA/TαRIIΔcαyt, MDA/pcDNA3, or parental MDA-MB-231 cells were inoculated into the right thigh of female nude mice. Tumor volume was measured with calipers and calculated by the formula of an ovoid where L equals midaxis length and W equals midaxis width: tumor volume = 4/3π × L/2 × W/2².

Analytical methods

Ca²⁺ measurement. Ca²⁺ concentrations were measured in whole blood using a Ciba Corning 634 ISE Ca²⁺/pH analyzer (Corning Medical and Scientific Medfield, Massachusetts, USA) as described previously (17).

PTHrP assay. PTHrP concentrations were measured in conditioned media and plasma using a two-site immunoradiometric assay (Nichols Institute, San Juan Capistrano, California, USA) that detects PTHrP(1–72) and has a calculated sensitivity of 0.3 pmol/l (27). PTHrP concentrations in conditioned media samples were calculated from a standard curve generated by adding recombinant PTHrP(1–86) to the specific type of medium (unconditioned) used and were considered undetectable if media concentrations were <0.3 pmol/l before correction for cell number.

Radiographs and measurement of osteolytic lesion area. Animals were x-rayed in a prone position against the film as described previously (17). All radiographs were evaluated without knowledge of treatment groups. The area of osteolytic bone metastases was calculated using a computerized image analysis system. Video images of radiographs were captured using a frame grabber board on a PC system. Quantitation of lesion area was performed using image analysis software (Java, Jandel Video analysis, Jandel Scientific, Corte Madera, California, USA).

Bone histology and histomorphometry. Forelimb and hindlimb bones were removed from mice at time of killing, fixed in 10% buffered formalin, decalcified in 14% EDTA, and embedded in paraffin wax. Sections were stained with hematoxylin, eosin, orange G, and phloxine. The following variables were measured in midsections of tibiae and femora, without knowledge of experimental groups, to assess tumor involvement: total tumor area and osteoclast number per millimeter of tumor/bone interface. Histomorphometric analysis was performed on an OsteoMeasure System (Osteometrics Inc., Atlanta, Georgia, USA).

Statistical analysis

Results are expressed as the mean ± SEM. Data were analyzed by analysis of variance followed by Tukey-Kramer post test. Log-rank test (Wilcoxon survival) was used to analyze survival data. P < 0.05 was considered significant.

Results

Effects of bone growth factors on PTHrP production in vitro by MDA-MB-231 cells. Because previous clinical and experimental studies demonstrated increased PTHrP expression by breast cancer cells in the bone microenviron-
ment, the effect of factors known to be present in bone matrix were tested on PTHrP production by human MDA-MB-231 breast cancer cells in vitro. Only TGF-β significantly increased PTHrP production by these cells in a dose-dependent manner (Fig. 1a). Other growth factors abundant in bone, such as FGF-1 and -2, IGF-1 and -2, and PDGF, had no effect on PTHrP secretion over a wide range of concentrations.

Expression of a truncated type II TGF-β receptor (TßRIIAcyt) in MDA-MB-231 cells. TGF-β was the only factor tested that affected PTHrP secretion, so its role in breast cancer metastasis to bone was investigated by transfecting MDA-MB-231 cells with the cDNA for a truncated type II TGF-β receptor (MDA/TßRIIAcyt) and generating clonal lines. This receptor is truncated at the extracellular domain and has 199 amino acids and a predicted protein size of 22 kDa. A hemagglutinin (HA) epitope was present in the extracellular domain. The mutant receptor binds TGF-β, but because it cannot phosphorylate the type I receptor, signal propagation does not occur (23). Thus, it acts in a dominant–negative fashion to block the biologic effects of TGF-β.

As shown in Fig. 1b, treatment of parental MDA-MB-231 cells and the MDA/pcDNA3 clone with TGF-β1 (5 ng/ml) significantly stimulated PTHrP secretion, while the same treatment of three different MDA/TßRIIAcyt resulted in minimal or no increase. Similar results were observed when the clones MDA/TßRIIAcyt and MDA/pcDNA3 were grown in the absence of the selective marker, G418, for four weeks and indicate that TßRIIAcyt was stably expressed. These data demonstrate that TßRIIAcyt acts as a dominant–negative to block the biologic effects of TGF-β to stimulate PTHrP production by MDA-MB-231 breast cancer cells. In two separate pilot experiments using a mouse model of bone metastases, all three MDA/TßRIIAcyt clones had smaller total osteolytic lesion area on radiographs compared with parental or empty vector controls (Fig. 1, c and d). Because MDA/TßRIIAcyt clone 2 was the least responsive to TGF-β, it was studied in further detail.

Receptor expression in stable clones was demonstrated by cross-linking to 125I-labeled TGF-β1. Figure 2a demonstrates the autoradiograph of 125I-labeled TGF-β1 cross-linking followed by immunoprecipitation with anti-HA antibody in clonal MDA-MB-231 cells expressing the TßRIIAcyt (MDA/TßRIIAcyt, clone 2) or the empty vector (MDA/pcDNA3). Two distinct bands, which were completely competed by cold TGF-β, were present in MDA/TßRIIAcyt. One band had a molecular weight corresponding to the predicted size of the truncated TGF-β type II receptor plus TGF-β dimer (47 kDa). The other band corresponded to the truncated type II TGF-β receptor plus TGF-β monomer, which has a predicted size of 34 kDa. No bands were evident by immunoprecipitation with the anti-HA antibody in the MDA/pcDNA3 clonal cells, which indicates that only MDA/TßRIIAcyt expressed the truncated type II TGF-β receptor. These results also indicate that in the MDA/TßRIIAcyt, this mutant TGF-β type II receptor was expressed at the cell surface and bound TGF-β. Endogenous type II TGF-β receptor was expressed by both cell lines, as demonstrated by 125I-labeled TGF-β1 cross-linking followed by immunoprecipitation with a polyclonal type II TGF-β receptor antibody (data not shown).

Effect of TGF-β on growth of MDA-MB-231, MDA/pcDNA3, and MDA/TßRIIAcyt clonal cells. Because TGF-β also regulates cellular functions of proliferation and differentiation, the effects of TGF-β on growth of parental MDA-MB-231, MDA/TßRIIAcyt (clone 2) and MDA/pcDNA3 clonal cells were studied. As illustrated in Fig. 2b, growth rates were similar for all three cell lines in the absence of TGF-β. TGF-β1 (20 ng/ml) inhibited the growth of parental MDA-MB-231 and MDA/pcDNA3 cells but had

Figure 4
Survival (a) and plasma PTHrP concentrations (b) in mice bearing MDA-MB-231, MDA/pcDNA3, or MDA/TßRIIAcyt tumors. (a) Survival of mice bearing MDA/TßRIIAcyt tumors was significantly longer than that of the controls. (b) Plasma PTHrP concentrations at sacrifice were significantly higher than respective concentrations before tumor inoculation (baseline) in mice bearing control tumors of MDA-MB-231 or MDA/pcDNA3. There was no significant difference between baseline and sacrifice values in mice bearing the MDA/TßRIIAcyt tumors.

Figure 5
Effect of TGF-β on PTHrP secretion by MDA-MB-231 clonal lines, TßRIIAcyt + TßRI(T204D), and TßRIIAcyt + pcDNA3.1zeo. Respective cells were plated onto 48-well plates and treated as in Fig. 1a. Values represent the mean ± SEM (n = 3 per group). *P < 0.01 and ***P < 0.001 compared with TßRIIAcyt + pcDNA3.1zeo (control or TGF-β-stimulated).
no effect on MDA/TβRII∆cyt. Thus, expression of the dominant–negative receptor blocked TGF-β–mediated growth inhibition of MDA-MB-231 cells.

Role of TGF-β in breast cancer metastases to bone. To determine the effects of TGF-β on breast cancer–mediated osteolysis, MDA/TβRII∆cyt (clone 2) or controls were studied in a mouse model of bone metastasis (17). Respective tumor cells were inoculated into the left cardiac ventricle of female nude mice, and serial radiographs were obtained. Representative radiographs from mice 24 days after tumor inoculation are illustrated in Fig. 3a. Osteolytic lesion number and area on radiographs were significantly less in MDA/TβRII∆cyt–bearing mice compared with that of MDA/pcDNA3 and MDA-MB-231 controls (Fig. 3b). These data were consistent with the pilot experiments in which two other MDA/TβRII∆cyt clones, in addition to clone 2, had significantly fewer and smaller bone metastases on radiographs compared with controls (Fig. 1, c and d).

Because radiographic methods assess only bone destruction and are not a direct measurement of tumor area in bone, histomorphometric analysis of forelimb and hindlimb bones was performed. Osteoclast number per millimeter of tumor/bone interface and tumor area in bone was significantly less in MDA/TβRII∆cyt–bearing mice compared with MDA/pcDNA3 and MDA-MB-231 controls (Fig. 3c).

In a separate experiment, survival of mice bearing MDA/TβRII∆cyt was significantly longer than that of the controls (Fig. 4a). In all experiments, there were no differences between MDA/TβRII∆cyt and control groups with regard to metastasis to nonbone sites. Gross and histological assessment of soft tissues revealed little or no metastasis to adrenal glands, lungs, liver, spleen, ovaries, brain, or kidneys. In this survival experiment, plasma PTHrP concentrations did not differ from baseline to sacrifice in the MDA/TβRII∆cyt group (Fig. 4b).

Effect of TβRII∆cyt on local tumor growth. Although there were no differences in metastasis to nonbone sites between MDA/TβRII∆cyt and control groups, there were insufficient metastases to determine whether expression of the dominant–negative receptor significantly affected tumor growth at sites other than bone. Thus, tumor-cell suspensions (10^7 cells/100 μl/mouse) of MDA/TβRII∆cyt, MDA/pcDNA3, or MDA-MB-231 cells were inoculated intramuscularly into the right thigh of female athymic nude mice. Tumors were excised, measured, and weighed at sacrifice, 21 days after tumor inoculation.
Restoration of TGF-β responsiveness into MDA/ TBRIIΔcyt by expression of a constitutively active type I TGF-β receptor. To confirm that the decrease in bone metastasis was due to blockade of TGF-β signaling, the effects of the dominant–negative type II receptor mutation were reversed by the introduction of a constitutively active form of the type I receptor, TBR-I. This receptor has a point mutation at amino acid 204 of the GS domain in which aspartic acid replaces threonine (T204D). In TGF-β signal transduction, TBR-II phosphorylates TBR-I, which then propagates the signal (28). TBRI(T204D) is constitutively phosphorylated and does not require ligand binding or interaction with the type II receptor to mediate TGF-β signaling (24). TGF-β did not stimulate PTHrP production in the TBRIIΔcyt + pcDNA3.1zeo control group, as were blood ionized calcium concentrations (data not shown). There was a fivefold increase in the basal production of PTHrP by the TBRIIΔcyt + TBRI(T204D) clone that was further enhanced by TGF-β (Fig. 5). Thus, expression of the constitutively active TGF-β type I receptor overcame the dominant–negative blockade and restored the TGF-β–stimulated PTHrP production.

Next, we studied the effect of the restoration of TGF-β responsiveness, MDA-MB-231 clonal lines of TBRIIΔcyt + TBRI(T204D) or TBRIIΔcyt + pcDNA3.1zeo control cells were inoculated into the left cardiac ventricle of female nude mice, and serial radiographs were obtained. Four weeks after tumor inoculation, mice bearing TBRIIΔcyt + TBRI(T204D) tumors had significantly more and larger osteolytic lesions on radiographs compared with those bearing TBRIIΔcyt + pcDNA3.1zeo tumors (Fig. 6, a and b). Histomorphometric analysis of hindlimbs supported the radiographic findings and revealed that the osteoclast number per millimeter of tumor/bone interface and tumor area in bone was significantly greater in the TBRIIΔcyt + TBRI(T204D) group when compared with the control group, TBRIIΔcyt + pcDNA3.1zeo (Fig. 6c).

Tumors from mice bearing TBRIIΔcyt + TBRI(T204D) demonstrate tumor replacing the marrow cavity and destruction of both trabecular and cortical bone, whereas bones from mice bearing TBRIIΔcyt + pcDNA3.1zeo had small foci of tumor in the bone marrow cavity with little bone destruction (Fig. 6d). Plasma PTHrP and blood ionized calcium concentrations were also significantly higher in mice bearing TBRIIΔcyt + TBRI(T204D) tumors compared with those bearing the control, TBRIIΔcyt + pcDNA3.1zeo (Fig. 7a). In a separate experiment, mice bearing the TGF-β–responsive tumor, TBRIIΔcyt + TBRI(T204D), had a significantly shorter survival time than mice bearing the tumors that were unresponsive to TGF-β (Fig. 7b).

Overexpression of PTHrP, under the control of a constitutive promoter, into MDA/ TBRIIΔcyt. To determine whether the effects of TGF-β to enhance bone metastases were mediated by PTHrP, the MDA/ TBRIIΔcyt clonal line was transfected with the cDNA encoding human proPTHrP(1–141) under the control of a constitutive CMV promoter. Figure 8a demonstrates that basal PTHrP secretion in vitro was greater in two different TBRIIΔcyt + PTHrP clones compared with the TBRIIΔcyt + pcDNA3.1zeo control. PTHrP secretion did not increase in response to TGF-β in any clone expressing the dominant–negative type II receptor (Fig. 8a). Next, the effect of PTHrP overexpression by MDA/ TBRIIΔcyt was studied in vivo. Thirty-one days after tumor inoculation, mice bearing either TBRIIΔcyt + PTHrP clone had significantly larger and more osteolytic lesions on radiographs compared with those bearing TBRIIΔcyt + pcDNA3.1zeo tumors (Fig. 8, b and c). At the time of sacrifice, plasma PTHrP concentrations (Fig. 8d) were significantly greater in both TBRIIΔcyt + PTHrP groups compared with the TBRIIΔcyt + pcDNA3.1zeo control group, as were blood ionized calcium concentrations (data not shown).

Discussion

Breast cancer metastasizes to bone in greater than 80% of patients with advanced disease and causes local bone destruction (2) with subsequent pain, fracture, hyper-
calcemia, and nerve compression syndromes. The data presented here suggest a central role for TGF-β in the pathogenesis of breast cancer metastasis to bone. Breast cancer cells that metastasizes to bone stimulate osteoclastic bone resorption (5–7, 29) and release of active TGF-β into the bone microenvironment (4). This increase in locally active TGF-β alters tumor-cell behavior to promote growth and bone destruction at the metastatic site. Thus, in the situation of breast cancer metastases to bone, it appears that host-derived TGF-β acts on the tumor cells, via a receptor-mediated mechanism, to endow metastatic capacity rather than a case of tumor-derived TGF-β acting on the mesenchyme to favor invasion. In the results reported here, TGF-β has the paradoxical effect of enhancing metastasis and bone destruction while inhibiting tumor-cell growth in vitro.

Dominant-negative blockade of the type II TGF-β receptor had no effect on cell growth in vitro in the absence of TGF-β or on cell growth of tumor cells inoculated intramuscularly, but it significantly decreased tumor growth in bone. In contrast, expression of the dominant-negative type II TGF-β receptor blocked the growth inhibitory effects of TGF-β in vitro. On the basis of the growth results in vitro, one might expect the MDA/TßRIIΔcyt cells to grow more rapidly in bone, when, in fact, these cells had a much slower growth rate in bone compared with the controls.

It has been suggested that breast cancer cells may progress from a growth-inhibited to a growth-stimulated response to TGF-β (30). Previous work in breast cancer has focused on the growth inhibitory effects of TGF-β, as exemplified by the findings that TGF-β1 suppresses mammary tumorigenesis in mouse mammary tumor virus/TGF-β1 transgenic mice (31). Furthermore, expression of a type II TGF-β receptor in the human breast cancer cell line, MCF-7, diminished tumorigenicity (32), whereas transgenic mice overexpressing a dominant-negative mutant type II TGF-β receptor had enhanced tumorigenesis in the mammary gland and lung in response to the carcinogen 7,12-dimethylbenz[a]-
Anthracene (33). The data presented here, however, suggest that the effects of TGF-β unrelated to growth inhibition in the context of tumor-host interaction can negatively affect the host. This paradigm goes against the simple idea that losing responsiveness to TGF-β is universally permissive for transformation. In fact, the results suggest that maintenance of TGF-β receptors on the tumor cells can adversely affect the host by an indirect effect on the metastatic process.

Although loss of TGF-β receptor function (34) or its signaling molecules (35–38) has been associated with malignant progression (28), there is growing evidence that TGF-β may enhance tumor growth and invasion. In a transgenic mouse model in which TGF-β1 expression was targeted to keratinocytes, TGF-β1 had a biphasic action during skin carcinogenesis by acting early as a tumor suppressor and later by enhancing a malignant phenotype (39). TGF-β has also been shown to induce an epithelial-mesenchymal transdifferentiation to an invasive phenotype (40, 41). Mammary epithelial cells transformed by Ras can become resistant to growth inhibition by TGF-β (42). Furthermore, in these cells, TGF-β enhances invasion and tumorigenesis by inducing a highly motile fibroblastoid phenotype. A possible basis for this is the ability of Ras-activated Erk to phosphorylate and inhibit Smad proteins (43). An oncogenic Ras mutation has been reported in MDA-MB-231 breast cancer cells (44). This may explain why these cells are only moderately growth-inhibited by TGF-β in vitro, as well as why cells expressing the constitutively active type I TGF-β receptor develop severe bone metastases, a phenotype that indicates excessive tumor growth in bone.

The effects of TGF-β in cancer may be tissue-specific. In recent work by Böttiger et al. (45), targeted expression of a dominant-negative type II TGF-β receptor in pancreas and liver resulted in pancreatic carcinoma without liver abnormalities. The results presented here are also consistent with a tissue-specific role of TGF-β in malignancy. In the situation of breast cancer metastasis to bone, TGF-β, released and activated as a result of tumor-stimulated osteoclastic bone resorption, is an important segment of a paracrine loop that may be responsible for the affinity with which breast cancer grows in bone. In tumor cells with oncogenic Ras mutations, TGF-β may promote further tumor development. The data presented here also suggest the possibility that PTHrP may be an effector of TGF-β in bone metastases, because overexpression of PTHrP into the breast cancer cells that expressed the dominant-negative TGF-β type II receptor resulted in accelerated bone metastases. The effect of TGF-β on tumor cells to stimulate PTHrP may result in adverse effects only when tumor cells are housed in bone rather than in soft tissue sites. Tumor cells in the bone microenvironment produce PTHrP and stimulate osteoclastic bone resorption, which in turn results in the release of active TGF-β. TGF-β then acts on the tumor cells to endow them with metastatic capacity and the ability to stimulate production of PTHrP. The net result is tumor growth, bone destruction, fracture, and the complications of osteolytic bone metastases.

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