

Endogenous human prolactin and not exogenous human prolactin induces estrogen receptor α and prolactin receptor expression and increases estrogen responsiveness in breast cancer cells[☆]

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Abstract

Prolactin (PRL) and estrogen act synergistically to increase mammary gland growth, development, and differentiation. Based on their roles in the normal gland, these hormones have been studied to determine their interactions in the development and progression of breast cancer. However, most studies have evaluated only endocrine PRL and did not take into account the recent discovery that PRL is synthesized by human mammary cells, permitting autocrine/paracrine activity. To examine the effects of this endogenous PRL, we engineered MCF7 cells to inducibly overexpress human prolactin (hPRL). Using this Tet-On MCF7hPRL cell line, we studied effects on cell growth, PRLR, ER α , and PgR levels, and estrogen target genes. Induced endogenous hPRL, but not exogenous hPRL, increased ER α levels as well as estrogen responsiveness in these cells, suggesting that effects on breast cancer development and progression by estrogen may be amplified by cross-regulation of ER α levels by endogenous hPRL. The long PRLR isoform was also upregulated by endogenous, but not exogenous PRL. This model will allow investigation of endogenous hPRL in mammary epithelial cells and will enable further dissection of PRL effects on other hormone signaling pathways to determine the role of PRL in breast cancer.

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Keywords: Prolactin; Prolactin receptor; Estrogen; Breast cancer

1. Introduction

In the normal mammary gland, the polypeptide hormone prolactin (PRL) and the steroid hormones, estrogen and progesterone, act synergistically to increase mammary gland growth, development, and differentiation (reviewed in [1–4]). Based on their roles in the normal gland, these hormones have been studied to determine their function in the development and progression of breast cancer. While the mitogenic activity of estrogen in mammary tumor cells has been well-established, appreciation of this activity for PRL is relatively recent [5–9]. Epidemiological studies that have evaluated circulating PRL levels in an attempt to show a correlation with breast cancer risk have been controversial.

While several studies suggest a positive correlation between circulating PRL levels and breast cancer risk factors (reviewed in [10]), drug therapies to reduce serum PRL levels, such as the use of bromocriptine, have been unsuccessful in treating the disease [11,12]. However, these studies evaluated only endocrine PRL and did not take into account the recent discovery that PRL is synthesized and secreted by human mammary cells, permitting autocrine/paracrine activity [13–15]. Study of this local PRL production is necessary to determine its contribution to breast cancer and potentially improve breast cancer treatment modalities.

Understanding regulation of the PRL receptor (PRLR) and the estrogen receptor (ER) within the mammary gland is key to determine how these hormones interact and exert their mitogenic effects. PRL has been shown to regulate responsiveness to ovarian hormones in several female reproductive tissues including the ovary and uterus, as well as the normal mammary gland [16,17]. In vivo studies of these relationships are complicated by indirect actions of PRL on both the ovary and the stromal compartment of the mammary gland, and have been shown to vary with genotype in the mouse [18]. PRLR^{-/-} and ER α ^{-/-} mice have defects in mammary

Abbreviations: hPRL, human prolactin; E2, 17 β -estradiol; iPRLR, long prolactin receptor isoform; iPRLR, intermediate prolactin receptor isoform; PgR, progesterone receptor; ER α , estrogen receptor α ; NIDDK, Institute of Diabetes and Digestive and Kidney Disease

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gland lobuloalveolar development, and ductal growth and differentiation, respectively [19,20]. Although levels of mammary ER α in PRLR $^{-/-}$ mice, and PRLR in ER α $^{-/-}$ mice have not been reported, the pattern of progesterone receptor (PgR) expression during mammary development in PRLR $^{-/-}$ mice was disrupted [2]. In a defined in vitro system, normal murine mammary epithelial cells responded to exogenous PRL with a small (25%) increase in ER and a three-fold increase in PgR [21]. However, despite the importance of ovarian hormone responsiveness in the design of therapeutic targets and prognostic indicators for breast cancer, the role of PRL in control of ER and PgR is less clear in mammary tumor cells. In primary neoplasms, several groups have described a positive correlation between ER and PgR, and PRLR levels [22–24]. However, other groups did not find a relationship [25–29]. In the breast cancer cell line, MCF7, Ormandy et al. did not see any effect of human prolactin (hPRL) treatment on ER mRNA levels or specific binding after 24 h, but did observe a small increase in PgR (<two-fold) [24]. In contrast, in an earlier report, Shafie and Brooks observed that hPRL induced a small increase (<two-fold) in ER in the same cell line, after 48 h [30]. Neither study addressed endogenous hPRL production, nor did they evaluate functional effects of PRL on estrogen responsiveness.

To examine the effects of endogenously synthesized mammary hPRL, we engineered MCF7 cells, an extensively studied hormone responsive breast cancer cell line, to inducibly overexpress hPRL. We used this Tet-On MCF7hPRL cell line to study effects of endogenous hPRL on cell growth, PRLR, ER α , and PgR levels, and estrogen target genes. Induced endogenous hPRL, but not exogenous hPRL added to the media, increased ER α levels as well as estrogen responsiveness in the Tet-On MCF7hPRL cells, suggesting that effects on breast cancer development and progression by estrogen may be amplified by cross-regulation of ER α levels by endogenous hPRL.

2. Materials and methods

2.1. Materials

The following antibodies were used for Western analyses: hPRL (anti-hPRL-IC-5) was obtained from the National Institute of Diabetes and Digestive and Kidney Disease (NIDDK) and Dr. Parlow; ER α (NCL-ER-6F11/2) from Novocastra Laboratories Ltd. (United Kingdom); IPRLR (PRLr-ECD 35-9200) and iPRLR (PRLr 34-4800) from Zymed (San Francisco, CA, USA); cyclin D1 (MS-210-P1) from Neomarkers (Fremont, CA, USA); progesterone receptor (NCL-PGR-312) from Novocastra (United Kingdom); and Bcl-2 (sc-492) from Santa Cruz Biotechnology Inc. (Santa Cruz, CA, USA). Recombinant hPRL (Lot AFP795) was obtained through the National Hormone and Pituitary Program, NIDDK, and Dr. Parlow. The ERE-luciferase construct contains three oxytocin receptor estrogen responsive

elements located upstream of a luciferase reporter [31]. All other reagents were obtained from Sigma–Aldrich Corp. (St. Louis, MO, USA).

2.2. Development of stable Tet-On MCF7hPRL cells

Tet-On MCF7 cells that had been selected for high responsive transcriptional activator (rtTA) activity were a gift from Dr. John Pink, Case Western University. Tet-On MCF7 cells were maintained in RPMI 1640 medium containing Tet System approved 10% fetal bovine serum (Clontech), penicillin and streptomycin, insulin, and geneticin (G418) 400 μ g/ml. hPRL cDNA [32] was blunt end cloned into the *Xba*I site of the Tet-On system pTRE vector (Clontech), and the construct confirmed by sequencing. Tet-On MCF7 cells were co-transfected with pTRE-hPRL and pTK-Hyg (Clontech), and stable transfectants were selected using hygromycin B (200 μ g/ml). The resulting clones were screened for inducible hPRL expression in response to doxycycline using RT-PCR and these were used for further experiments. Tet-On MCF7hPRL cells were maintained in the same media as for Tet-On MCF7 cells with the addition of hygromycin B. Four to five days prior to experiments with estrogen, cells were changed to steroid free, phenol-red free RPMI 1640 containing 5% Tet System approved FBS (Clontech) that had been stripped three times with dextran coated charcoal ($3 \times$ CSS) to remove endogenous steroids.

2.3. RT-PCR and Western analysis

For hPRL mRNA analysis, cells were plated in six well plates at 1×10^5 cells per well, serum starved for 24 h, and treated $-/+$ doxycycline as indicated. RNA was isolated using the RNeasy kit (Qiagen) and RT-PCR was performed as previously described [7]. For Western analysis, Tet-On MCF7hPRL cells were plated at 1×10^6 cells per 100 mm dish, serum starved for 24 h, and treated $-/+$ 0.5 μ g/ml doxycycline for different times, or cells were treated with 0.5 μ g/ml doxycycline for 48 h followed by treatment with 1 nM E2 or 4 nM hPRL as indicated. Cell lysates were harvested and analyzed as previously described [7]. Primary antibody concentrations were as follows: hPRL 1:10,000; ER α 1:1000; IPRLR 1:2000; iPRLR 1:6250 (0.04 μ g/ml); cyclin D1 1:500; PgR 1:1000; and Bcl-2 1:1000. Signals were quantitated using Molecular Dynamics, Inc. (Sunnyvale, CA) Densitometer and ImageQuant, version 4.2a software.

2.4. Secreted hPRL detection and quantitation

Conditioned media was collected from Tet-On MCF7 parental cells and Tet-On MCF7hPRL cells that were treated without or with doxycycline as indicated. Nb2 cells obtained from Dr. Lewis Sheffield, were employed to measure bioactive secreted hPRL in the Tet-On MCF7hPRL conditioned

media [33–35]. Nb2 cells were maintained in Fischer's Media, containing 10% horse serum, 10% FBS, and 0.1 mM β -mercaptoethanol. Nb2 cells were blocked in the cell cycle for 24 h prior to the assay in the media above without 10% FBS. Cells were then plated at 10^5 cells per well in a 96 well plate and incubated with the conditioned media or hPRL standards (0–3 ng/ml) for 72 h. After 72 h, total cell number was determined by the MTS assay (Promega, Madison, WI, USA) according to manufacturer's protocol. The amount of secreted hPRL was calculated from the hPRL standard curve. For detection of hPRL protein in Tet-On MCF7hPRL cell conditioned media by Western, 500 μ l of conditioned media was harvested. Protein from conditioned media was precipitated with 1.5 ml of ice-cold acetone, incubated at -20°C for 2 h, and centrifuged at $14,000 \times g$ for 10 min. Precipitated protein pellets were allowed to dry slowly, resuspended in 10 μ l of protein sample buffer, and analyzed by Western as described above.

2.5. Tet-On MCF7hPRL cell proliferation

Tet-On MCF7hPRL cells were grown in estrogen free media for 4 days then plated at 5×10^5 cells per 60 mm tissue culture dish. Cells were changed to serum free media for 24 h and treated for 48 h with or without vehicle, 0.5 μ g/ml doxycycline, and/or 1 nM E2. Cell number was analyzed as previously described [7].

2.6. Reporter gene assay

Tet-On MCF7hPRL cells were grown in estrogen free media for 4 days then cells were plated into 12 well tissue culture plates at 3×10^5 cells per well. Cells were serum starved for 24 h, then transiently transfected using SuperFect (Qiagen) as previously described [36]. After 3 h, the transfection complex was replaced with serum free media $-/+$ doxycycline and $-/+$ E2 at different concentrations for 6 h. Cells lysates were harvested and analyzed for luciferase and β -galactosidase activity as described [36].

3. Results

3.1. Development and characterization of a mammary epithelial tumor cell line that inducibly overexpresses hPRL

To study the local production of hPRL in human mammary epithelial tumor cells, we developed a MCF7hPRL inducible cell model using the Tet-On Gene Expression System (Clontech). As diagrammed in Fig. 1, when the Tet-On MCF7hPRL cells are treated with doxycycline, the rtTA recognizes the tet responsive element (TRE) on the responsive plasmid and activates hPRL transcription. To confirm that doxycycline treatment of Tet-On MCF7hPRL cells increased hPRL transcription and hPRL protein expression,

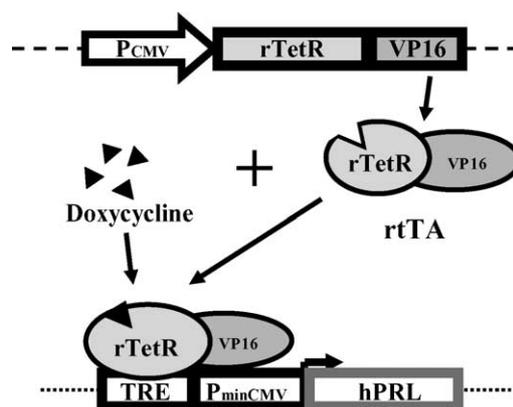


Fig. 1. Schematic representation of induction of hPRL production in the Tet-On MCF7hPRL cell model. MCF7 cells were stably transfected with the reverse Tet responsive transcriptional activator regulator plasmid. The rtTA consists of the reverse Tet repressor (rTetR) fused to the VP16 transactivation domain of herpes simplex virus, under the control of a cytomegaloviral promoter (PCMV). Next, the Tet-On MCF7 cells were stably transfected with the pTRE responsive plasmid expressing human prolactin, under the control of a tet responsive element followed by a minimal cytomegaloviral promoter (PminCMV). In the presence of doxycycline, the rtTA binds to the TRE in the responsive plasmid and activates transcription of hPRL.

cells were treated without and with doxycycline and analyzed by RT-PCR and Western blots. Tet-On MCF7hPRL cells treated with doxycycline showed a marked induction of hPRL mRNA (Fig. 2A) and protein expression (Fig. 2B) compared to the untreated control, indicating that the Tet-On MCF7hPRL cells are functioning properly.

In order to confirm that this induced hPRL could be secreted and was therefore capable of paracrine activity, conditioned Tet-On MCF7hPRL cell media was collected and analyzed for bioactive hPRL using the rat pre-T lymphoma cell line Nb2. Nb2 cells express elevated levels of a high affinity PRLR isoform, and depend on PRL for growth, making them extremely sensitive to PRL and ideal for detection of secreted active PRL [33–35]. To optimize doxycycline treatment conditions, the Tet-On MCF7hPRL cells were subjected to increasing concentrations of doxycycline for variable times. Following treatment, conditioned media was harvested and secreted hPRL was detected by the Nb2 cell bioassay. As shown in Fig. 2C, hPRL secretion was detected in response to low concentrations of doxycycline, reaching a maximum at 0.5 μ g/ml. We chose this concentration for further experiments. Although we detected hPRL mRNA, and protein in cell lysates as early as 24 h, hPRL activity was not appreciably increased in the media using this assay until 48 h after doxycycline treatment.

Using this bioassay, we determined the concentration of hPRL secreted by the doxycycline-induced cells by comparing the conditioned media to hPRL standards. Conditioned media from Tet-On MCF7hPRL cells treated with doxycycline for 72 h contained about 10 ng/ml of hPRL (100 ng PRL/ 1×10^6 cells/72 h) and untreated cells contained about 0.5 ng/ml of hPRL (5 ng hPRL/ 1×10^6 cells/72 h) (Fig. 3A).

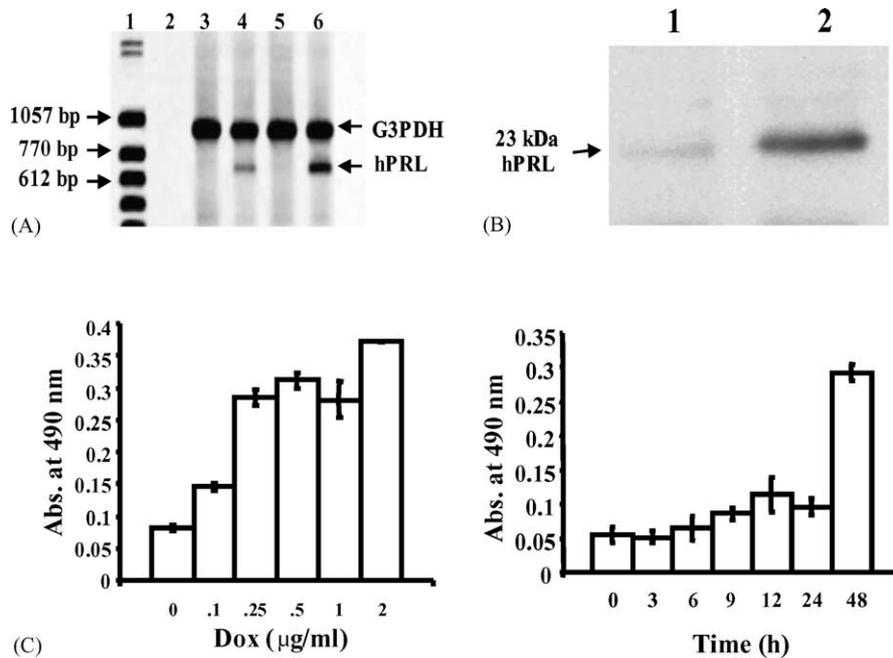


Fig. 2. Doxycycline induces hPRL mRNA, protein expression, and secretion in Tet-On MCF7hPRL cells. (A) RT-PCR analysis of mRNA isolated from cells cultured for 24 and 48 h without or with 0.5 µg/ml doxycycline. Lane 1: lambda DNA ladder; lane 2: negative control without the addition of cDNA to the PCR reaction; lane 3: 24 h without doxycycline; lane 4: 24 h with 0.5 µg/ml doxycycline; lane 5: 48 h without doxycycline; lane 6: 48 h with 0.5 µg/ml doxycycline. G3PDH was used as a positive internal control in all samples. (B) Representative Western analysis of hPRL protein induction by doxycycline. Cells were treated without (lane 1) or with (lane 2) 0.5 µg/ml doxycycline for 24 h. Cell lysates were harvested and 30 µg of protein was analyzed. (C) Dose-response and time-dependence of doxycycline-induced hPRL secretion by Tet-On MCF7hPRL cells. Cells were washed and serum starved for 24 h, then treated with increasing concentrations of doxycycline for 48 h in serum free media (left) or treated with 0.5 µg/ml doxycycline in serum free media for different times (right). Ten microliters of harvested conditioned media was diluted in 40 µl of serum free media; bioactive secreted hPRL was detected using Nb2 cell proliferation (see Section 2). Each value is the mean ± S.D. of three replicates.

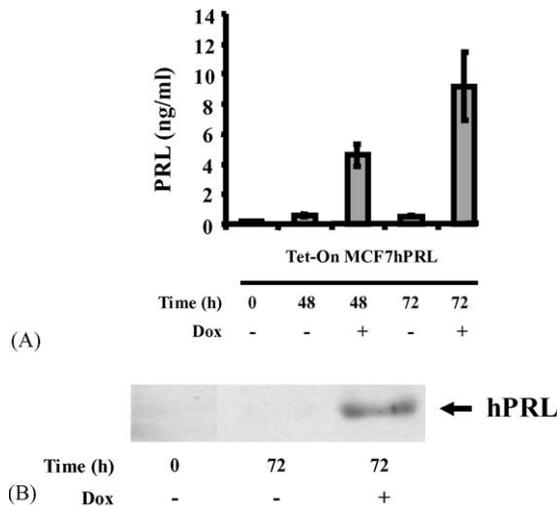


Fig. 3. Quantitation of dox-inducible hPRL secretion by Tet-OnMCF7 hPRL cells. (A) Tet-On MCF7hPRL cells were serum starved for 24 h then treated without or with 0.5 µg/ml doxycycline for 48 or 72 h. The amount of bioactive secreted hPRL was quantitated using Nb2 cell proliferation compared to hPRL standards. Each value is the mean ± S.D. of triplicates. (B) Representative Western analysis of hPRL in the conditioned media of Tet-On MCF7hPRL cells. Cells were serum starved for 24 h then treated without or with 0.5 µg/ml doxycycline in serum free media. Five hundred microliters of conditioned media was harvested at 0 and 72 h and precipitated with acetone prior to analysis.

The latter is similar to the amount of hPRL secreted by cultured breast cancer cells in previous reports [13,14]. In this system, doxycycline treatment induced a 20-fold increase in hPRL secretion compared to basal levels. hPRL secretion by the control Tet-On MCF7 parental cell line, without the pTRE responsive plasmid, was similar to uninduced Tet-On MCF7hPRL cells (data not shown). To confirm the induction of secreted hPRL, protein in conditioned media from doxycycline treated Tet-On MCF7hPRL cells was precipitated with acetone and analyzed by Western blot (Fig. 3B). These studies indicate that the Tet-On MCF7hPRL cells express high levels of hPRL mRNA, hPRL protein, and secrete hPRL into the media when induced with doxycycline.

3.2. Tet-On MCF7hPRL cells proliferate faster in response to doxycycline-induced hPRL and E2

Next we examined the effect of doxycycline-induced hPRL on cell growth in the presence and absence of 17β-estradiol (E2). Tet-On MCF7hPRL cells were grown in 3 × CSS serum for 5 days prior to the experiment to minimize exposure to estrogens. Cells were plated in 3 × CSS and then serum starved for 24 h ($t = 0$), prior to treatment with doxycycline, E2 or both together. Viable cells were then counted 48 h after treatment (Fig. 4). The rate of apoptosis was extremely low (data not shown); therefore,

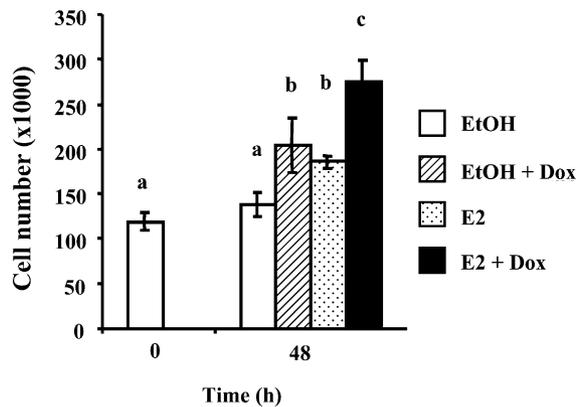


Fig. 4. Growth of Tet-On MCF7hPRL cells in response to doxycycline and E2 treatment. Representative experiment. Cells were serum starved for 24 h then treated without or with vehicle, 0.5 $\mu\text{g}/\text{ml}$ doxycycline, and 1 nM E2. Viable cells were counted at 0 and 48 h. Results are expressed as the mean \pm S.D. of triplicate plates. Different letters denote significantly different values analyzed by one-way ANOVA, followed by Student–Newman–Keuls multiple comparison test ($P < 0.05$).

these changes represent primarily proliferation. Numbers of doxycycline treated cells increased 1.5-fold over untreated controls ($P < 0.05$), similar to E2 treatment, indicating that increased endogenous hPRL production can stimulate Tet-On MCF7hPRL cell proliferation. Cells treated with both doxycycline and E2 together showed a further increase ($P < 0.05$) over doxycycline or E2 treatment alone. These results are consistent with previous data showing breast cancer cells proliferate in response to these hormones [7,9] and confirm that endogenous PRL has mitogenic effects in this system.

3.3. Doxycycline-induced hPRL increases ER α and IPRLR, but not iPRLR expression in Tet-On MCF7hPRL cells

MCF7 cells express multiple PRLR isoforms generated by alternative splicing, including the long PRLR (IPRLR) and the intermediate PRLR (iPRLR), each with different signaling capabilities. The predominant isoform in these cells, the IPRLR, is the most studied PRLR isoform and is a classic type I cytokine receptor. It mediates signals to many pathways, including the well-studied Jak/Stat pathway (reviewed in [10]). The iPRLR isoform has a truncated cytoplasmic domain and a unique 13 amino-acid sequence after the extended box 2 region. It also activates the Jak/Stat pathway, but it is less effective in mediating signals to Fyn [37]. MCF7 cells also express ER α , although they have undetectable levels of ER β (Wade Welshons, personal communication).

To investigate the effect of increased endogenous hPRL on expression of these receptors, we treated Tet-On MCF7hPRL cells with or without doxycycline for 48 or 72 h and analyzed the cell lysates for ER α , IPRLR, and iPRLR by Western blot (Fig. 5A). After 48 h of doxycycline treatment, cells showed a three-fold increase in ER α and a two-fold increase in IPRLR expression, indicating that induced hPRL

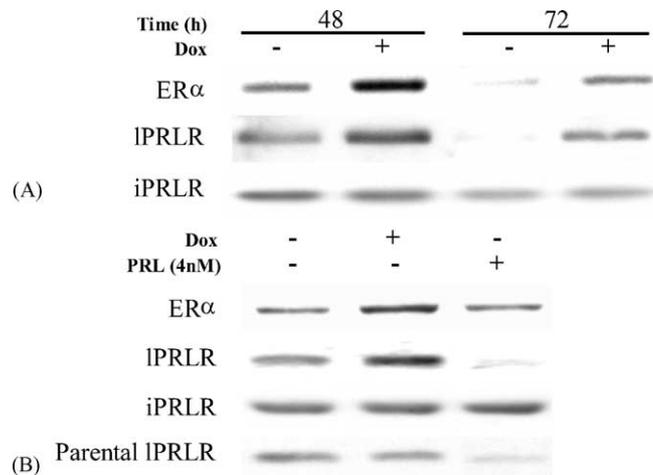


Fig. 5. Dox-induced endogenous hPRL increases ER α and IPRLR, but not iPRLR in Tet-On MCF7hPRL cells. (A) Representative Western analysis of ER α , IPRLR and iPRLR in cell lysates that were serum starved for 24 h then treated without or with 0.5 $\mu\text{g}/\text{ml}$ doxycycline for 48 or 72 h. (B) Representative Western analysis of ER α , IPRLR, and iPRLR in Tet-On MCF7hPRL cell lysates, and IPRLR in Tet-On parental cell lysates. Cells were serum starved for 24 h then treated without or with 0.5 $\mu\text{g}/\text{ml}$ doxycycline, and vehicle or 4 nM hPRL for 48 h.

upregulated these receptors. Doxycycline treatment had no effect on iPRLR levels, indicating that this effect was isoform specific. A similar trend was seen at 72 h, although levels of receptors were uniformly lower. Levels of ER α , IPRLR, and iPRLR were not altered in the parental Tet-On MCF7 cells treated with doxycycline (data not shown).

Because PRL has been reported to up- or down-regulate PRLR at different targets depending on dose and experimental system (reviewed in [38–40]), we compared the effects of exogenous hPRL and doxycycline-induced endogenous hPRL. As shown in Fig. 5B, IPRLR levels decreased and ER α expression was unchanged when cells were treated with 4 nM exogenous hPRL for 48 h, in contrast to the upregulation of both receptors following increased endogenous synthesis. The iPRLR isoform was not changed by either treatment. Treatment with lower concentrations (0.04 and 0.4 nM) of exogenous hPRL also had no effect on ER α , IPRLR, or iPRLR levels (data not shown). The parental Tet-On MCF7 cells responded similarly to exogenous hormone. IPRLR was downregulated (Fig. 5B), and ER α levels were not altered by any of the treatments (data not shown).

3.4. Doxycycline-induced hPRL increases Tet-On MCF7hPRL cell responsiveness to E2

The number of estrogen receptors expressed within a cell has been shown to be a limiting factor for E2 responsiveness [41]. Because we observed an increase in ER α in response to increased endogenous hPRL, we examined the ability of the Tet-On MCF7hPRL cells to mediate the classical E2 response, activation of transcription via an

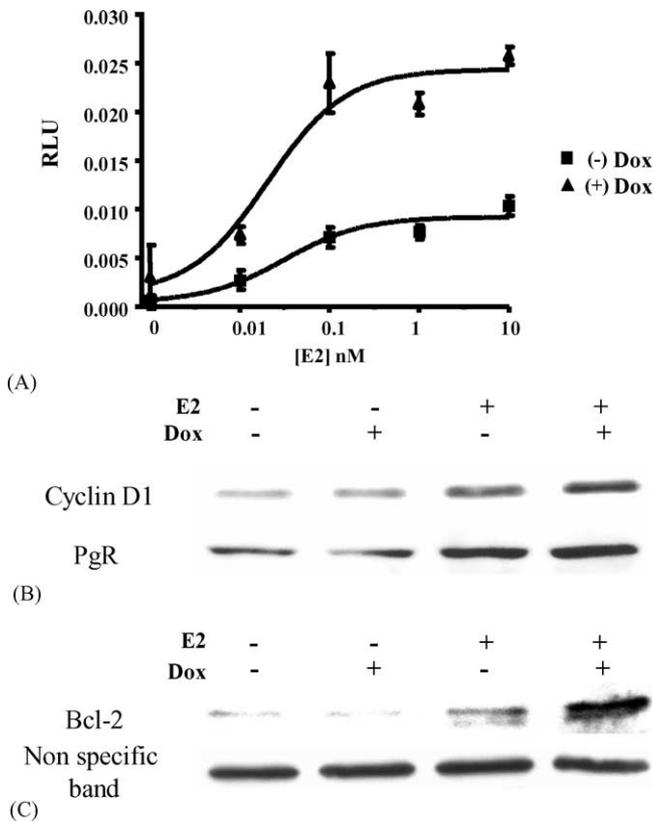


Fig. 6. Dox-induced hPRL production increases E2 sensitivity in Tet-On MCF7hPRL cells. (A) Representative experiment. Tet-On MCF7hPRL cells were plated in 5% $3 \times$ CSS for 24 h then serum starved for 24 h in the presence of 0.5 μ g/ml doxycycline. Cells were transiently transfected with an ERE-luciferase reporter and β -galactosidase control construct, then treated with different concentrations of 17 β -estradiol (E2) for 6 h. Cells lysates were assayed for luciferase activity, and β -galactosidase activity was used to correct for transfection efficiency. The graph shows relative luciferase units (RLU) as a function of E2 concentration on a logarithmic scale. Data were analyzed using a nonlinear regression equation (Prism 3.0). Points represent the mean \pm S.D. of triplicates. (B) and (C) Representative Western analysis of cyclin D1 and PgR (B) or Bcl-2 (C) in Tet-On MCF7hPRL whole cell lysates. Cells were serum starved for 24 h, treated without or with 0.5 μ g/ml doxycycline for 48 h and then treated with vehicle control or 1 nM E2 for 6 h (B) or 48 h (C).

ERE enhancer (Fig. 6A). Cells that were pre-treated with doxycycline for 48 h showed a three-fold increase in maximal ERE-luciferase activity compared to the vehicle treated cells, although the ED₅₀ was not significantly different. We also examined effects on expression of other known E2 targets including the cell cycle regulator, cyclin D1; the classic E2 responsive protein, progesterone receptor; and the anti-apoptotic protein, Bcl-2 [3,42–46]. Cells were estrogen withdrawn, pre-treated with doxycycline for 48 h, and then treated with or without 1 nM E2 (Fig. 6B and C). Doxycycline treatment alone modestly increased cyclin D1 levels, consistent with our previous data showing that PRL increases cyclin D1 transcription [7,36]. E2 treatment alone increased cyclin D1 levels 3.5-fold, also consistent with previous reports [42,43,47]. Interestingly, treatment with

E2 and doxycycline together increased cyclin D1 levels 4.5-fold over untreated control, showing an increased efficacy of E2 treatment following doxycycline. Doxycycline treatment did not alter PgR, but slightly enhanced the effect of E2. When Tet-On MCF7hPRL cells were treated with both E2 and doxycycline together, they showed almost a 10-fold increase in Bcl-2 levels compared to the lack of change when treated with doxycycline alone and a three-fold increase when treated with E2 alone. Together, these data indicate that doxycycline-induced hPRL not only increases ER α levels, but also in doing so, increases the responsiveness of the Tet-On MCF7hPRL cells to E2.

4. Discussion

The complex interactions between hPRL and other hormones within the mammary gland important for normal development (reviewed in [1–4]) suggest a role for these hormones in the development and progression of breast cancer. The discovery that mammary epithelial cells produce and secrete hPRL [13–15] has redirected breast cancer research in this area. However, studying autocrine/paracrine actions of PRL in the mammary gland has been challenging. Here we developed a hormonally responsive, mammary tumor cell line that inducibly expresses and secretes hPRL as a tool to study endogenous hPRL and how it may regulate receptor levels as well as responsiveness to other hormones in breast cancer.

One mechanism of hormone interaction is cross-regulation of their receptors. Endogenous hPRL increased ER α protein levels substantially, which was associated with a robust increase in E2 responsiveness, measured by ERE-luciferase activity, and activation of E2 target genes, including cyclin D1 and Bcl-2. These targets are important for cell cycle progression and inhibition of cell death, respectively, and both are implicated in breast cancer [48–50]. The inability of induced PRL to similarly augment the PgR response was unexpected, and may be explained by maximal stimulation of this pathway by E2 alone. PRL induces transcription of ER α through the Jak2/Stat5 pathway in the rodent corpus luteum (reviewed in [16]). In light of the well characterized ability of this pathway to mediate many PRL actions (reviewed in [10]), and its recognized importance in mammary development [51,52], the Jak2/Stat5 pathway is likely to be involved in PRL regulation of ER α at this target as well.

The status of ER α expression in breast cancer is critical for prognosis and therapeutic strategies. However, few models for studying regulation of ER α expression are available. Expression of a PRL transgene within murine mammary epithelium resulted in ER α +, as well as ER α - mammary tumors [53]. While these findings in vivo do not demonstrate direct induction of mammary ER α expression by PRL, the lack of this tumor population in response to other mammary oncogenes suggests that this relationship also may be important in tumorigenesis. These models of locally produced

PRL provide tools to dissect this potentially very significant relationship in breast cancer development and progression.

Endogenous hPRL also selectively increased the major PRLR isoform, IPRLR, without affecting iPRLR protein expression. Multiple factors have been shown to govern total PRLR expression at different target tissues [38–40]. Receptor levels may increase due to changes in transcription, translation, or protein stability, or decrease due to receptor degradation and/or processing. While long-term PRL treatment in vivo upregulated mammary PRLR [54,55], and higher circulating PRL during various physiologic states can be associated with elevated mammary PRLR [56–58], these models are complicated by complex indirect effects and feedback loops. In vitro, treatment with exogenous ligand generally results in PRLR downregulation, following internalization and subsequent degradation [59–61], similar to the effect of exogenous PRL in the present studies. A very low concentration (0.04 nM) of PRL slightly increased (50%) PRL binding in another breast cancer cell line, but the effect was lost at 0.4 nM PRL [62]. Interestingly, engineering the ER α –breast cancer cell line, MDA-MB-435, to constitutively overexpress PRL, also resulted in elevated IPRLR [63].

It is likely that transcription plays a major role in the increase in IPRLR seen here. Exogenous PRL increased activity of one of the PRLR promoters via Stat5 in pancreatic β -cells in vitro [64]. Analyses of the PRLR promoter in both rodents and humans have revealed additional elements that have been shown to mediate a response to PRL in other systems, including AP-1 and Sp1 sites, as well as sites for interaction with C/EBP β [65–69]. However, additional studies are required to determine the effective mechanism for this response in both normal and neoplastic epithelial cells.

The substantial increase in ER α in response to increased endogenous hPRL compared to the modest, or lack of effect, using exogenous hPRL in the parent cell line [24,30], and the marked disparity between the effects of exogenous and increased endogenous hPRL on both ER α and IPRLR levels in the present study, are consistent with important differences in the mechanism of action of endogenously produced and exogenously added hPRL. When compared to normal physiological and commonly used experimental concentrations, the amount of hPRL secreted into the media by the Tet-On MCF7hPRL cells is very low, accumulating to concentrations near the K_d of the PRLR only after 72 h [70]. This phenomenon is similar to that described for autocrine production of hGH by MCF7 cells [71]. One possible explanation for this difference is that the locally synthesized PRL may be post-translationally modified, resulting in a different spectrum of activity compared to the unmodified hPRL from NIDDK. For example, phosphorylated hPRL has been shown to inhibit growth promoting effects of unmodified PRL, while potently stimulating effects associated with differentiation (reviewed in [72]); the latter might include increased ER α expression. The approximate agreement between mitogenic PRL activity measured by the Nb2 assay,

and immunoreactive PRL in the conditioned media in our studies, does not support this as a major factor. However, our data do not rule out some modification of a minor component that could account for our findings. Alternatively, endogenous hPRL may have access to different pathways than PRL secreted by the pituitary and presented to the plasma membrane. Such “intracrine” signaling by PRL has been suggested in other cell types [73–75], as well as by other growth factors including epidermal growth factor, platelet derived growth factor, and renin-angiotensin [76–78]. These possibilities are currently under investigation.

We have developed a model to study the effects of endogenous hPRL in mammary tumor cells to better understand the autocrine/paracrine signaling of PRL in these cells. Endogenous hPRL, but not exogenous hPRL, upregulated both IPRLR and ER α protein levels, and substantially increased the responsiveness of these cells to E2. This model will allow investigation of endogenous hPRL in mammary epithelial cells and will enable further dissection of PRL effects on other hormone signaling pathways to determine the role of hPRL in breast cancer. These studies may help to develop new prognostic indicators and treatments for the disease.

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