Parathyroid hormone–related protein (PTHrP) is a secreted factor expressed in almost all normal fetal and adult tissues. It is involved in a wide range of developmental and physiological processes, including serum calcium regulation. PTHrP is also associated with the progression of skeletal metastases, and its dysregulated expression in advanced cancers causes malignancy-associated hypercalcemia. Although PTHrP is frequently expressed by breast tumors and other solid cancers, its effects on tumor progression are unclear. Here, we demonstrate in mice pleiotropic involvement of PTHrP in key steps of breast cancer — it influences the initiation and progression of primary tumors and metastases. *Pthrp* ablation in the mammary epithelium of the PyMT-MMTV breast cancer mouse model caused a delay in primary tumor initiation, inhibited tumor progression, and reduced metastasis to distal sites. Mechanistically, it reduced expression of molecular markers of cell proliferation (Ki67) and angiogenesis (factor VIII), antiapoptotic factor Bcl-2, cell-cycle progression regulator cyclin D1, and survival factor AKT1. PTHrP also influenced expression of the adhesion factor CXCR4, and coexpression of PTHrP and CXCR4 was crucial for metastatic spread. Importantly, PTHrP-specific neutralizing antibodies slowed the progression and metastasis of human breast cancer xenografts. Our data identify what we believe to be new functions for PTHrP in several key steps of breast cancer and suggest that PTHrP may constitute a novel target for therapeutic intervention. […]
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Introduction
Metastases to bone, lung, and other organs are common and catastrophic consequences of breast cancer progression; most patients do not die from the primary tumor, but because of cancerous invasion to distal sites (1, 2). Once breast cancer metastases are established in bone or lung, the condition is generally considered incurable. There is therefore an urgent need to improve current therapies that address cancer spread, and an ideal solution will target upstream signaling molecules to prevent compensatory mechanisms that can result from blockade of individual downstream signaling points (3, 4).

Parathyroid hormone–related protein (PTHrP, also referred to as parathyroid hormone–like protein [PTHLP]) is a secreted factor expressed in almost all normal fetal and adult tissues. The 13 N-terminal amino acids of PTHrP are highly homologous to those of parathyroid hormone (PTH), a characteristic that allows PTHrP to act through the type 1 PTH receptor (PTH1R) (5). The rest of the PTHrP amino acid sequence is unique, however, and confers to the molecule many properties resulting from signal transduction cascades and nuclear translocation distinct from those of PTH (6). PTHrP acts as an autocrine, paracrine, or intracrine factor in a wide range of developmental and physiological processes (7, 8), it has growth-promoting and antiapoptotic properties (6), and it plays a crucial role in the development of the mammary gland and skeleton (8–10). Of special interest is the association of PTHrP with oncogenic transformation; ref. 34) was used to generate animals with a Cre-loxP–mediated (35) hetero- or homozygous Pthrp gene ablation specifically targeted to the mammary epithelium (ME) (Supplemental Figure 1, A and B; supplemental material available online with this article; doi:10.1172/JCI46134DS1). All animals used in the present study were confirmed by marker analysis to possess more than 99% FVB/NJ background. In standard PyMT-MMTV animals, tumors appeared spontaneously, approximately 100% of these tumors expressed PTHrP (55 tumors tested by RT-PCR), and their PTHrP expression increased with age (Figure 1A). In contrast, in Pthrp-ablated animals, spontaneous breast tumors showed and gastric carcinomas (25, 26). Circulating levels of PTHrP generally correlate with the more advanced stages of cancer (20, 27–32), and PTHrP regulates the expression of several tumor-relevant genes (33). Despite the frequent association of PTHrP dysregulation with many tumor types, a precise and direct role for PTHrP in cancer development and progression has been difficult to prove, and its involvement in tumor initiation in vivo and in critical steps of malignant conversion is not clear.

Here, we demonstrate PTHrP implication in key steps of breast cancer initiation, progression, and metastasis. We show that PTHrP plays a major role in stimulation of breast tumor growth rates and metastatic spread to distal organs through its effects on several crucial control molecules, including prosurvival signal molecule AKT and chemokine receptor CXCR4.

Results
Pthrp ablation occurring after birth allows normal mammary development. To clarify the role of PTHrP in tumorigenesis, the human breast cancer mouse model PyMT-MMTV (where the mT oncogene drives oncogenic transformation; ref. 34) was used to generate animals with a Cre-loxP–mediated (35) hetero- or homozygous Pthrp gene ablation specifically targeted to the mammary epithelium (ME) (Supplemental Figure 1, A and B; supplemental material available online with this article; doi:10.1172/JCI46134DS1). All animals used in the present study were confirmed by marker analysis to possess more than 99% FVB/NJ background. In standard PyMT-MMTV animals, tumors appeared spontaneously, approximately 100% of these tumors expressed PTHrP (55 tumors tested by RT-PCR), and their PTHrP expression increased with age (Figure 1A). In contrast, in Pthrp-ablated animals, spontaneous breast tumors showed
Figure 1
Cre-mediated Pthrp ablation in ME allows normal mammary development. (A) Confocal images of IF staining with anti-PTHrP antibody in spontaneous breast tumors from standard PyMT mice and Western blot quantification showing increasing PTHrP expression in these tumors with respect to age. (B) Confocal images of IF staining for DAPI (blue) and PTHrP (green) in tumor tissues from control (Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-}) and heterozygous (Pthrp\textsuperscript{flox/+};Cre\textsuperscript{-}) transgenic animals showing an incremental reduction in PTHrP expression with allele ablation. Lower panel: Western blot quantification for PTHrP expression in tumors from the various genotypes. (C) Confocal images of IF staining with DAPI (blue), Cre recombinase (green), and PTHrP (red) showing colocalization of Cre and PTHrP. (D) Whole-mount staining analysis (Neutral Red) of mammary glands showing ductal outgrowth at 3, 5, and 7 weeks for control, heterozygous, and homozygous female virgin mice. (E) Western blot of PTHrP expression in mammary glands of virgin 7-week-old mice. Scale bars: 50 μm (A–C); 5 μm (D).

incremental reduction in PTHrP expression from Pthrp\textsuperscript{flox/+};Cre\textsuperscript{-} (control) to Pthrp\textsuperscript{flox/+};Cre\textsuperscript{-} (heterozygous) to Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} (homozygous) (Figure 1B). Pthrp\textsuperscript{WT/WT};Cre\textsuperscript{-} animals were generated to test potential artifactual side effects caused by expression of Cre recombinase, but showed no difference from other controls throughout all experiments.

When tumor cells were cultured in vitro, radioimmunometric assays for PTHrP in the conditioned medium indicated low residual expression of PTHrP in heterozygous ablated cells as follows: Cre\textsuperscript{-} control cells, 178.7 ± 33.6 pg/ml; Cre\textsuperscript{-} tumor-derived cells, 10.1 ± 2.3 pg/ml; mean ± SD, n = 10 and 13 mice, respectively. In tumor-bearing mice, circulating PTHrP was undetectable, and calcium serum concentrations were not significantly different between control (2.28 ± 0.39 mmol/l) and Pthrp- ablated mice (2.23 ± 0.23 mmol/l).

PTHrP expression in the normal gland was localized both in luminal epithelial cells and in myoepithelial cells (Supplemental Figure 2, A–C, and ref. 36). PTH1R expression was unaffected by Pthrp ablation (Supplemental Figure 3).

Immunofluorescence (IF) staining confirmed Cre expression in the luminal epithelium of control mice (Figure 1C). The Cre gene was expressed under the control of the MMTV-LTR terminal repeat (MMTV-LTR), and its expression was detectable from 6 days postpartum (37). Pthrp ablation therefore occurred after birth, and mammary glands from 3-, 5-, and 7-week-old virgin females presented normal ductal outgrowth as well as normal lactation capacity with no detectable differences among genotypes (Figure 1D). PTHrP in the mammary gland was substantially reduced in 7-week-old homozygous mice in comparison with control animals (Figure 1E). These data show that ME-targeted Pthrp deletion can be achieved without hindering normal mammary gland development.

Pthrp ablation delays breast cancer initiation and primary tumor progression. Tumor hyperplasia was detectable in the mammary gland of control animals as early as 27 days after birth, adenomas were visible by 35 days, and early adenocarcinomas were visible by day 45 (Figure 2A). In contrast, ablation of both Pthrp alleles significantly delayed tumor initiation (Figure 2A). Tumor growth over time was reduced in homozygous Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} animals in comparison with Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} and Pthrp\textsuperscript{WT/WT};Cre\textsuperscript{-} controls (Figure 2B), and tumor weight/mouse at sacrifice (13 weeks) was 70% lower in ablated animals than in controls (Figure 2C). While 100% of control mice presented palpable tumors (2-mm diameter) around day 55, heterozygous and homozygous animals reached this percentage by days 75 and 92, respectively (Figure 2D). Homozygous mice reached tumor size requiring humane sacrifice at a much later age than control and heterozygous mice (Figure 2E). Tumors from hetero- and homozygous mice were 50% and 75% smaller than their respective Cre\textsuperscript{-} controls, and there were 35% and 60% fewer tumors in heterozygous and homozygous mice at 13 weeks (Figure 3, A and B, and Supplemental Table 1). The expression of Cre was not involved in the changes observed in tumor delay, as demonstrated by the Pthrp\textsuperscript{WT/WT};Cre\textsuperscript{-} controls, which did not differ from the other controls (Figure 2, B–E, and Figure 3, A and B). These results show that ablation of the Pthrp gene in the ME of PyMT-MMTV mice significantly delays primary breast cancer initiation events, even before tumors reach a palpable stage, and reduces their subsequent growth.

More complete Pthrp ablation by Cre-carrying adenovirus further delays breast cancer initiation and progression. Because some residual PTHrP expression was detectable in approximately 20% of primary tumors from Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} animals (Figure 1B), a more complete Pthrp knockout was achieved by transfecting isolated cells from Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-}, Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-}, and Pthrp\textsuperscript{WT/WT};Cre\textsuperscript{-} tumors with an adeno virus containing a CreGFP sequence or with a control adenovirus containing GFP only. Transfected adenoGFP control or adenoCre\textsuperscript{-} cells purified by cell sorting were transplanted into the fourth mammary fat pad (MFP) of healthy FVB syngeneic mice (5 × 10\textsuperscript{6} cells). PTHrP expression was eliminated in the adenoCre\textsuperscript{-}-derived tumors (Figure 4, A and B), tumor volume was greatly decreased (Figure 4C), and tumor load at injection site was dramatically reduced (Figure 4, D and E). These results show that more stringent Pthrp ablation conditions enhance tumor inhibition.

Pthrp ablation modifies cell-cycle, apoptosis, and angiogenesis events. Decreasing PTHrP expression reduced Ki67 (cell proliferation), factor VIII (angiogenesis), and cyclin D1 staining (Figure 5, A and B). Propidium iodide flow cytometry analysis revealed that 53.7% ± 1.19% of control Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} remained in G0/G1 compared with 95.2% ± 3.70% of homozygous Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-}. In contrast, 18.34% ± 1.80% of control Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} and 8.10% ± 4.00% of homozygous Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} tumor cells were in S phase, an observation consistent with the cyclin D1 decrease (not shown). Importantly, in cultured cells isolated from tumors, cyclin D1 expression colocalized specifically with cells that escaped PTHrP ablation (Figure 5C), suggesting that in this system, PTHrP is crucial for cyclin D1 expression. Pthrp ablation was accompanied by an increase in TUNEL apoptotic staining in tumors and tumor-derived cultured cells (Figure 5D) and by a decrease in Bcl-2 expression (Figure 5B). Western blots showed no difference in expression levels for factor VIII, cyclin D1, and Bcl-2 between Pthrp\textsuperscript{flox/flox};Cre\textsuperscript{-} and Pthrp\textsuperscript{WT/WT};Cre\textsuperscript{-} controls (Supplemental Figure 4), indicating that decreases in signaling molecule levels can be attributed to Pthrp ablation. Overall, these data indicate a pleiotropic effect of PTHrP in cell-cycle, apoptosis, and angiogenesis events.

Pthrp is involved in CXCR4 and AKT expression control. Pthrp ablation was accompanied by inhibition of metastasis marker CXCR4 expression in primary breast tumors of the same size (controls: 13 weeks; homozygous animals: 18 weeks; Figure 6, A and B). Furthermore, CXCR4 in cells isolated from primary tumors appeared exclusively in cells that avoided Pthrp ablation (Figure 6B). Pthrp ablation also decreased Akt1 total protein and increased total Akt2 protein in tumor tissues (Figure 6, C and D), and...
most residual AKT1 colocated with residual PTHrP in Cre- cultured cells isolated from tumors (Figure 6E). Again, the expression of Cre was not involved in changes seen for AKT or CXCR4 (Supplemental Figure 4). Phosphorylation of AKT1 (Ser473) was also inhibited by Pthrp ablation (Figure 6F). siRNA experiments targeting AKT1 effectively decreased its levels in control (Cre-) and Pthrp-ablated (Cre+) tumor cells (Figure 7A). Cell proliferation was reduced by 38.9% ± 8.0% by Pthrp ablation alone (Figure 7B), by 46.7% ± 4.1% through AKT1 knockdown alone, and by 80.2% ± 2.2% in Pthrp-ablated cells combined with AKT1 siRNA. These data indicate that PTHrP is involved in the control of expression of survival molecules AKT1 and AKT2 and of the chemokine receptor CXCR4.

**Figure 2**
Pthrp ablation delays breast cancer initiation and progression. (A) H&E staining of breast tissues from control (Pthrp<sup>flox/flox</sup>;Cre<sup>-</sup>) and homozygous (Pthrp<sup>flox/flox</sup>;Cre<sup>+</sup>) animals at 27, 35, and 45 days. Hyperplasias (arrowheads) are not detectable in homozygous animals before 45 days. Scale bars: 200 μm. (B) Tumor volume over time for control (Pthrp<sup>flox/flox</sup>;Cre<sup>-</sup> and Pthrp<sup>WT</sup>;Cre<sup>+</sup>) and homozygous mice (Pthrp<sup>flox/flox</sup>;Cre<sup>+</sup>) showing that the delay in tumor growth is not related to the expression of the Cre gene. Inset: early time points. Values shown represent mean ± SD, n = 12 mice per group. ***P < 0.001. (C) Tumor weight per mouse at sacrifice (13 weeks) for control and homozygous animals. Values shown represent mean ± SD, n = 12 mice per group. ***P < 0.001. (D) Kaplan-Meier analysis of tumor onset for mice of all genotypes illustrating allelic effect for Pthrp ablation. (E) Kaplan-Meier analysis showing that control mice reach age requiring sacrifice much earlier than homozygous animals.

**PTHrP drives metastatic spread to peripheral blood, bone marrow, and lungs, and distal metastases comprise PTHrP/CXCR4 double-positive cells.** Cells from control tumors possessed more than twice the Matrigel invasiveness potential of cells from Pthrp-ablated mice (Figure 8A), and their motility after surface wounding was higher (Figure 8B). There were fewer circulating tumor cells in homozygous than in control animals (same-size tumors, controls: 13 weeks; homozygous: 18 weeks; Figure 8C). Tumor cells flushed from bone marrow were detectable only in control animals (average of 31.5 ± 3.4 cells per animal, n = 4; Figure 8D).

Similarly, the incremental decrease in PTHrP lowered the number of micro- and macroscopic metastatic lesions in lung tissues. At 13 weeks, 100% (45/45) of control mice presented lung metastases compared...
When tumor cells were transfected with adenCreGFP or adenGFP virus, purified, and implanted (5 × 10^5 cells) in the fourth MFP of syngeneic mice, same-size primary tumors in the MFP were achieved at 8 weeks in control, 11 weeks in heterozygous, and 16 weeks in homozygous mice (not shown). At those stages, lung metastases were present in 70% (16/23) of control, 55% (6/11) of heterozygous, and 0% (0/22) of homozygous mice (Figure 8, H and I).

Importantly, in the spontaneous PyMT tumor model, lung metastases cells in control and homozygous mice (same-size tumors, 13 weeks and 18 weeks, respectively) were almost uniformly PTHrP and CXCR4 positive (Figure 9), indicating that lung metastases were mainly derived from PTHrP-expressing tumor cells that escaped gene ablation (Supplemental Figure 5). These results point to a driving role for PTHrP in invasion and metastasis in this system and suggest that combined expression of PTHrP and CXCR4 is crucial to metastatic spread.

Anti-PTHrP neutralizing mAbs inhibit breast cancer progression in vitro and in vivo. mAbs were produced against human PTHrP and CXCR4 positive (Figure 9), indicating that lung metastases were mainly derived from PTHrP-expressing tumor cells that escaped gene ablation (Supplemental Figure 5). These results point to a driving role for PTHrP in invasion and metastasis in this system and suggest that combined expression of PTHrP and CXCR4 is crucial to metastatic spread.

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Discussion

To investigate the involvement of PTHrP in breast cancer initiation, progression, and metastasis, we constructed an ME-specific Pthrp knockout animal in the PyMT-MMTV transgenic mouse model of breast tumorigenesis, using mice homozygous for a floxed Pthrp allele (2 loxP sites flanking exon 4, which encodes most of the PTHrP protein) (38). The PyMT mouse is a model with complete penetrance and approximates the 4 identifiable stages of tumor progression observed in human breast tumors (hyperplasia, adenoma, early carcinoma, and late carcinoma). These stages are followed in PyMT animals by a high frequency of distal metastasis, and the morphological characteristics and expression of biomarkers parallel those in the human cancer process (39). We obtained mice presenting various degrees of ablation of the Pthrp gene, demonstrating that reduction of PTHrP expression in ME brings substantial delays in breast cancer initiation without affecting mammary development. In Pthrp-ablated animals, primary tumor hyperplasia was considerably delayed, and palpable tumors appeared much later and were smaller and fewer than in control mice. Our observations are in agreement with several clinical studies in which patients with tumors produc-
ing high PTHrP levels presented higher rates of metastasis and increased or earlier mortality (40–43). A recent report reached opposite conclusions, but relied on animal models with a very late-onset oncogenic system (neu based), which is more relevant to tumors arising in older animals (44), and used mice with a heterogeneous genetic background. The age at which oncogenic induction occurs and the host’s genetic background are 2 factors that significantly affect the biological behavior of tumors (45, 46) and likely account for the
discrepancy observed with our results. Clinical studies reporting a good prognostic value for PTHrP in breast cancer have concentrated on cohorts in which neo-adjuvant cases were eliminated, therefore focusing on the less aggressive cases (47). In contrast, our animal model is more relevant to early-onset, pregnancy-independent, highly aggressive breast tumorigenesis, corresponding to a category of patients likely to require neo-adjuvant treatment.

In our mouse model, Pthrp ablation was accompanied by a near-complete disappearance of the CXCR4 receptor, a member of the chemokine superfamily that regulates cell migration and target-
CXCR4 depletion were accompanied by a significant decrease in total levels of the prosurvival AKT1 isoform and by an increase in the AKT2 isoform. AKT2 ablation accelerates tumor induction, and the 2 AKT isoforms possess opposite oncogenic effects (59). In striking similarity with our observations, a knockout of Akt1 interferes with mammary tumor initiation and growth in PyMT-MMTV mice, without hindrance to mammary development (59), but it is difficult to speculate whether PTHrP acts through acute or adaptive effect on AKT and CXCR4 signaling. Pthrp ablation has repercussions on the cell cycle (G0/G1 to S transition, cyclin D1, and Ki67) and enhances apoptotic events. Similarly, in the PyMT-Akt1 knockout mice, the delay in tumorigenesis is related to inhibition of cell proliferation, to a decrease in Ki67 and cyclin D1, and to promotion of apoptosis (59). Interestingly, cotargeting Pthrp and Akt1 by gene ablation and siRNA techniques provides a stronger inhibition of proliferation over either blockade alone (Figure 7, A–B), suggesting the existence of PTHrP-independent signaling through AKT. This is supported by the presence of residual AKT1 in Pthrp-ablated cells (Figure 6E) and suggests that a combined therapeutic approach could provide enhanced efficacy.

The CXCR4/CXCL12 interaction drives metastatic progression by activation of angiogenesis, chemotaxis, adhesion, and invasion processes (50), and neutralization of CXCR4/CXCL12 action in vivo by antibodies (48), inhibitory peptide (60), or siRNA (61) attenuates metastases in mouse models. Here, Pthrp-targeted ablation also reduces the number of tumor cells in peripheral blood and bone marrow and delays metastasis in lungs, the preferred metastatic site in the PyMT-MMTV mouse. Importantly, the metastatic tumors that eventually develop in homozygous animals present a pattern of positive expression for both PTHrP and CXCR4 (55).

In breast and prostate cancer cells, the CXCR4 receptor/CXCL12 ligand interaction results in activation of the PI3K/AKT pathway, which drives cell growth and survival (56) and is a key signaling route for CXCR4 (50, 57). The PI3K/AKT pathway is also essential to the transmission of oncogenic signaling from the middle T oncoprotein and to development of mammary adenocarcinomas in the PyMT system (39, 58). Here, Pthrp ablation and consequent CXCR4 expression, with consequences to survival and metastasis. Since almost all residual expression of PTHrP and CXCR4 in homozygous primary tumors colocalizes to the same cells, PTHrP appears to be an important control for CXCR4 expression, with consequences to survival and metastasis. An involvement of PTH in CXCL12 production by osteoblasts (62) reinforces the concept that as a member of the PTH family, PTHrP plays a role in CXCR4/CXCL12 signaling.

Previous work with anti-PTHrP antibodies revealed inhibition of bone turnover and osteolytic metastases (63, 64). Here, treatment of nude mice bearing PTHrP-positive human breast cancer
xenografts with anti-PTHrP mAbs reduced both primary growth and lung metastatic tumor development, which suggests a PTHrP role distinct from bone turnover control and calcium homeostasis. Overall, our data demonstrate that in a model of early-onset, pregnancy-independent, highly aggressive breast tumorigenesis, PTHrP plays a tumor-promoting role in initiation, progression, and metastasis, in part through upregulation of the CXCR4 and AKT pathways (Figure 11). Autocrine, intracrine, and paracrine effects have been reported for PTHrP (8, 27, 65) and are likely involved in its pleiotropic cancer-promoting activities, since the PTH1 receptor is present throughout the tumor tissue and is unaffected by Pthrp ablation. In carcinomas, there is frequent alteration in the expression of many growth factors involved as autocrine and paracrine mediators of stromal-epithelial interactions (66). Our results support the idea that some PTHrP-mediated effects observed in vivo cannot be observed in vitro, which points to important tumor/stromal interactions that must be considered when examining PTHrP functions during breast cancer progression. Finally, the effective inhibition of PTHrP action in human cancer xenografts by our neutralizing antibodies constitutes a proof of principle for therapeutic consideration.

Methods

Conditional ablation of the Pthrp gene in mouse ME. C57BL/6 Pthrp floxed mice (Pthrp flox/flox) (38) were backcrossed 9 generations to FVB animals. Marker analysis (Jackson Laboratory) confirmed a more than 99% FVB/NJ background. Crossing these mice with PyMT-MMTV strain PyMT 634 (34) or MMTV-Cre (35) mice (FVB background, Cre7 type) produced PyMT-MMTV; Pthrp flox/flox and PyMT-MMTV; Pthrp flox/+ and MMTV-Cre; Pthrp flox/+ mice. Crossing these animals produced PyMT-MMTV; Pthrp flox/flox; Cre + (homozygous), PyMT-MMTV; Pthrp flox/+; Cre + (heterozygous), PyMT-MMTV; and Pthrp flox/+; Cre − (controls) mice.
Figure 10 
Anti-PTHrP neutralizing mAbs inhibit breast cancer progression in vitro and in vivo. (A) Proliferation in Matrigel (24 hours) of human MDA-MB-435 breast cancer cells showing growth-inhibition effect of neutralizing antibodies 158 and M45 in vitro. *P < 0.05. (B) Tumor volume over time after injection of MDA-MB-435 cells into the MFPS of BALB/c nu/nu mice and treatment with anti-PTHrP mAbs, showing the tumor-reducing effect in vivo. Data are expressed as means of 8 mice in each group. *P < 0.05; **P < 0.01. (C) IF confocal images of mammary tumors 6 weeks after injection of MDA-MB-435 cells in MFPS of nude mice showing decrease of CXCR4 (top panels) and AKT1 (bottom panels) in treated animals. Shown are DAPI (blue), CXCR4 (green), and AKT1 (red). (D) H&E staining of lung metastases 6 weeks after injection of MDA-MB-435 in MFPS. Treatment with anti-PTHrP mAbs reduces the size and numbers of lung metastases. (E–G) Fewer mice present lung metastases after treatment with either mAb, and the metastases are smaller and fewer in numbers in treated animals. Mean ± SEM. *P < 0.05 (E); *P = 0.013 (F); *P = 0.045 (G). (H) IF confocal images of lung metastases in nude mice injected with MDA-MB-435 cells treated (6 weeks) or not with anti-PTHrP mAbs. Lung metastases are CXCR4 positive irrespective of treatment. Shown are DAPI (blue) and CXCR4 (green). Scale bars: 100 μm (A); 50 μm (C and H); 200 μm (D).

Immunohistochemistry and IF. Antibodies used were as follows: mouse monoclonal N-terminal anti-PTHrP antibody (1-34, AE-0502, IDS), mouse monoclonal PTH1R antibody (Upstate), mouse monoclonal anti-Cre recombinase and anti-phospho-Akt1 (Ser473) (Abcam), goat anti–total Akt1, Akt2, and Ki67, mouse cyclin D1, Bcl-2, actin, and rabbit polyclonal antibodies for the PTH1 receptor, PTHrP1–38 C-terminal end, tubulin, anti-rabbit and anti-mouse peroxidase conjugates (Santa Cruz Biotechnology Inc.), Alexa Fluor 555–conjugated goat anti-rabbit IgG and Alexa Fluor 488–conjugated goat anti-mouse IgG (Invitrogen), guinea pig anti-mouse cytokeratin 8, donkey anti-guinea pig IgG CY3-labeled, and rabbit anti-mouse cytokeratin 5 (PROGEN Biotechnik GmbH).

Tumors and other tissues were fixed, embedded, sliced, and stained with H&E according to standard protocols. The SK-4100 kit (Vector) was used for immunohistochemistry (IHC), and slides were reacted for diamobenzidine and analyzed with a Leica DMR microscope and BIOQUANT Nova Prime software (Bioquant). IF staining was conducted on deparaffinized tissue sections or cells fixed on slides. Results were analyzed with an LSM 510 Meta confocal microscope (Carl Zeiss Microimaging).

Whole-mount staining. MFPS were removed, spread on a glass slide, air dried, fixed, delipidated, and stained with 0.5% Neutral Red (Sigma-Aldrich).

Western blotting. Proteins were extracted from tissues, and samples (30–50 μg) were fractionated by SDS-PAGE electrophoresis, transferred to PVDF membranes, reacted with primary and secondary antibodies, and developed by enhanced chemiluminescence according to standard methods. Whole mammary glands were prepared according to Ackler et al. (67).

Cultivation of mouse breast tumor cells from primary tumors. Tumors were harvested, minced, and incubated in 2.4 mg/ml collagenase B and dispase II (Roche) in DMEM (no FBS) at 37°C for 2 hours. Floating cells were washed, pelleted, resuspended, and propagated in complete DMEM.

Adenovirus vector. Tumor cells in culture were infected with a recombinant adenovirus vector (10^6 viral particles/ml) containing a GFP-Cre transgene for the Cre recombinase. Forty-eight hours later, cells were selected on a BD FACSaria II cell sorter. GFP-positive cells (1 × 10^6 cells) were injected into the fourth MFP of anesthetized 5-week-old syngeneic FVB mice (Charles River). Cell-cycle analysis, PTHrP, and calcium assays. Cultured cells were stained with propidium iodide and analyzed for cell cycle by flow cytometry (FACSCalibur). Levels of PTHrP in culture medium or serum were measured by 2-site immunoradiometric assay (Beckman-Coulter). Serum calcium was determined with a Synchron 67 Autoanalyzer (Beckman). siRNA transfection for Akt1 knockdown. Cells isolated from mammary tumors were transfected with siRNA Akt1 targeting sequences (s62216) as follows: sense: CUCAGAUCUACUUCAGArtt, antisense: UCUUG-GAAUGAGUACUUGAGg (bases in lower case are overhangs not corresponding to the original target sequence), and negative control (Ambion). Cell proliferation was assayed by hemocytometer counting.

Tumor cell detection in peripheral blood and in bone marrow. Blood was collected from sacrificed animals by cardiac puncture. Cytosin-concentrated samples were stained with the Epimet Epithelial Cell Kit (Micromet). Bone marrow flushed with PBS from tibias and femurs cut at the knee joint was trituated through a 25-gauge needle. Samples (2 tibias and 2 femurs/animal) were counted with a hemocytometer and spread onto Cytospin Immunoselect Adhesion Slides (Squarix). Tumor cells were detected with pan-cytokeratin mAb A45-B/B3 or cultured in DMEM with 10% FBS.

Matrigel invasion and wound-healing assays. MDA-MB-435 cells were tested for invasion in Matrigel for 20–24 hours with or without mAbs or IgG (10 μg/ml) according to standard methods. Invading cells were counted after paraformaldehyde fixation and hematoxylin staining. For motility assays, cells were grown on poly–l-lysine–coated plates (Millipore). The cell carpet was scraped and the growth of cells into the scraped area monitored.

Lung metastases measurements. Photomicrographs of H&E-stained slides of lung metastases were analyzed with ImageJ software (http://rsweb.nih.gov/ij/index.html) to assay metastatic surface/lung (n = 9 mice).

Anti-PTHrP P1-38, mAbs. We generated 2 PTHrP-specific mAbs against hPTHrP1–38 peptide: hybridomas 158 and M45 produced mAbs (subclasses IgG3 for mAb 158 and IgGx for mAb M45) with strong binding to hPTHrP1–38. The mAbs were highly specific (no reaction with PTH), and no cross-reactivity between antibodies and other fragments of PTHrP was observed. The 158 and M45 hybridomas have been deposited at the International Depositary Authority of Canada (accession numbers 060808-02 and 060808-01).

Figure 11 
PTHRP influences several key steps in breast cancer. Interactions are described here for PTHrP in tumor cell proliferation; through its effects on cell proliferation factor Ki67, cell-cycle progression regulator cyclin D1, and the G1/S to S transition, PTHrP is involved in very early steps of oncogenesis. PTHrP influences breast tumor cell survival, apoptosis, and angiogenesis through control of levels of expression for crucial signaling molecules such as AKT1/AKT2, Bcl-2, and factor VIII. Of great interest is the observation that PTHrP is involved in the control of CXCR4 expression and consequently also plays a role in metastatic spread.


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Treatment of nude mice with anti-PTHrP mAbs. 1 x 10^6 human cancer cells (MDA-MB-435) were injected into the fourth MFP of 4- to 5-week-old BALB/C nu/nu females (Charles River). Starting the next day, antibodies were injected subcutaneously (200 μg) every 48 hours for 6 weeks. Mouse IgG; isotype control was from R&D Systems. Primary tumor growth rates were determined by plotting the means of 2 orthogonal diameters of the tumor measured at 5-day intervals. Tumor volume was measured once weekly. Animals were sacrificed 6 weeks after tumor implantation.

PyMT-MMTV animal handling. Tumor-bearing animals were examined by palpation twice a week until they were 13 weeks of age. Mice with excessive tumor burden (> 1.5 cm) were euthanized. For Kaplan-Meier, animals were sacrificed when total tumor load exceeded 20% of animal weight. All experiments were carried out in compliance with regulations of the McGill University Institutional Animal Care Committee. All animal surgeries were conducted in accordance with principles and procedures dictated by the highest standards of humane animal care.

Statistical analysis of tumor progression and tumor growth. Numerical data are presented as the mean ± SD. The data were analyzed by ANOVA followed by a Bonferroni’s post-test to determine the statistical significance of differences. All statistical analyses were performed using Instat Software (GraphPad Software), and P < 0.05 was considered statistically significant. Specific tests are mentioned in the text.

Study approval. These animal studies were approved by the McGill University Animal Care Compliance Office.

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