

PROSPECTS

Modulation of Prostate Cancer Growth in Bone Microenvironments

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Abstract Bone remains one of the major sites, and most lethal host organs, for prostate cancer metastasis. Prostate cell spread and establishment in bone depends on multiple reciprocal modifications of bone stromal and epithelial cancer cell behaviors. This review focuses on recent advances in the characterization of cell–cell and cell–matrix interplay, effects on cell growth, adhesion and invasion, and several therapeutic possibilities for co-targeting prostate cancer cells and bone stroma. We address the topic from three main perspectives: (1) the normal and aging bone stromal environment, (2) the “reactive” bone stromal environment, and (3) the cancerous prostate epithelial cells themselves. First, normal, and especially aging, bones provide uniquely rich and “fertile soil” for roaming cancer cells. The interactions between prostate cancer cells and insoluble extracellular matrices, soluble growth factors, and/or sex steroid hormones trigger bone remodeling, through increased osteoclastogenesis and further matrix metalloproteinase activity. Second, after cancer cell arrival and establishment in the bone, host stromal cells respond, becoming “reactive” in a process again involving extracellular matrix remodeling, together with growth factor and steroid receptor signaling this process ultimately enhances cancer cell migration, stromal transdifferentiation, and invasion of the cancer tissues by stromal, inflammatory, and immune-responsive cells. Third, prostate cancer cells also respond to supportive bone microenvironments, where soluble and matrix-associated molecules affect cancer cell growth and gene expression, especially altering cancer cell surface receptor and integrin-mediated cell signaling. We discuss both integrin cell–matrix and gap junctional cell–cell communication between cancer cells and their microenvironments during prostate cancer progression. *J. Cell. Biochem.* 91: 686–705, 2004. © 2003 Wiley-Liss, Inc.

Key words: prostate cancer bone metastasis; stromal–epithelial interaction; androgen-independent progression; osteoblast; osteoclast and bone re-modeling; reactive stroma; gap junctional communication; growth factors; extracellular matrices; growth factor receptors; integrin profile; connexin

Like many other metastasizing cancer cells, cancerous prostate epithelial cells interact with multiple distinct organ microenvironments, including the surrounding prostate gland before and during their escape, the lymphatic and

vascular endothelia during transit, and the bone upon establishment within the skeleton. To successfully treat patients with prostate cancer bone metastasis and its associated complications, such as spinal cord compression, pathological bone fractures, chronic pain, anemia, and infection, a better understanding is needed of the interactions between prostate cancer cells and their environments, especially those of the bone. This review focuses on the recent characterization of this cell biology, in particular bone regulation of prostate cancer cell growth, adhesion and invasion, as well as the reciprocal effects of the stromal environment on cancer cells.

It has long been observed that tumors of different origins appear predisposed to metastasize to different organs, depending primarily on the circulatory system to gain access. In the 1800s, cancer cells were detected in the blood

Grant sponsor: NIH/NCI; Grant numbers: CA-76620, CA098912; Grant sponsor: NASA grant (to LWC); Grant number: NCC89-144; Grant sponsor: American Cancer Society IRG-90-016-11, NCC9-144 and Georgia Cancer Coalition (grants to ME).

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Received 25 August 2003; Accepted 2 September 2003

DOI 10.1002/jcb.10702

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stream [Cruveilhier, 1829; Ashworth, 1869] and identified as metastatic emboli in the 1950s [Engell, 1955; Roberts et al., 1958]. As early as 1889, Dr. Stephan Paget [Paget, 1889] examined the autopsy records of 735 women who died of breast cancer and proposed the popular “seed and soil hypothesis,” arguing that the tumor cells (the seeds) had specific affinities for the growth milieu provided by certain organs (the soil). More recent experiments have revealed that distinct locations of metastases can be predicted and traced following injection of melanoma cells into mice [Fidler and G, 1976]. The skeleton is fertile ground for many tumor types. Approximately 95–100% of myelomas, 65–75% of breast and prostate cancers, 60% of thyroid cancer, and 15–45% of bladder, lung, and renal cancers and melanomas end up in bone [Cifuentes and Pickren, 1979]. When considering skeletal metastasis, researchers have focused on such topics as: the tropism of cancer cells to bone, the roles played by the bone marrow’s reticulate vasculature, its provision of uniquely rich oxygen, nutrients and survival factors to tumor cells, the routes taken by heterogeneous cancerous cell types, and their migration and invasion into marrow stroma, and finally, subtle differences between bone microenvironments. Prostate cancer cells may employ a unique portal-like venous drainage system between the prostate and the lower spine [Bastón, 1940; Resnick, 1992; Bubendorf et al., 2000]. The size and frequency of metastatic tumors found in 1,589 prostate cancer autopsies revealed that not only do spine metastases precede those to lung and liver, but there is an upward metastatic spread along spinal veins after initial lumbar metastatic deposits are established [Bubendorf et al., 2000]. Although many metastases remain unpredictable, suggesting complex and multi-step mechanisms for cancer cell spread and establishment [Liotta and Kohn, 2001], bone remains the major organ for prostate cancer metastasis. This review is subdivided into three parts: (1) the normal and aging bone stromal environment as a “rich soil,” (2) the reactive, or activated bone stromal environment and its effects on the interplay between prostate cancer and bone stromal cells, and (3) the responsive character of prostate cancer cells, themselves, especially in terms of integrin-mediated cell–matrix, and gap junction-mediated cell–cell communications.

NORMAL AND AGING BONE STROMA: A RICH SOIL FOR CANCER CELLS

Bone heterogeneity and the complexity of microenvironments are key to an understanding of restriction and selectivity in cancer cell spread within a non-uniform environment. Prostate cancer cells are known to “prefer” trabecular bone, also called spongy or cancellous bone, a bone type composed of a network of tiny strands of bone trabeculae. Such bone is found at the ends of long bone, in ribs, in the pelvis, vertebrae, and the skull and is interspersed with marrow and a rich vasculature. For successful metastasis, the cells of trabecular bones must attract cancerous prostate cells, allow them entry into the tissue, and provide optimal growth conditions once inside.

Entry into the bone marrow is generally believed to depend upon the arrest of circulating cancer cells in the vascular beds, with single cells arresting in the capillaries, while multicellular aggregates arrest in larger vessels [Liotta et al., 1976]. Recent evidence argues that only endothelium-attached cancer cells are able to give rise to metastases [Al-Mehdi et al., 2000]. Cell “docking” on the vasculature involves recognition of protein receptors, such as selectins, integrins, cadherins, and immunoglobulin superfamily members. These proteins may be expressed initially by the cancer and/or vascular endothelial cells in response to disease progression and local cues, such as growth factors, cytokines, chemokines, or reactive oxygen species. Along with the heterotypic cell interactions between cancer cells and endothelia, homotypic interactions within multicellular cancer cell aggregates also occur. In several instances, *in vivo* selection for tumor cells of high metastatic potential has resulted in the selection of cells with increased homotypic aggregation properties [Saiki et al., 1991; Glinsky and Glinsky, 1996]. This correlation appears to depend upon selectins and β -galactoside-mediated adhesive interactions, at the sites of primary attachment to the microvascular endothelia [McEver, 1997; Glinsky et al., 2003]. Beyond specific cell adhesions, bone endothelial sites may also be favored for biomechanical reasons. Unlike other bones, the vascular beds within trabecular bones form sinusoids of large diameter and reduced blood flow rate (up to ten times slower), possibly allowing increased cancer cell attachment to

vessel surfaces, and their access across only a few microns into the stromal marrow compartment itself [Schnitzer et al., 1982]. This unique vasculature may help to resolve conflicting results from calculations of the times required for sufficient cancer cell attachments to endothelia prior to appearance in the bone [Haier and Nicolson, 2001a,b].

A prostate epithelial cells ability to intercalate among bone cell's may be of greater importance than the strength of their initial adhesion; *in vitro*, prostate carcinoma cells preferentially interact with bone marrow endothelial cells (HBME), adhering to HBME over umbilical vein, aortic, dermal, or lung endothelial cells (HUVEC, HAEC-1, HDMVEC, and Hs888Lu, respectively) [Lehr and Pienta, 1998; Scott et al., 2001; Sikes et al., 2003 submitted]. Shortly after prostate (and breast) cancer cells relocate to vessel walls, they express factors that are known to stimulate endothelial retractions and are believed to induce "reactive" stroma. One such factor is osteonectin. This small Ca^{2+} ion-binding glycoprotein is normally expressed by osteoblasts, endothelial cells, and megakaryocytes [Rodan and Noda, 1991; Kelm et al., 1992; Thomas et al., 2000; Brekken and Sage, 2001; Lin et al., 2001]. Cancer cell expression of osteonectin is involved in Human Epidermal growth factor Receptor 2 (HER2, also referred to as Her2/neu or c-erbB2) tissue remodeling and the breaching of vessel walls [Sanchez et al., 2002; Holbro et al., 2003]. Porous vessel walls give the cancer cells access to both chemotactic factors from the underlying tissue compartments and new extracellular matrices.

Once attached, cancer cells encounter varying levels of chemotactic, adhesion, and growth factors provided by marrow stromal cells [Hart, 1982; Liotta and Kohn, 2001]. Not only do prostate cancer cells adhere to bone marrow endothelial cells, osteoblasts, and prostate stromal fibroblasts [Haq et al., 1992; Lehr and Pienta, 1998], but media conditioned by these cells can stimulate prostate cancer cell growth [Lang et al., 1995]. Further evidence that bone stromal cells may prepare their soil with unique, even species-specific "fertilizers" comes from Tsingotjidou et al. [2001], who demonstrated that human prostate cancer cells home to experimentally implanted human bone, rather than to the mouse bone within their athymic mouse hosts. The identities and roles of

the released factors responsible for such homing are not yet clear [Tsingotjidou et al., 2001].

A recurring statement in the literature is that bone matrix is composed of more than 90% type I collagen fibrils secreted by osteoblasts, but bone is a dynamic tissue, constantly being remodeled and altered through both local and systemic cues. Maintenance of compartmentalization and multiple delicate balances in collagenous and non-collagenous components of the bone extracellular matrix are key to tissue homeostasis, bone remodeling, and disease, but have proven difficult to study, as they depend upon many cell types and regulation by cytokines, growth factors, and matrix metalloproteinases. Isotype-specific collagen antibodies have revealed that collagens are compartmentalized. Although collagens I, III, and X are all matrix-localized, collagen III is the most abundant in the marrow [Reddi et al., 1977], where it is produced by cells of fibroblast lineage [Castro-Malaspina et al., 1980]. Type IV collagen is produced by endothelial cells [Jaffe et al., 1976] and is found in the basement membranes of the vascular sinuses [Gay et al., 1984], where the endothelial cells of the vessel walls are separated from the adventitial cells by the collagen IV and laminin basement membrane [Inoue and Osmond, 2001]. Likewise, non-collagenous matrix components, including fibronectin, vitronectin, laminin, and bone sialoprotein, may also be compartmentalized and shift levels or localization during bone remodeling or cancerous invasion. For example, laminin 5 (LM-5) protein occurs in a novel bone-specific isoform (with the chains $\alpha 5$, $\beta 2$, and $\gamma 2$) in multiple splice versions, and is markedly absent in cancerous prostate cells before their establishment in bone [Siler et al., 2000, 2002; Hao et al., 2001].

Much of the bone matrix is produced by osteoblasts, whose activities are closely coupled to those of the osteoclasts and invading cancer cells. *In vitro* osteoblasts are known to be able in the presence of ascorbic acid and β -glycerophosphate, to form a mineralized bone matrix closely resembling woven bone [Quarles et al., 1992]. Metastasizing prostate cancer cells, once inside the bone, actually mimic osteoblasts, not only in their synthesis and deposition of non-collagenous bone matrix proteins (including osteopontin, osteonectin, osteocalcin, and bone sialoprotein) [Koeneman et al., 2000], but also by participating directly in osteoclastogenesis

and the formation of mineralized bone. This ability to take part in osteoclastogenesis has been linked to prostate epithelial cell expression of RANKL (receptor activator of NF- κ B ligand) and sRANKL (soluble RANKL), both of which are capable of activating osteoclasts through the cell surface receptor, RANK [Boyle et al., 2003; Lee et al., 2003]. Activated osteoclasts degrade the bone, releasing a number of crucial growth factors and cytokines, including TGF- β , IGF-1, and IL-6, and directly or indirectly promoting proliferation of both osteoblasts and prostate cancer cells [Brown et al., 2001a,b]. Together, invading prostate cancer cell matrix deposition and growth factor regulation are likely to have profound effects on cell adhesion, motility, proliferation, and survival.

Given that bone stromal-derived growth factors affect cancer cell response to chemokines, in a trans-activation process [Mira et al., 2001], a better understanding is needed of both systemic and local cues for bone remodeling, as well as shifts in bone microenvironmental composition, especially in the context of aging bone. It is generally accepted that bone turnover rates and bone loss are accelerated in response to shifts of hormone level, but there are differences in bone loss at different skeletal sites and between cortical and trabecular bone. Aging is accompanied by a reduction in both trabecular bone volume and numbers of endosteal and osteocyte cells. Additionally, hematopoietic tissue decreases, while fat cells increase. Although there is great variability in the scale of these changes, depending upon both sex and age, two mechanisms involving matrix proteins have been outlined to explain them. The first is based upon the extracellular matrix becoming more rigid, due to cross-linking of matrix proteins and changes in surface charge, either by glycation or by oxidation of lysyl and hydroxy lysyl residues to aldehydes [Miyata et al., 1997; Bailey et al., 1998; Hadley et al., 1998]. These changes are accompanied by degradation of the collagen and other matrix proteins [Termine, 1990]. The second mechanism involves decreased production of the most common matrix proteins, as seen in cells from donors of increasing age [Fedarko et al., 1990, 1992, 1995]. Such decreases are likely to alter the soluble growth factor and cytokine cues normally contained within the matrix.

The intersection between research findings on hormones, bone remodeling, aging, and prostate cancer is complex, but rich, especially

because of well known hormonal effects on extracellular matrix composition, growth factors, and cytokines. Balances of estrogenic and androgenic actions on bone have been studied in animal models, where estrogen functions to reduce longitudinal growth rate, whereas androgens actually stimulate chondrocyte maturation and metaphyseal ossification [Lebovitz and Eisenbarth, 1975]. Although androgen deficiency can reduce these processes, androgen elevations are also able to reduce growth, possibly indirectly, through aromatization to form estrogen [Iannotti, 1990; Hermann et al., 2000]. The increased estrogen levels observed in obese patients, together with altered metabolism in the fat cells, likely support trans-differentiation of prostatic fibroblasts into smooth muscle cells, an important event early in the development of benign prostatic hyperplasia (BPH), among aging males. Increases in femoral fracture incidence, which also correlate with aging, may be at least partly attributable to shifts in the rates of bone turnover, although changes in bone mass are not as well documented [Boonen et al., 1997]. While some have correlated androgen levels inversely with bone loss, others have disputed these findings and claim that bone mass is unrelated to serum testosterone and adrenal androgens.

In addition to circulating sex steroids, some cytokines directly regulate bone cell behavior, while others appear to act indirectly through growth factors. Shifts in both cytokine and growth factor levels are known to accompany aging. For example, interleukin-6 (IL-6) levels increase in the serum of older individuals, particularly after menopause or andropause. IL-6 not only regulates inflammation, but has been implicated in rheumatoid arthritis, osteoporosis, multiple sclerosis, and most notably has been identified as a direct regulator of osteoblast function [Gimble et al., 1994; Jay et al., 1996]. Other factors linked to aging and changing estrogen levels are interleukin-1 (IL-1) and tumor necrosis factor (TNF- α). IL-1 and TNF- α are among the most powerful stimulators of bone resorption known, and well-recognized inhibitors of bone formation. However, systemic levels should not be considered the only measures of a regulator's effectiveness. Local concentrations need not necessarily change as systemic levels do, and osteoblast responses to growth regulators may shift as well. Such is the case for osteoblast response to insulin-like

growth factor-1 (IGF-1). IGF-1 receptor levels do not change, but osteoblast response to the growth factor decreases with age [Pfeilschifter et al., 1993, 2000; D'Avis et al., 1997; Ankrom et al., 1998]. In this way, age-associated shifts in any given factor could be either further enhanced or ameliorated quite locally [Rosen, 2000].

TUMOR-ASSOCIATED STROMA: ACTIVE PARTICIPANTS IN CANCER PROGRESSION

Stromal cells are now known to actively modify cancer cell behavior, instead of providing only passive support for local cancer growth. In fact, cancer and stromal cells reciprocally modify each other's microenvironments, using feedback cycles to support cell growth and differentiation, during both normal development and disease [Chiquet-Ehrismann et al., 1986; Yee et al., 1989; Basset et al., 1993; Wright et al., 1994; Singer et al., 1995]. In the LNCaP human prostate cancer progression model, we demonstrated a change in cell phenotype from marginally tumorigenic behavior to tumorigenic and bone metastatic behavior, following cell following cell co-culture with other human prostate or bone stromal cells in vitro [Rhee et al., 2001] and in vivo [Thalmann et al., 1994]. Organ-specific stromal cell treatment can irreversibly determine genomic organization of cancer epithelia, resulting in cancer cell progression to androgen independence and increased bone metastatic potential [Thalmann et al., 1994, 2000].

Fibroblasts and smooth muscle stromal cells in the human prostate gland synthesize both the structural and regulatory components of the extracellular matrix, forming a meshwork of fibrillar, adhesive glycoproteins, and proteoglycans. This matrix serves as a reservoir of active and latent growth factors [Tuxhorn et al., 2001], such that stromal cells are greatly affected by matrix remodeling during cancer progression. In prostate and other cancers, stromal cells at the invasion fronts have been shown to change both morphologically and biochemically, converting from fibroblasts to myofibroblasts, and expressing vimentin, versican, hyaluronic acid, MMP2, MMP9, fibroblast activation protein (FAP), pro-collagen Type I, and tenascin [Tuxhorn et al., 2002a,b,c]. These changes are termed stromal activation, and correlate negatively with patient survival [Tuxhorn et al., 2001, 2002a]. Conversion to myofibroblasts

occurs in colon, liver, lung, breast, pancreas, and prostate cancers, where it is localized to the invasion fronts [Miura et al., 1993; Neaud et al., 1995; Ronnov-Jessen et al., 1995; Rowley, 1998; Doucet et al., 2000; Lohr et al., 2001]. In breast cancer, myofibroblast progenitors have been identified as fibroblasts (100%), vascular smooth muscle cells (40%), and pericytes (10%) [Bissell et al., 1999; Bissell and Radisky, 2001]. In the last case, pericytes would need to migrate from the blood vessel basement membrane to the interstitial collagenous stroma [Ronnov-Jessen et al., 1995].

Thus, when activated, stromal cell populations change both phenotypically and genotypically [Moinfar et al., 2000], but questions remain about how many of the stromal myofibroblasts are endogenous (but altered by association with the cancer cells) and how many are "recruited" from other cellular compartments. A three-dimensional co-culture study in our program, using multiple stromal cell lines (the human osteosarcoma cell line MG-63, a human prostatic stromal cell line, and the marginally tumorigenic LNCaP cell line), resulted in permanent, non-random genotypic changes of the LNCaP cells, together with associated phenotypic changes in tumorigenicity and metastatic potentials [Rhee et al., 2001]. The reciprocal is also true, as androgen-independent, metastatic C4-2 tumor cells can genotypically and phenotypically alter co-cultured stromal cells (Sung et al., unpublished observation). These in vitro studies effectively argue for transdifferentiation of pre-existing, tumor-associated stroma, rather than cell recruitment during cancer progression.

Molecular Regulation of Stromal Conversion

Myofibroblast conversion in primary prostate tissue can be regulated in vivo and in vitro by transforming growth factor β 1 (TGF- β 1) [Tuxhorn et al., 2002a,b,c], with necessary TGF- β 1 concentrations decreasing as prostate cancer cell malignancy status increases [Tuxhorn et al., 2002a,b,c]. As a candidate regulator for the genesis of reactive stroma at Prostatic Intraepithelial Neoplasia (PIN) sites, TGF- β 1 increase in high-grade PIN lesions [Tuxhorn et al., 2002a,b,c]. TGF- β 1 regulates the stromal response to wound repair, a response commonly compared to tumor progression (see below), and is also known to regulate angiogenesis. When TGF- β 1 expression is blocked, angiogenesis

decreases dramatically [Tuxhorn et al., 2002a,b,c]. This effect may be linked to myofibroblast expression of vascular endothelial growth factor (VEGF), in response to hypoxia or other growth factor release [Orlandini and Oliviero, 2001]. In fact, TGF- β 1 is known to act not only with VEGF, but also platelet derived growth factor (PDGF), and fibroblast growth factor (bFGF) to drive the reactive stromal response during cancer progression [Tuxhorn et al., 2001; Sung and Chung, 2002; De Wever and Mareel, 2003]. TGF- β 1 is secreted as a latent complex that is subsequently activated by proteinase cleavage, involving furin, plasmin, cathepsin, human mast cell chymase, leucocyte elastase, matrix metalloproteinases MMP2, MMP9, MT1-MMP, $\alpha_v\beta_6$, and $\alpha_v\beta_8$ integrins, thrombospondin 1, and environmental pH change [Lyons et al., 1990; Taipale et al., 1995; Crawford et al., 1998; Munger et al., 1999; Yu and Stamenkovic, 2000; Mu et al., 2002].

Because TGF- β 1 receptors are down-regulated in carcinoma cells, but remain high in stromal cells, TGF- β 1 is thought to act directly on stroma [De Wever and Mareel, 2003]. Despite the many factors that have now been shown to act during TGF- β 1 regulation of stroma, downstream responses within the stromal cells are less well understood. TGF- β 1 treatment does appear to affect the smads pathways. Smads are named after *C. elegans* Sma and *Drosophila* Mad proteins, the first identified members of this class of signaling effectors, and are known to increase production of fibronectin, plasminogen activator I, and cyclin-dependent kinase (CDK) inhibitors [De Wever and Mareel, 2003]. The TGF- β 1 signal pathway is also linked to the MAPK pathways, which in turn affect other regulators, including extracellular regulated kinase (ERK), c-jun amino-terminal kinase (JNK), and p38 [Akhurst and Balmain, 1999; ten Dijke and Heldin, 1999].

The Shifting Balance Between Osteoblasts and Osteoclasts

Bone extracellular matrices contain many embedded factors that are released when the balance of osteoblast and osteoclast functions is shifted and the matrix remodeled. Prostate cancer skeletal metastases are characterized primarily as osteoblastic with an underlying osteoclastic component [Keller, 2002]. Advances in the understanding of osteoclastogen-

esis and prostate cancer bone metastasis have defined several promising therapeutic targets for reduction of tumor-induced osteolysis. One factor found to be important in tumor-induced promotion of osteoclast activity is the RANKL protein (mentioned above; Table I). The soluble form of this protein (sRANKL) is required during normal osteoclastogenesis, but both RANKL and sRANKL are produced by prostate cancer cells within the bone, where they activate osteoclasts and induce osteolysis. Another factor, osteoprotegerin, a soluble decoy receptor for RANKL, inhibits RANKL-induced osteoclastogenesis [Zhang et al., 2001], and in murine models, has been found to inhibit tumor-induced osteolysis. Parathyroid hormone-related protein and interleukin-6 (IL-6) are also produced by prostate cancer cells and can promote osteoclastogenesis (Table I). Other key factors orchestrating stromal reactions are TGF- β 1 [Tuxhorn et al., 2002a,b,c], platelet-derived growth factor (PDGF); [Roodman, 2003], and factors responsible for activation of latent TGF- β 1 (such as matrix metalloproteinases) or co-signaling within the TGF- β 1 pathway (such as Wnt-catenin and Ras); [Rowley, 1998]. Drugs that specifically inhibit TGF- β 1's activities are particularly promising; these include soluble TGF- β 1 antagonists, TGF-receptor antagonists, inhibitors of TGF- β 1 intracellular signaling, and metalloproteinase inhibitors that block latent TGF- β 1 activation [De Wever and Mareel, 2003].

Matrix metalloproteinases (MMPs) warrant special attention in the context of bone remodeling, as these zinc-dependent proteinases are secreted by prostate cancer cells and are known to promote osteolysis by degrading bone matrix [Nemeth et al., 2002]. MMP2 and MMP9 have both been associated with prostate cancer [Dong et al., 2001; Nemeth et al., 2002] and are found at increased levels in the blood plasma and urine of patients with prostate metastases [Gohji et al., 1998; Moses et al., 1998]. Membrane-bound proteinase MT1-MMP levels also increase during progression from benign prostate disease to PIN, and further to cancer [Upadhyay et al., 1999; Nagakawa et al., 2000; Udayakumar et al., 2003]. MT1-MMP is involved in the proteinase activation of pro-MMP-2, pro-collagen 1, and laminin [Ohuchi et al., 1997; Udayakumar et al., 2003]. Matrix metalloproteinases are also active during osteoclast recruitment to sites of

TABLE I. Secreted Factors From Prostate Epithelial Cells That Stimulate Bone Remodeling by Modulating Osteoblast and Osteoclast Activity

Factors	Bone formation	Bone resorption
Bone resorption		
sRANKL		+(1)
Osteoprotegerin (OPG)		+(1)
IL-1		+(2)
Cathepsin K		+(3)
Collagen I amino-terminal telopeptide (NTX)		+(4)
α -Collagen I carboxy terminal telopeptide (α -CTX)		+(4)
β -Collagen I carboxy terminal telopeptide (β -CTX)		+(4)
Collagen I cross-linked carboxy terminal telopeptide (ICTP)		+(4)
TNF- α		+(2)
ET-1		-(5)
Bone formation		
Galectin-1	+(6)	
Cyclophilin A	+(7)	
BMP-2	+(2)	
BMP-4	+(2)	
BMP-6	+(8)	
Bone alkaline phosphatase (BAP)	+(4)	
Pro-collagen I amino-terminal propeptide (PINP)	+(4)	
ET-1	+(9)	
PTHrP	+(10)	
IL-6	+(2, 11)	
Placenta bone morphogenic protein (PLAB)	+(12)	
Bone sialoprotein (BSP)	+(13)	
Osteoprotegerin (OPG)	+(2)	
IGFBP-3	-(14)	

(1) [Brown et al., 2001a,b], (2) [Lee et al., 2003], (3) [Brubaker et al., 2003], (4) [de la Piedra et al., 2003], (5) [Chiao et al., 2000], (6) [Andersen et al., 2003a], (7) [Andersen et al., 2003b], (8) [Autzen et al., 1998], (9) [Guise et al., 2003], (10) [Dougherty et al., 1999], (11) [Garcia-Moreno et al., 2002], (12) [Thomas et al., 2001], (13) [Waltregny et al., 2000], (14) [Fizazi et al., 2003].

bone remodeling [Keller, 2002; Nemeth et al., 2002].

Both mineralized and non-mineralized bone matrices are affected by a variety of osteoclast-associated enzymes, including cysteine proteinases and MMPs. A number of synthetic MMP inhibitors are able to slow bone resorption in bone culture systems, including chemical inhibitors of MMP-2 and MMP-9 [Hill et al., 1994, 1995]. If a “vicious cycle” is initiated during prostate cancer metastasis, such that cancer cells stimulate bone matrix turnover, which in turn stimulates cancer growth [Nemeth et al., 2002], MMPs may make ideal therapeutic targets for the cycle’s disruption. In murine models, MMP inhibitors have been shown to diminish tumor establishment in bone. The regulation of MMP activation and activity is strongly dependent on the levels of Tissue Inhibitor of Metalloproteinases (TIMPs); [Liotta et al., 1991; Nagase, 1997; Brew et al., 2000]. Hence, balance between proteinases and inhibitors may determine the net enzymatic activity present in the tissue. Significant down-regulation of TIMP1 and TIMP2 correlate with the induction of pro-MMP9 expression, in co-culture of prostate cancer and stroma [Dong et al., 2001];

this effect is likely due to the stromal cells, as they are the major producers of TIMPs. Cancerous progression could, then, be supported by both increased MMP expression and decreased TIMPs, and the resultant changes in MMP concentrations and half-lives. Other promoters of osteolysis, derived from prostate cancer metastases, may now also be considered as new therapeutic targets [Dong et al., 2001].

Stromal Invasion of Epithelia: Wounds That do not Heal

During organogenesis and normal maintenance of adult organs, stromal and inflammatory cells intercalate among epithelial cells, regulating morphogenesis and cytodifferentiation [Ronnov-Jessen et al., 1996; Bissell and Radisky, 2001]. As active participants in carcinogenesis, stroma can: (1) increase deposition of extracellular matrices [Sung and Chung, 2002], (2) recruit other “reactive” stromal fibroblasts or myofibroblasts to the tumor [Tuxhorn et al., 2002a,b,c; Zidar et al., 2002], and (3) alter microenvironments adjacent to the tumor, through inflammatory and immune-responsive cell secretion of cyto- and chemokines [Svennevig, 1980; Heinrich et al., 2003; Wang et al.,

2003]. Because this sequence of events mimics classic wound-healing cascades (albeit without normal levels of apoptosis), cancers have been referred to as “wounds that do not heal” [Tuxhorn et al., 2001, 2002a]. In this context, the stromal reaction to tumor epithelia could be viewed as a host defense mechanism, initially intended to curtail or restrict tumor expansion. Sadly, the stromal reaction ultimately increases stromal cell number, alters stromal differentiation, and produces the extracellular matrices, growth factors, and metalloproteinases of a uniquely “fertile soil” for the support of tumor cell growth and invasion.

THE CANCER CELL: A MOVING TARGET

Not only the bone stromal cells, but also the prostate cancer cells change irreversibly during disease progression and metastasis. Because of feedback loops, it is difficult to separate early soil changes from those in the seeds. As human prostate carcinoma cells change phenotype they actively participate in bone turnover, thus affecting both the stroma and their own subsequent regulation. Attempted time-courses and causal relationships are often unclear on whether shifts in protein expression by the prostate cells occur before extravasation or are the consequence of tumor microenvironment interaction at the new location. To obtain a true developmental time-course, one would have to

analyze the circulating prostate cells in patients before metastases occur [Ts'o et al., 1997; Wang et al., 2000; Ellis et al., 2003] and compare expression profiles of prostate cancer cells from primary tumors and marrow biopsy specimens. Unfortunately, such measurements have not yielded clinically significant results, suggesting that the location of the prostate cancer cells may not be as telling as the cells' specific phenotypes. For example, over 50% of all prostate patients with clinically localized disease have prostate cells in their bone marrow, but only 10–30% are expected to develop recurrence after radical prostatectomy, and even fewer to develop detectable skeletal metastases. Thus, it seems likely that proliferative cancer cell phenotypes principally develop *after* re-localization to the bone environment.

Adhesion to Extracellular Matrices at Sites of Metastasis

Cancer cell adhesion to new extracellular matrices depends upon a variety of cell membrane receptors, including the integrins. Previous studies of integrin expression in various epithelial carcinomas have found that increased levels of the $\alpha_v\beta_3$ integrin heterodimer correlate well with metastatic potential, but otherwise many published observations conflict. Table II shows that integrin expression can change transiently during disease progression; thus, some of the published discrepancies may not

TABLE II. Differences in Integrin Expression Between Cell Lines of Different Metastatic Potential

Integrin	Cell line DU-145	PC3	LNCaP	RWPE-1	Change in carcinoma	Change in metastasis
$\alpha_2\beta_1$		0 (2)	+(3)		–(6)	+(6)
$\alpha_3\beta_1$		–(2)	+(3)		0 (7)	
$\alpha_5\beta_1$		0 (2)				
$\alpha_6\beta_1$	+(1)		+(3)		0 (7–9); +(6)	+(6)
$\alpha_6\beta_4$	+(1)	+(2)	–(3)		–(8–11)	
$\alpha_v\beta_3$		0 (2)	+(3)		+(12)	
α_2			+(4)		–(7)	
α_3	0 (1)		+(4)		–(7)	
α_4					–(7)	
α_5	0 (1)		+(4)		–(7)	
α_6	+(1)		0 (4)	– (5)	0 (13); –(7)	
α_v					–(6)	
β_1	0 (1)		+(4)		0 (13,14); – (7); +(15)	
β_{1A}					0 (16)	
β_{1C}					–(14,16,17)	
β_2					0 (7)	
β_3					0 (7)	+(18)
β_4			+(4)		–(7,9,10,11,13)	

+, increases; 0, stable; –, decreases. (1) [Rabinovitz et al., 1995], (2) [Dedhar et al., 1993], (3) [Edlund et al., 2001], (4) [Freedland et al., 2003], (5) [Bello-DeOcampo et al., 2001], (6) [Bonkhoff et al., 1993], (7) [Cress et al., 1995], (8) [Nagle et al., 1994], (9) [Nagle et al., 1995], (10) [Allen et al., 1998], (11) [Davis et al., 2001], (12) [Zheng et al., 1999], (13) [Knox et al., 1994], (14) [Perlino et al., 2000], (15) [Murant et al., 1997], (16) [Fornaro et al., 1996], (17) [Fornaro et al., 1999], (18) [Hartstein et al., 1997].

only be attributed to differences in methodology and antibodies, but to cell stage. In prostate cancer research, one α integrin subunit, α_6 , has received the majority of attention. The α_6 subunit pairs with β_1 or with β_4 in hemidesmosomal structures. β_4 is not found in metastatic cells [Allen et al., 1998], whereas α_6 expression decreases more slowly during disease progression, and actually increases remarkably in lymph node metastasis [Bonkhoff et al., 1993]. This pattern is somewhat similar to that of α_2 , although decreases in α_2 are detectable at much earlier stages. Further complexity in protein pattern was revealed by findings in the LNCaP prostate cancer progression system; we discovered that as prostate cancer cells progress, differences in the regulation of integrin subunit usage and heterodimerization are much more marked than differences in the surface expression of most integrins [Edlund et al., 2001].

Unlike the other integrin heterodimers, expression and activation of the vitronectin receptor $\alpha_v\beta_3$ integrin has been clearly linked to the metastasis and progression of many cancers [Juliano and Varner, 1993], most specifically to cell migration through basement membranes, in studies using function-blocking antibodies *in vivo* [Brooks et al., 1994a,b]. $\alpha_v\beta_3$ integrin is also expressed by proliferating vascular endothelial cells during angiogenesis [Brooks et al., 1995; Drake et al., 1995]. Monoclonal antibodies or small molecular antagonists for $\alpha_v\beta_3$ are known to inhibit blood vessel formation (including tumor angiogenesis) in a variety of *in vivo* models [Eliceiri and Cheresch, 2000, 2001]. These promising results, however, are confounded by the finding that mice lacking both $\alpha_v\beta_3$ and $\alpha_v\beta_5$ not only support enhanced tumor growth, but also present increased neovascularization [Reynolds et al., 2002].

Despite probable redundancy during angiogenesis and extravasion, antibodies, peptides, and chemical compounds have all been tested as integrin antagonists, with clinically significant results in some cases [Gutheil et al., 2000]. Furthermore, small, non-peptide inhibitors of both $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrin heterodimers inhibit tumor angiogenesis [Kumar et al., 2001; Kumar, 2003], and in our hands have successfully blocked previously established tumor growth in the femur, when systemically delivered (unpublished observation). The actual mechanism behind integrin-based decreases in tumor growth remains unclear, but could be

direct regulation of cell proliferation and/or indirect regulation involving growth factors and angiogenesis. One intriguing indirect possibility is the cooperation between integrins and matrix metalloproteinases (MMP's). The binding of MMP-2 to integrin $\alpha_v\beta_3$ in invasive cells appears to act together with MT1-MMP to present activated MMP-2 on the cell surface [Hofmann et al., 2000]. Inhibitors used to block β_3 -integrin function also inhibit MMP-2 by decreasing the expression of MT1-MMP [Pasco et al., 2000].

In addition to the integrins, Ig-domain cell adhesion molecule L1 (L1CAM) is also expressed in neuroblastoma, melanoma, and several carcinomas [Gabrielsen et al., 1988; Linnemann et al., 1989; Reid and Hemperly, 1992; Pancook et al., 1997]. L1 was originally identified as a neural cell adhesion molecule involved in axon guidance and neuronal migration [Fransen et al., 1997, 1998; Kadmon and Altevogt, 1997; Hoffman, 1998; Bliss et al., 2000; Thelen et al., 2002]. L1 protein interacts directly with both $\alpha_v\beta_3$ integrin and the fibronectin receptor, integrin $\alpha_5\beta_1$, as well as the cytoskeletal linker protein ankyrin [Bennett and Chen, 2001]. When expressed in the seminal vesicles and prostate of the urogenital tract [Kujat et al., 1995], L1 is normally restricted to non-proliferating epithelial cells. However, gene expression profiling in transgenic mouse models implicates L1 in both prostate and breast cancer cell progression to metastasis [Gutwein et al., 2000; Calvo et al., 2002]. Similarly, the L1 expression profiles of commonly used prostate cancer cell lines also correlate with high metastatic potential (Dr. Chia-Ling Hsieh, personal communication). Although normal prostatic tissue does not detectably express L1, the protein is readily found by immuno-staining in the stromal matrices adjacent to prostate tumors. L1's functions are still unknown, but interestingly, ARCaP cell expression and shedding of L1 correlates with the cell line's ability to penetrate endothelial monolayers [Zhou et al., 1996]. L1 likely influences epithelial-endothelial interactions and/or decreases vessel wall rigidity.

Altered Extracellular Matrix

Once in new environments, cancer cells not only bind extra-cellular matrix differently, but are known to alter the extracellular matrix adjacent to prostate, breast, and colon carcinomas, a key behavior in cancer

progression [Redler and Lustig, 1968, 1970]. Distal and proximal stromal fibroblasts differ from one another in adhesion, migration, and growth factor receptor expression. Likewise, production of woven bone (with random, loosely packed collagen strands) increases in prostate cancer patients relative to that of lamellar bone [Boyce et al., 1999]. These changes are regulated by secretion of osteoblastic factors by the prostate cells, regulating the core binding factor-1 (Cbfa1), a transcription factor known to control expression of osteocalcin, bone sialoprotein (BSP), osteopontin, and type I collagen in vivo [Ducy, 2000; Karsenty, 2000; Lin et al., 2001; Yang et al., 2001]. In vitro expression of antisense oligo-nucleotides for Cbfa1 decreases expression of these same osteoblastic genes [Banerjee et al., 1997; Ducy et al., 1997]. A similar transcription factor interruption strategy may be applicable against prostate cancer because of the cells' osteomimicry [Koeneman et al., 2000].

Cbfa1 affects Map Kinase (MAPK) and cyclic AMP (cAMP) signaling pathways, which in turn feedback on Cbfa1 protein levels. MAPK appears to activate Cbfa1, while cAMP reduces Cbfa1 protein levels [Xiao et al., 1998; Tintut et al., 1999; Xiao et al., 2000]. Cbfa1's actions on the MAPK pathway may involve other known MAPK stimulatory factors, including epidermal growth factor (EGF), fibroblast growth factor (FGF), and integrin adhesion molecules. In vitro, Collagen I has been found to regulate Cbfa1 [Xiao et al., 1998]. Not only does long-term cell culture on collagen I substrata induce osteoblastic differentiation of bone marrow cells, but this induction is abolished by addition of integrin function-blocking, collagen-specific peptides. Regulation of integrin-collagen interactions may be quite complex, with cryptic sites in the collagen molecules becoming exposed during matrix remodeling [Petitclerc et al., 2000; Xu et al., 2001], together with altered secretion levels of collagens I and III [Klein, 1995; Billiard et al., 2003].

Once cancer cells enter the bone microenvironment, one of the mechanisms used by cancer cells to modify their surroundings and regulate their own phenotype may be the shedding of membrane vesicles. Shed membrane vesicles are implicated in cancer cell escape from immune responses, the induction of angiogenesis, and the spread of metalloproteinase activity [Wood et al., 1997; Kim et al., 2002]. Membrane

vesicles, derived of specific regions of the plasma membrane [van Blitterswijk et al., 1979, 1982; Lerner et al., 1983] and enriched in tumor cell surface antigens are shed both in vitro and in vivo. Interestingly, vesicle quantities in patient serum appear to correlate with cell invasiveness and tumor load [Dolo et al., 1994, 1995, 1998; Ginestra et al., 1998, 1999; Dolo et al., 1999].

Chemotactic Factors

In addition to osteonectin and osteopontin (the bone matrix protein scaffolds bound by $\alpha_v\beta_3$ integrins during migration; van der Flier and Sonnenberg, 2001), chemokines are small molecular weight cytokine-like peptides known to affect cytoskeletal arrangement and induce directional cell migration. Many chemokines are secreted, and more than 20 receptors identified. The 50 chemokines so far identified in humans are all structurally related, mostly basic, small (8–14 kDa) molecules, whose receptors form a large family of seven-pass transmembrane, G protein-coupled molecules. These receptors show considerable overlap in ligand specificity, and act in a complex regulatory network [Muller et al., 2001; Taichman et al., 2002; Pfitzenmaier et al., 2003; Sun et al., 2003]. Chemokines were originally found to act during the directed movement of leukocytes across the endothelial layers of blood vessels into tissue [Muller et al., 2001]. Both bone and lung stromal cells secrete the chemokine CXCL12, whose receptor CXCR4 is expressed on the cell surfaces of many breast tumor cells. Antibodies against this CXCR4 receptor are able to significantly reduce breast cell metastasis to regional lymph nodes and lung [Muller et al., 2001]. Expression levels of both CXCL12 (also called chemokine stromal factor-1, SDF-1) and its receptor were found to be elevated in localized and metastatic cancers [Sun et al., 2003], where such chemotactic factors could play key roles in directing cancer cell relocation. Further studies are needed to assess whether levels of the same chemotactic factors that stimulate metastasis to the bone are altered physiologically during aging.

Gap Junctional Communications Among and Between Epithelial and Stromal Cells

Assessments of the mechanisms of bi-directional cell-cell communication—from stromal cell to cancer cell and cancer cell back to stromal cell [Gleave et al., 1991; Chung, 1993, 1995;

Sokoloff et al., 1996; Rowley, 1998; Tuxhorn et al., 2001, 2002a; Sung and Chung, 2002], have focused primarily on extracellular matrix protein expression, growth factor secretion, angiogenesis stimulation, and altered cell proliferation and survival [Rowley et al., 1994; Gregoire and Lieubeau, 1995; Martin et al., 1996; Noel and Foidart, 1998; Rowley, 1998; Tuxhorn et al., 2001, 2002a,b]. Gap junctions between cells, and the restrictive, direct cytoplasmic transfer they allow, should not be overlooked as uniquely powerful players in the reciprocal cell–cell communication that accompanies cancer progression.

Gap junctions are formed of connexin molecules linked between opposing cells. Electron microscopy has revealed their existence in most tissues [Shivers and McVicar, 1995], including human bone [Cancelas et al., 2000; Durig et al., 2000; Tuxhorn et al., 2002a,b,c]. If cancer treatments are most effective when aimed at both stromal cells (soil) and cancerous prostate epithelial cells (seeds), then gap junctions between and among these cell compartments deserve special attention for therapy. Indeed, connexins have recently been found to be involved in the “bystander effect” in gene therapy experiments [Carystinos et al., 1999, 2001]. Gap junctional connections are not only important for normal cell growth regulation [White et al., 1995; Nicholson and Bruzzone, 1997], but connexin expression levels negatively correlate with cancer progression; that is, down-regulation of connexin proteins accompanies increased metastatic potential [Trosko and Ruch, 1998; Sulkowski et al., 1999; Carystinos et al., 2001; Naus, 2002; Trosko, 2003]. Levels of connexins 26, 32, and 43 are all known to decrease during cancer progression [Grossman et al., 1994; Tsuda et al., 1995; Gee et al., 2003], and some connexins have even been found to function as tumor suppressor genes [Cunha et al., 2002]. Furthermore, when certain connexins are overexpressed, a decrease in tumorigenicity has been observed [Hirschi et al., 1996; Mehta et al., 1999]. However, little is known about subtle cancerous changes in connexin usage, and even less about the direct mechanisms by which connexins affect the development of primary tumors or tumors at distant sites. Most metastases retain some connexin expression, raising the possibility that these proteins are involved in intercellular communication between cancer cells

and the cells in their new habitat. Heterotypic gap junction formation is possible, both between connexins of different isoforms and between cell types [Bruzzone et al., 1996; Goodenough and Paul, 2003]. Heterotypic gap junctional communication does occur in both the bone marrow and hematopoietic tissues, and multiple connexins are known to be involved both during bone development and tissue homeostasis, in particular connexin 43 [Steinberg et al., 1994; Lecanda et al., 1998, Lecanda et al., 2000a,b]. In prostate cancer, several connexins have been analyzed, although with mixed results, as only a portion of the cells express connexins. Some researchers have detected connexins 32 and 40 in normal prostate, but not connexin 43 [Mehta et al., 1996; Kucuk et al., 2001]. Others have found decreased expression of connexins 32 and 43 in prostatic carcinoma [Tsai et al., 1996; Hossain et al., 1999a,b,c; Habermann et al., 2001], suggesting that they are still present in the metastatic tumors.

We are not aware of any studies correlating androgen levels, connexin expression, and tumor invasion and metastasis. Evidence that androgens regulate the expression of the connexin 43 gene in prostate tissue comes from studies in normal and castrated rats, where castration is associated with a dramatic and specific increase in connexin 43 mRNA and protein expression, and is correlated with increased apoptosis [Huynh et al., 2001]. Supplementing castrated animals with testosterone or DHT reverses this process. Thus, in castrated hosts, it is conceivable that connexins may actually play a role in increasing the ability of tumor cells to invade bone tissues, particularly through elevations of connexin 43 in men treated with androgen deprivation therapy.

CONCLUDING REMARKS

Recently, across cancer research fields, the interplay between cancer cells and their micro-environments has enjoyed increased attention, as reflected in the topic’s designated status as an “extraordinary opportunity” by America’s National Cancer Institute. Evidence is accumulating that prostate cancer depends especially strongly upon shifts in cell environment, or that acquired, rather than inherited, genetic alterations are key to its development and progression. At the time of disease diagnosis, prostate cancer is multifocal and heterogeneous,

with an average of five apparently independent cancer lesions and numerous other high-grade PIN lesions [Bastacky et al., 1995], and with each focus and lesion presenting separate genetic constitutions. Furthermore, the associations between prostate cancer development and diet, lifestyle, and/or levels of sex steroid are, in fact, stronger than those for sporadic and even familial forms of prostate cancer [Haenszel and Kurihara, 1968; Shimizu et al., 1991], although certain inherited genetic loci have been shown to contribute to familial prostate cancer [Ahlbom et al., 1997; Page et al., 1997; Lichtenstein et al., 2000].

Three future developments, having to do with tumor–microenvironment interactions, are particularly needed. First, improved methodologies for cell culture and in vivo molecular imaging will allow better recapitulation of normal development and cancer–stroma interplay, as well as imaging of proteins during cancer cell growth and locomotion. Second, the development of additional relevant cancer cell lines (including luminal and basal epithelial cell lines, as well as fibroblast, smooth muscle, myofibroblast, endothelial, and inflammatory cell lines), each representing different stages of prostate cancer development and progression, would provide invaluable tools for dissecting the features and requirements of invasive prostate cancer, and for differentiating between prostate cancers based on their degree of virulence. Third, molecular profiling of the signaling networks associated with soluble growth factors, insoluble extracellular matrices, and sex steroid hormones will aid in the identification, testing, and validation of selective anti-neoplastic agents.

Given the power of the microenvironment to direct prostate cancer progression, and the known preference of prostate cancer cells for the bone, we have reviewed here recent studies that address shifting prostate and bone stromal cell–cell and cell–matrix interactions during cancer progression. Increased knowledge of normal and aging bone stroma, reactive bone stroma, and the prostate cancer cell response to bone stroma, will ultimately reveal opportunities for improved therapy in the prevention and treatment of prostate cancer metastasis.

ACKNOWLEDGMENTS

The authors thank Amanda Tate for editorial and graphic help.

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