

Mammary gland tumor formation in transgenic mice overexpressing stromelysin-1

Carolyn J. Sympson†, Mina J. Bissell* and Zena Werb†*

*Life Sciences Division, Lawrence Berkeley Laboratory
Berkeley, CA 94720

†Laboratory of Radiobiology and Environmental Health and Department of Anatomy,
University of California, San Francisco, CA 94143-0750

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Proteinases have been shown to play a role in tissue remodeling during the normal processes of embryo implantation, wound healing, ovulation, and involution of the uterus, prostate and mammary gland (for review see ref 1). Extracellular matrix (ECM)-degrading proteinases have also been implicated in pathological processes such as rheumatoid arthritis and in all stages of tumor progression, including growth, angiogenesis, invasion and metastasis (for review see ref 2). In order for tumor cells to spread to other parts of the body, they must gain entry to the stroma, where they penetrate the lymph nodes or blood vessels (intravasation) and thereby the systemic circulation. Once carried to a distant site from the primary tumor, surviving cells may exit the circulation (extravasation) into the target organ and form a secondary tumor colony (metastases). ECM-degrading proteinases have been implicated in the intra- and extravasation of tumor cells across basement membranes (BM) (for review see refs 3,4). A variety of ECM-degrading enzymes, including matrix metalloproteinases and serine proteinases, have been shown to be induced or activated in tumor cells.⁴⁻⁶

We have recently described the production and characterization of transgenic mice that overexpress the matrix metalloproteinase stromelysin-1 (SL-1) in the mammary gland.⁷ The mammary gland of prepubertal female mouse is essentially quiescent. However, with the onset of ovarian activity (between 20–25 days), primary ducts, once confined to the immediate area of the teat, form large, club-shaped terminal end buds and infiltrate the fat pad. These ducts usually arrive at the end of the fat pad at about day 70 (mature virgin), and the gland remains dormant until pregnancy. Examination of glands taken from normal 70-day virgin females revealed the typical branching pattern (Figure 1A). However, the mammary glands from SL-1 transgenic mice, which express very low levels of the transgene, showed precocious alveolar development and maturation, with primary and secondary ducts that filled the fat pad (Figure 1C). This phenotype is both morphologically and functionally (as measured by their ability to express α -casein) very similar to a normal 9–12-day pregnant gland. An intermediate phenotype was captured at day 50 of virgin transgenic development (Figure 1B). The right side of the panel represents the most mature part of the gland, near the teat structure, which displays alveolar development similar to the transgenic gland in panel C. However, the most immature part of the gland, on the left side of the panel, resembles the normal virgin gland that is still branching as evidenced by the large end buds.

The mechanism by which SL-1 acts as a morphogen, leading to hyperbranching, proliferation and differentiation, has not been elucidated but is currently under investigation. We have hypothesized three possibilities for how the proteolytic action of SL-1 could directly or indirectly result in a proliferative response: (a) generation or activation of a growth stimulator; (b) destruction of a growth inhibitor; or (c) release of growth factors sequestered by the BM. It is known from our own studies that an intact BM is inhibitory for growth of both murine and human mammary epithelial cells. In rodent epithelial cells, it was shown that BM strongly down-regulates the expression of transforming growth factor (TGF)- β and TGF- α .^{8,9} For normal human cells, which grow at the same rates as malignant cells on tissue culture plastic, embedding in BM will lead to a complete arrest of growth.^{10,11} Alternatively, the cleavage of BM molecules by ECM-degrading proteinases could unmask ECM molecules or generate fragments of ECM molecules that may in turn act as growth stimulators (for review see ref 12).

Proteinases appear to play a role in the branching morphogenesis of the salivary gland and kidney, as well as in capillary sprouting. Fukuda *et al*¹³ have shown that branching of the salivary gland organoids can be severely affected by the addition of exogenous collagenase, suggesting a role for matrix metalloproteinases in this process. Madin-Darby canine kidney (MDCK) epithelial cells can undergo branching morphogenesis in three dimensional collagen or fibrin gels.¹⁴ This morphogenic effect can be inhibited by the addition of exogenous serine proteinase inhibitors and stimulated by hepatocyte growth factor (HGF), also known as scatter factor. It has been shown that recombinant human HGF causes a dramatic increase in both urokinase-type plasminogen activator (uPA) and its receptor.¹⁵

The formation of capillaries requires focal dissolution of the BM, followed by the migration of endothelial cells, which then form blood vessels upon canalization. Pepper *et al*¹⁶ found that capillary formation is dependent upon the balance between uPA and its inhibitor, PAI-1. Angiogenic factors, basic fibroblast growth factor (bFGF) and TGF- β 1, stimulate this response by causing an increase in uPA expression. This proteinase activity may be focal, because endothelial cells have functional uPA receptors, and the number of these receptors also increases upon stimulation with bFGF.^{17,18} Using amniotic invasion assays, others have found that invasion of bFGF stimulated bovine endothelial cell invasion requires both plasmin and collagenase activity.¹⁹ To add to this ever-growing cascade of events, others have shown that TGF- β 1-stimulated macrophages secrete increased levels of uPA, which in turn releases sequestered bFGF from the BM.²⁰ In addition, BMtethered TGF- β 1 has been shown to be released by the action of proteinases.²¹ It can be deduced from these experiments, using a variety of model systems, that extracellular proteolysis is most probably required for branching morphogenesis.

As the SL-1 transgenic mice progress through the pregnancy stage of mammary gland development, many of the alveoli appear collapsed and surrounded by a thin layer of BM.^{7,22} This morphological phenotype is accompanied by a decrease in the expression of milk-specific genes, β -casein and whey acidic protein. Further characterization of the mid-pregnant gland has revealed that the mammary epithelial cells are undergoing programmed cell death, as measured by in-situ labelling of fragmented DNA.²³ In-vitro evidence also suggests a role for an intact BM for cell survival and maintenance of tissue-specific morphology and function. These mice are capable of lactating even though the BM is severely disrupted and the alveoli are collapsed. Despite gross abnormalities during the growth and differentiation phases of the transgenic glands, the involution process appears to be normal.

We have now observed a dramatic incidence of tumors in four independent lines of these mice. All transgenic mice, thus far, display reactive stromas and phenotypic abnormalities ranging from severe hyperplasia, resembling a lactational phenotype, to adenocarcinomas (Figure 2). Hormonal stimulation does not appear to be absolutely necessary for tumor formation because tumors have developed in glands of virgin mice as young as four months old.

The mechanism by which disruption of the BM leads to tumor formation is not yet known. It is clear from our culture studies that the ECM is a potent inhibitor of growth and an important regulator of cell survival and tissue-specific differentiation of mouse mammary epithelial cells. The signal transduction pathway by ECM is only now being unravelled.²⁴ We have hypothesized

that the massive disruption of the BM could lead to induction of growth factors, leading to rapid proliferation of the epithelial cells and thus to increased genetic errors. This process in turn could give rise to a select population of cells that have a growth advantage. The availability of this breast tumor model should lead to an elucidation of the early steps in the malignant process and to an understanding of how an abnormal microenvironment in epithelial tissues could lead to cancer induction and progression.

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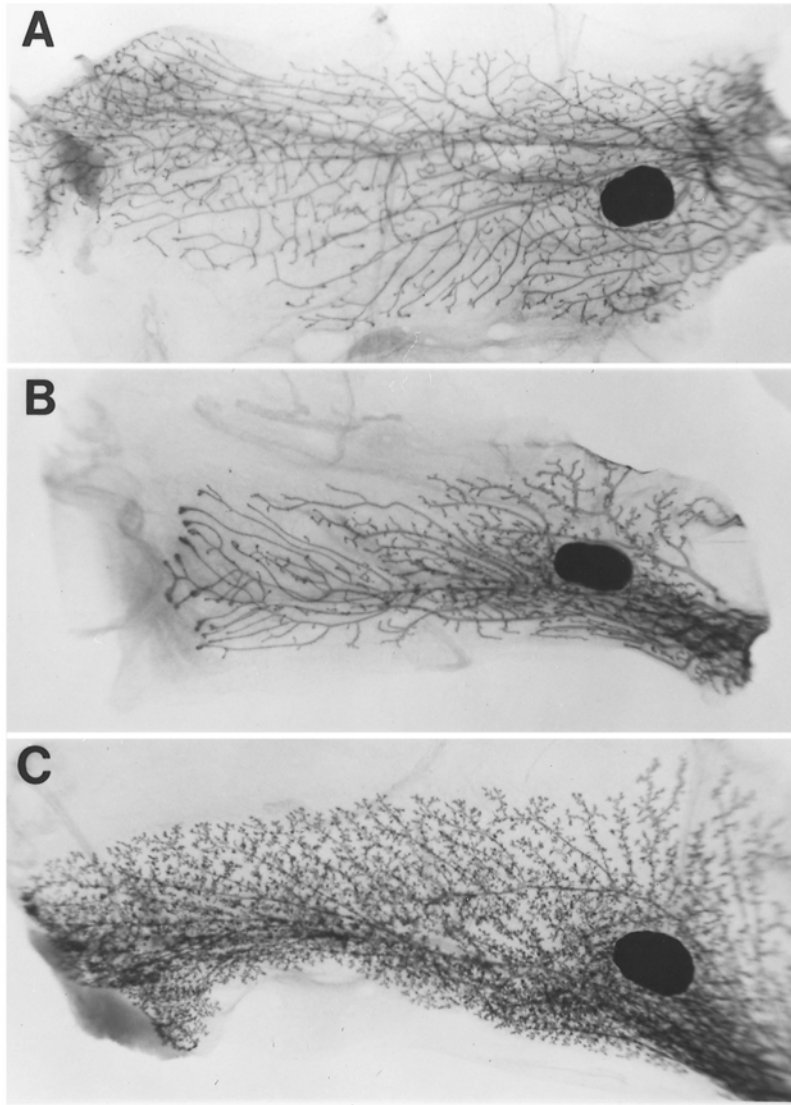
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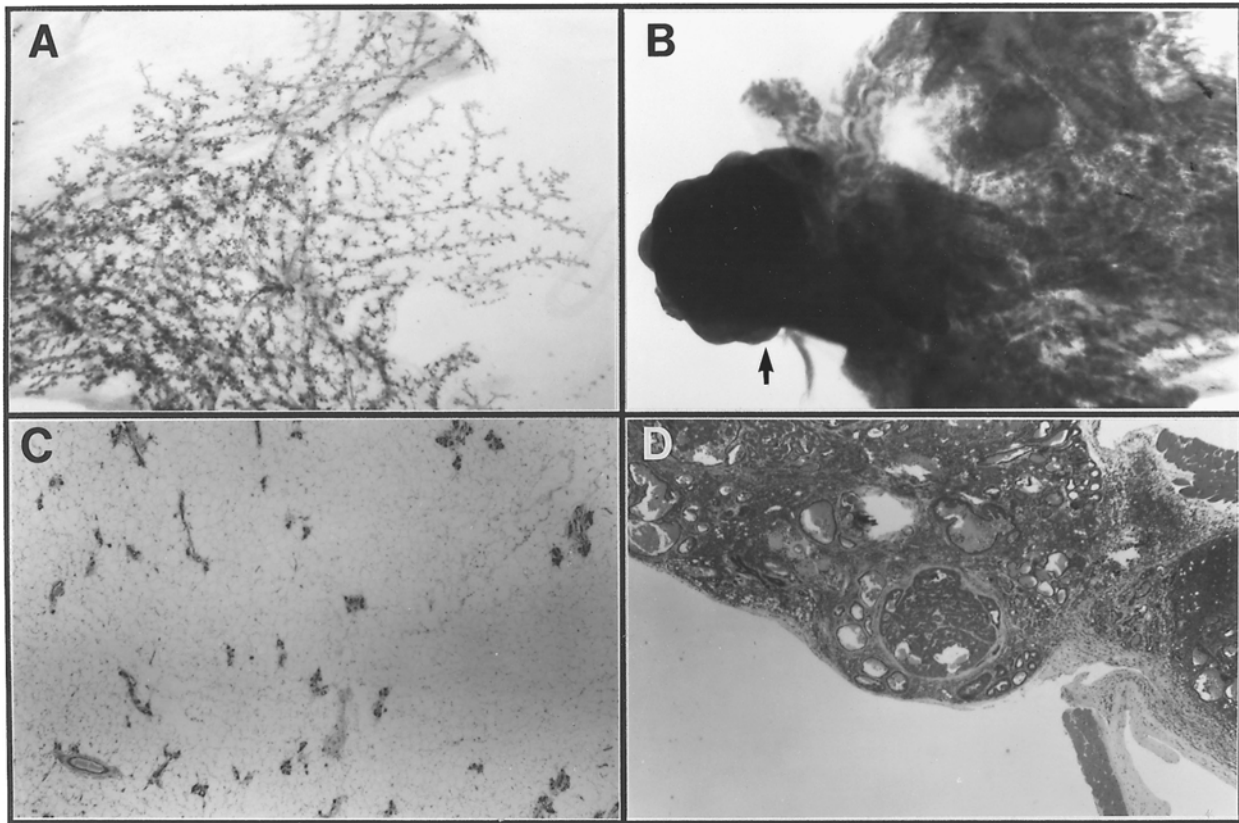
Figures

FIGURE 1



Phenotype of normal and transgenic mammary glands in virgin mice. Whole-mount carmine-stained glands from normal CD-1 mice: 70-day virgin (A), 50-day virgin transgenic (B), and 70-day virgin transgenic (C). The 70-day transgenic gland (C) is highly branched with prominent alveolar structures and closely resembles a normal 9–12-day pregnant gland. An intermediate phenotype can be captured at day 50 of virgin transgenic development (B). The darkstaining oval structures are the lymph nodes. Photographs are 6x magnification. See Simpson *et al*, 1994, for details.

FIGURE 2



Morphology of mammary glands from normal and SL-1-overexpressing transgenic mice. Whole-mount carmine-stained glands from normal (A) and transgenic tumor-bearing (B) mice and the corresponding hematoxylin-eosin-stained cross sections, C and D, respectively. The arrow in panel B corresponds to the enlarged cross-section in panel D, which demonstrates an encapsulated adenocarcinoma. Both glands are the thoracic (nos 2/3) taken 4 months after the first lactation. Photographs in A and B are 6x magnification and C and D are 40x magnification.