

CHAPTER 7

Progesterone Receptor Action: Translating Studies in Breast Cancer Models to Clinical Insights

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Abstract

Progesterone receptors (PR) are useful prognostic indicators of breast cancers likely to respond to anti-estrogen receptor (ER) therapies. However, the role of progesterone, therapeutic progestins, or unliganded or liganded PR in breast cancer development or progression remains controversial. PR are ligand-activated transcription factors that act in concert with intracellular signaling pathways as “sensors” of multiple growth factor inputs to hormonally regulated tissues, such as the breast. The recently defined induction of rapid signaling events upon progestin-binding to PR-B provides a means to ensure that receptors and coregulators are appropriately phosphorylated as part of optimal transcription complexes. PR-activated kinase cascades may provide additional avenues for progestin-regulated gene expression independent of PR nuclear action. Herein, we present an overview of progesterone/PR and signaling cross-talk in breast cancer models and discuss the potential significance of progestin/PR action in breast cancer biology using examples from both in vitro and in vivo models, as well as limited clinical data. Kinases are emerging as key mediators of PR action. Cross-talk between PR and membrane-initiated signaling events suggests a mechanism for coordinated regulation of gene subsets by mitogenic stimuli in hormonally responsive normal tissues. Dysregulation of this cross-talk mechanism may contribute to breast cancer biology; further studies are needed to address the potential for targeting PR in addition to ER and selected protein kinases as part of more effective breast cancer therapies.

Introduction

Normal breast development requires estrogen receptor (ER α), progesterone receptor (PR) and peptide growth factors. Estrogen stimulates ductal elongation and progestins induce ductal sidebranching and alveologenesis.¹ Epidermal growth factor (EGF), in addition to promoting the proliferation of terminal end-buds, augments estrogen-induced ductal outgrowth and progesterone-induced sidebranching.² Indeed, estrogen induces PR isoform expression only in the presence of EGF,³ suggesting the existence of important cross-talk between EGFRs and both steroid receptors (SRs). Ligand-activated PRs and ERs are potent mitogens in the developing breast and mammary epithelial cells express PR as well as ER α . Moreover, estrogen is usually required to induce the expression of PR. PR and ER are normally expressed by only ~7-10% of nondividing epithelial cells in the lumen of the mature mammary gland. This nonproliferative condition

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appears to be sustained by such inhibitory molecules as TGF-beta or high levels of p27, a CDK inhibitor (reviewed in G.W. Robinson et al ⁴). In response to communication between stromal and epithelial compartments, SR-positive epithelial cells express and secrete pro-proliferative molecules, such as Wnts or IGF-II, thereby inducing the proliferation of adjacent SR-negative epithelial cells.^{4,5} Recent data indicate that SR-positive cells in the breast may support the activity of nearby stem-like progenitor cells via the expression of secreted factors.⁶

In contrast to the normal breast, where proliferating cells are devoid of SRs, the majority of newly diagnosed breast cancers (~70-80%) express ER and PR. The existence of SR-positive proliferating cells in breast cancer indicates that SR-positive cells may undergo an early switch to autocrine stimulation and/or continue to divide. Breast cancer is not the only setting where PR-containing cells divide. In an *in vivo* model of the mammary gland during pregnancy, the PR-B isoform colocalizes with cyclin D1 in BrdU-stained (dividing) cells.⁷ Thus, signaling pathways involved in normal mammary gland growth and development are likely reactivated during breast cancer progression.

Progestins are recognized as mediators of increased post-menopausal breast cancer risk when taken as part of combined hormone replacement therapy relative to estrogen alone or placebo.⁸ Experimental animal models of the effects of hormones on the postmenopausal mammary gland indicate that progestins stimulate proliferation.^{9,10} While progestins are not carcinogens, progesterone might induce recently initiated precancerous breast cell populations to inappropriately reenter the cell cycle or stimulate dormant stem cells to undergo self-renewal (discussed below). Breast tumors develop resistance to endocrine-based treatments (anti-estrogens and/or aromatase inhibitors) as they progress. However, the majority (65%) of resistant breast cancers retain high levels of SRs (ER α and PRs). In these resistant, SR-positive cancers, the rapid action of SRs at the membrane might begin to inappropriately trigger the classical transcriptional activities of SRs. In this way, PRs activated by extremely low or sub-threshold concentrations of hormone or PRs phosphorylated in the absence of hormone can activate membrane-associated signaling pathways, including *c*-Src kinase, EGFR and the p42/p44 MAPK pathway. Elevation of MAPK activity and downstream signaling frequently occurs in breast cancer, providing a strong survival and proliferative stimulus to breast cancer cells. MAPK signaling downstream of EGFR or Her2 (erbB2) is also associated with resistance to endocrine therapies.¹¹

This chapter focuses specifically on the role of progesterone and progesterone receptors (PR) in the pathophysiology of breast cancer. We review the literature describing PR-initiated genomic and nongenomic signaling pathways in breast cancer progression with the purpose of highlighting key kinases involved in the integration of rapid cytoplasmic signaling events and PR nuclear actions. We also discuss the clinical findings relevant to the use of PR status in the prediction of breast cancer behavior, evidence for PR action in breast cancer and the potential for PR ligands as therapeutic agents.

Classical Actions of PRs

PRs are activated through binding with the ovarian steroid ligand, progesterone. PRs are classically defined as ligand-activated transcription factors that regulate gene expression by binding directly or indirectly to DNA. Three PR isoforms are the product of a single gene located on chromosome 11 at q22-23 that undergoes transcription via the use of alternate promoters and internal translational start sites.¹² PR isoforms consist of the full length PR-B (116 kDa), N-terminally-truncated PR-A (94 kDa) and PR-C-isoforms (60 kDa). PR-positive cells usually co-express PR-A and PR-B isoforms; these receptors have different transcriptional activities within the same promoter context, but can also recognize entirely different promoters.^{13,14} PR-B is required for normal mammary gland development,¹⁵ while PR-A is essential for uterine development and reproductive function.¹⁶ PR-C is devoid of classical transcriptional activity and instead functions as a dominant inhibitor of uterine PR-B in the fundal myometrium during labor.¹⁷ In the absence of progesterone, PRs are complexed with several chaperone molecules including heat shock protein (hsp) 90, hsp70, hsp40, Hop and p23; these interactions are requisite for proper protein folding

and assembly of stable PR-hsp90 heterocomplexes that are competent to bind ligand.¹⁸ Hsps also function to connect PRs to protein trafficking systems. After binding to progesterone, the receptors undergo restructuring, dimerization and hsp dissociation. Activated receptors bind directly to specific progesterone response elements (PREs) and PRE-like sequences in the promoter regions of such target genes as *c-myc*,¹⁹ *fatty acid synthetase*,²⁰ and *MMTV*.²¹ Treatment with progestin also results in an upregulation of regulatory molecules without classical PREs in their proximal promoter regions, such as *EGFR*^{22,23}, *c-fos*^{24,25} and *cyclin D1*.^{26,27} PR regulation of genes without canonical PREs can occur through indirect DNA-binding mechanisms, as in the example of PR binding to Specificity protein 1 to promote p21 transcription in the presence of progestin.²⁸ PRs can also regulate genes by tethering to activating protein 1²⁹ or signal transducers and activators of transcription (STATs).^{25,30}

When either directly or indirectly bound to DNA, PRs regulate the basal transcription machinery in conjunction with nuclear receptor coregulatory molecules. Coregulators modulate transcription through chromatin remodeling and recruitment of transcriptional machinery (e.g., RNA Polymerase-II). Histone acetyl transferases (HATs) and histone deacetylases (HDACs) function as coactivators and corepressors, respectively. Both HATs and HDACs coordinate transcriptional activity with other regulator proteins, including the ATP-dependent chromatin remodeling complexes (SWI/SNF), arginine methyltransferases (CARM1 and PRMT1) and histone kinases (reviewed in N.J. McKenna, B.W. O'Malley³¹).

Direct PR Phosphorylation in Breast Cancer Models

Similar to other SR family members, phosphorylation-dephosphorylation events add multi-functionality to PR action (Fig. 1). Several protein kinases phosphorylate PR isoforms primarily on serine residues within the amino-termini and, to a lesser degree, on serine residues throughout the receptor.^{12,32} PR contains a total of 14 known phosphorylation sites (reviewed in C.A. Lange³³). Serines at positions 81, 162, 190 and 400 appear to be constitutively phosphorylated in the absence of hormone³⁴ (Fig. 1). One to two hours after progestin treatment, serines at positions 102, 294 and 345 are maximally phosphorylated.³⁵ Specific kinases have been identified that are responsible for phosphorylation of selected sites. Serines at positions 81 and 294 are phosphorylated by casein kinase II³⁶ and mitogen-activated protein kinase (MAPK),^{37,38} respectively. Progestins can also stimulate Ser294 phosphorylation independently of MAPKs by activation of an unknown kinase(s).³⁹ Eight of the total 14 sites (i.e., serines 25, 162, 190, 213, 400, 554, 676 and Thr430) are phosphorylated by cyclin A/cyclin-dependent protein kinase 2 (CDK2) complexes in vitro.^{34,40} Only five of these sites (i.e., serines 162, 190, 213, 400, 676) are proven in vivo phosphorylation sites.^{34,36,40}

While the function of PR phosphorylation is incompletely understood, it might influence aspects of transcriptional regulation, such as interaction with coregulators, as reported for ER- α ⁴¹ and recently for PR.⁴² PR phosphorylation is also involved in the regulation of ligand-dependent³⁸ and -independent^{43,44} PR nuclear localization, receptor turnover, hormone sensitivity and transcriptional activities.^{37,38,45,46} As has been reported for ER α ,^{47,48} phosphorylated PRs are hypersensitive relative to their underphosphorylated counterparts.⁴⁹ For example, following a brief (5-15 min) pretreatment with EGF, phosphorylated nuclear PR-B receptors are transactivated by sub-physiologic progestin levels. EGF and progestins synergistically upregulate mRNA or protein levels for a number of growth regulatory genes,²⁵ including cyclin D1 and cyclin E;²² the regulation of cyclins by progestins is MAPK-dependent. Cyclins, in turn, regulate progression of cells through the cell cycle by interaction with cyclin-dependent protein kinases. Progestins activate CDK2,²⁷ which predominantly phosphorylates PRs at proline-directed (S/TP) sites,^{34,40} perhaps allowing for the coordinate regulation of PR transcriptional activity during cell cycle progression. In support of this idea, Narayanan and coworkers^{42,50} report that PR activity is highest in the S phase and lower in the G0/G1 phases of the cell cycle, but this activity is impaired during the G2/M phases, concomitant with lowered PR phosphorylation. Overexpression of Cyclin A or CDK2 enhanced PR transcriptional activity. While cyclin A interacts with the N-terminus of PR, CDK2 seems to

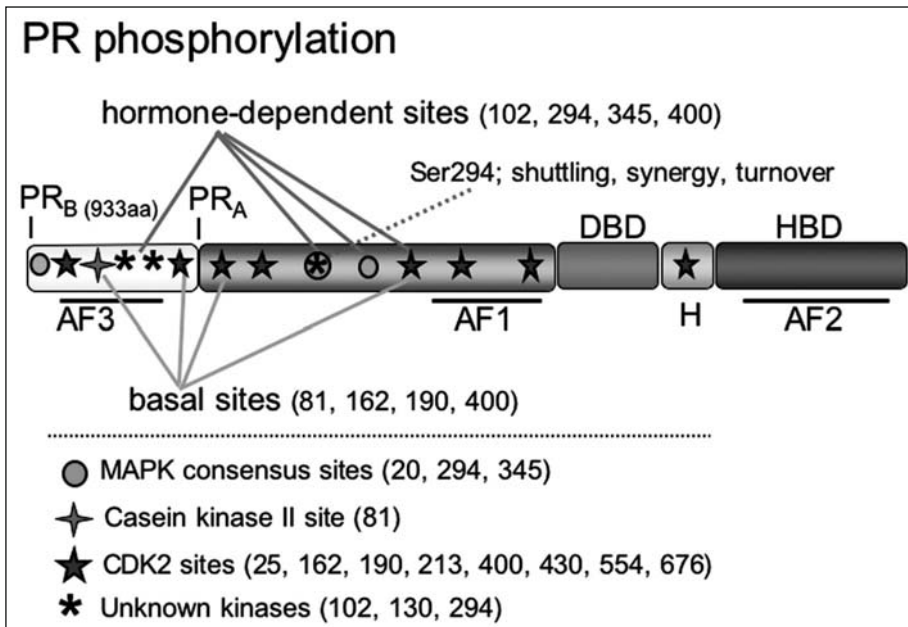


Figure 1. Phosphorylation sites in human PR. PR phosphorylation. Thirteen serine residues and one threonine residue in human PR are shown, to represent basal (constitutive) and hormone-induced phosphorylation sites⁴⁰ and may contribute to PR regulation by MAPK,³⁷⁻³⁹ casein kinase II,³⁶ and CDK2.^{34,40} Individual PR phosphorylation sites may be regulated by multiple protein kinases³⁹ and/or in a sequential manner,¹⁴³ illustrating the complexity of PR regulation by phosphorylation.

alter PR function indirectly by increasing the phosphorylation and recruitment of steroid receptor coactivator-1 (SRC-1) to liganded PR.

PR Ser294 Phosphorylation in Breast Cancer Models

PR Ser294 is rapidly phosphorylated upon exposure to ligand.³⁵ Ser294 is also a proline-directed or MAPK consensus site (PXXSP). Progestin-induced Ser294 phosphorylation occurs within 30-60 min independently of MAPK activation, whereas growth factor-induced Ser294 phosphorylation occurs within 3-5 mins in a MAPK-dependent manner.³⁹ PR Ser294 is considered a significant site for PR regulation by multiple kinases.^{37-39,49} Ser294 phosphorylation appears to mediate increased PR nucleo-cytoplasmic shuttling.³⁹ Rapid nuclear translocation of unliganded PR and nuclear export of liganded PR requires MAPK-dependent phosphorylation of PR Ser294.³⁹ PR nuclear sequestration in response to MAPK activation might serve to protect inactive or active receptors from degradation in the cytoplasm or upon nuclear export.³⁹ Following ligand binding, PR undergoes rapid downregulation.⁵¹ Phosphorylation of Ser294 greatly augments PR downregulation by making liganded PR a cytoplasmic target for ubiquitination and degradation by the 26S-proteasome pathway.^{37,39} In several recent reports, it has been shown that reversible phosphorylation of PR Ser294 couples increased transcriptional activity to rapid down-regulation of the PR protein by the ubiquitin-proteasome pathway.^{37-39,49,52} Further investigation is required to determine whether the link between these events involves regulation of transcriptional events by components of the ubiquitin pathway and/or participation of nucleo-cytoplasmic shuttling factors or chaperones.

In the absence of progestins, however, EGF-induced nuclear accumulation of PR is required for transcriptional activation. Labriola et al⁴³ report that exposure of T47D breast cancer cells to the EGF family member, heregulin, can stimulate PR nuclear localization, DNA binding and transcriptional activity in the absence of hormone. Heregulin exposure also resulted in activation of MAPK and PR Ser294 phosphorylation. Qiu et al³⁹ report that PR Ser294 phosphorylation results in similar nuclear activity. However, growth factors alone failed to stimulate PR transcriptional activity or alter PR downregulation in T47D cell variants.³⁸ However, in the presence of ligand, MAPK activation greatly augmented both of these events.^{38,39} One explanation for these apparently conflicting results is that differential expression of EGFR family members expressed on the cell surface between T47D cell line clones might lead to differences in the activation of downstream intracellular kinases, such as CDK2.⁴⁴ Indeed, regulation of PR by alternate signaling pathways may contribute to dysregulated gene expression and changes in cell growth and/or survival. For example, PR-B regulation of IRS-2 expression in breast cancer cells requires phosphorylation of PR Ser294 and occurs in the absence of ligand.⁴⁹ In any case, these exciting data^{39,43} suggest a continuum between PR hypersensitivity to extremely low ligand concentrations and complete ligand-independence, a phenomenon that is well-documented for androgen receptor (AR) and ER α .

Extranuclear Actions of PR

While the genomic effects of steroid hormone treatment are delayed by several minutes to hours (i.e., following transcription and translation), the extranuclear or nongenomic effects occur rapidly in only a few minutes. Progestin treatment of breast cancer cells causes a rapid and transient activation of MAPK signaling that is ER-dependent, but independent of PR transcriptional activity.^{53,54} Migliaccio et al were the first to report that estradiol activates p60-Src kinase and MAPK in MCF-7 cells⁵⁵ and that PR and ER α interact to stimulate p60-Src kinase in T47D cells.⁵³ Maximal activation of p60-Src kinase is observed within 2-5 minutes and downstream activation of p42/p44 MAPKs occurs within 5-10 minutes of progestin treatment.^{53,54}

Human PR contains a proline-rich (PXXP) motif that mediates direct binding to the Src-homology three (SH3) domains of signaling molecules in the p60-Src kinase family in a ligand-dependent manner.⁵⁴ In vitro experiments demonstrate that purified liganded PR-A and PR-B activate the c-Src-related protein kinase, HcK; PR-B but not PR-A activates c-Src and MAPKs in vivo. PR-B with a mutated PXXP sequence prevents c-Src/PR interaction and blocks progestin-induced activation of c-Src (or HcK) and p42/p44 MAPKs. Furthermore, mutation of the PR-B DNA-binding domain (DBD) abolished PR transcriptional activity without affecting progestin-induced c-Src or MAPK kinase activation. Therefore, nongenomic MAPK activation by progestin/PR-B/c-Src complexes probably occurs by way of a c-Src-dependent mechanism involving Ras activation via phosphorylation of the c-Src substrate adaptor proteins p190 and/or Shc and followed by Grb-2 and Sos binding (Fig. 2).

Ballare et al⁵⁶ report that MAPK activation by progestins is blocked by antiprogestins and anti-estrogens in COS-7 cells transfected with PR and ER α . They propose that c-Src/MAPK activation by PR is mediated indirectly by the interaction of the Src-homology two (SH2) domain of c-Src with phosphotyrosine 537 of ER α .⁵⁶ In their model, activation of c-Src and the MAPK pathway by progestins depends upon the presence of unliganded ER α , which interacts constitutively with PR-B via two domains that flank the proline-rich sequence of PR. Deletion of either of these two ER-interacting domains in PR-B blocked c-Src/MAPK activation by progestins in the presence of ER α .⁵⁶ Mutation of PR-B's PXXP domain had no effect. In contrast, Boonyaratanakornkit et al⁵⁴ report that ectopic PR expression increased basal c-Src activity in COS-7 cells in the absence of progestins and independently of added ER; co-expression of both PR-B and ER α reduced basal levels of c-Src activity. Under these conditions (i.e., low basal c-Src activity), progestin binding to PR-B clearly activated c-Src. In addition, progestins activated c-Src in PR-null MCF12A cells transduced with wild-type PR but not the PXXP-mutant PR adenoviruses. Both groups found

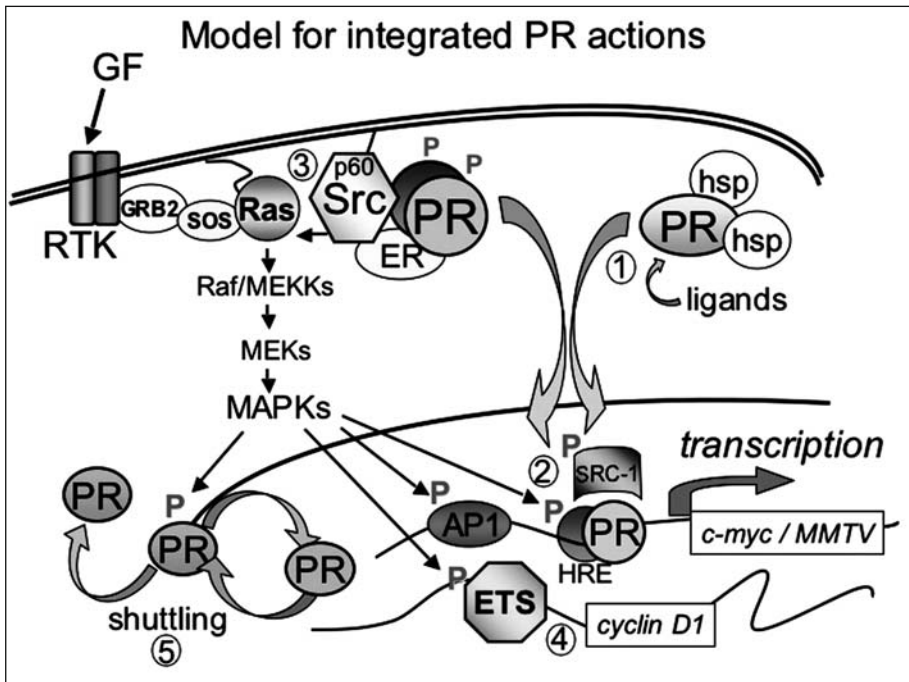


Figure 2. Functional significance of PR phosphorylation. Phosphorylation (P) of specific sites in PRs couple multiple receptor functions, including transcription, nuclear-cytoplasmic shuttling and PR downregulation. 1) Ligand-binding mediates dissociation of heat-shock proteins and nuclear accumulation of PR dimers. 2) Nuclear PRs mediate gene regulation; phosphorylated PRs recruit regulatory molecules that include phospho-proteins and likely function in inter-connected processes (transcription, elongation, localization and turnover). 3) PRs and growth factors activate MAPKs independently via a c-Src kinase-dependent pathway, resulting in positive regulation of PR action via "feed-back" regulation (i.e., direct phosphorylation of liganded PRs or coactivators). 4) Activation of MAPKs by PRs provides for regulation of gene targets whose promoters do not contain PREs and are otherwise independent of PR-transcriptional activities but utilize PR or SR-activated MAPKs. 5) MAPK regulation of PRs mediates nuclear accumulation/shuttling and nuclear export coupled to regulation of PR transcription.

that ER α interacts with the SH2-domain of c-Src, but neither group tested the effects of estrogen on the ability of progesterone to activate c-Src or MAPKs.^{54,56}

Although discrepancies between these two models must be resolved, it is possible that overexpression of SRs in COS-7 cells leads to concentration-dependent effects resulting in the formation of different signaling complexes depending on the presence of other signaling and adaptor molecules. In support of this idea, Wong et al⁵⁷ identified an additional ER-interacting "adaptor" protein, termed MNAR (modulator of nongenomic activity of estrogen receptor), that contains both LXXLL (nuclear receptor binding) and PXXP (SH3-domain binding) motifs. MNAR is essential for ER-Src interaction, but it is not required for progestin/PR-dependent activation of c-Src (D.P. Edwards, personal communication). Taken together, these data indicate that multiple interactions contribute to direct protein kinase activation by SRs and suggest that at least some nongenomic signaling functions of amphibian PR have been conserved in mammals. Interestingly, a separate gene product encoding the putative mammalian homologue of membrane progesterone receptor (mPR), a progesterone-binding G-protein coupled receptor first identified in spotted sea trout oocytes,⁵⁸ has been described. Further studies are needed to determine if mPR plays a role

in progestin-induced “rapid” signaling or if mPR interacts with classical PRs. However, studies with mPR underscore the important concept that binding proteins other than classical steroid receptors may regulate some nongenomic steroid-mediated signaling events.

Integration of Rapid Signaling and Nuclear SR Actions

While its role in mammalian physiology remains unclear, SR-mediated activation of cytoplasmic signaling molecules could theoretically serve to potentiate several nuclear functions of activated SRs (Fig. 2). One mechanism by which amplification of SR nuclear functions might occur is through rapid, direct phosphorylation of SRs and/or their coregulators in response to activation of SR-induced cytoplasmic pathways that coincide with ligand binding. Clearly, such a positive feedback loop would explain the dramatic influence of activated signaling pathways on PR nuclear function. For example, several progestin-dependent functions of PR are MAPK-dependent, including upregulation of cyclins D1 and E, CDK2 activation and S-phase entry.^{22,38,44,59}

Following ligand-binding, most SRs stimulate a transient (3-10 min) activation of MAPKs. However, mitogenic signaling requires sustained (hrs to days) MAPK activation in fibroblast cell models.⁶⁰ Recently, Faivre et al⁶¹ found that in addition to rapid and transient activation of MAPK by progestin/PR-B (5-15 min), progestin-bound PR-B induced subsequent oscillations in MAPK activity that culminated in a sustained (hrs to days) phase of MAPK activation that was EGFR- and c-Src-dependent. Further studies revealed the creation of an autocrine signaling loop, in which PR-B triggered transcriptional upregulation of Wnt-1, leading to activation of frizzled-dependent MMPs and shedding of EGF ligands from the cell surface. This signaling cascade implicates Wnt-1-dependent transactivation of EGFR in response to progestins; PR induced transcriptional upregulation of Wnt-1 and EGFR mRNA was sensitive to inhibition of MAPKs. Additional experiments demonstrated that progestin-induced cyclin D1 upregulation, S-phase entry, or soft-agar growth of T47D breast cancer cells was either blocked by shRNA targeted to Wnt-1 or inhibitors of MAPK, c-Src and EGFR. Finally, progestins failed to stimulate S-phase entry in MCF-7 cells that stably express a PXXP-mutant PR-B, which is unable to bind to the SH3-domain of c-Src and activate MAPK.⁵⁹ Soft-agar growth of T47D cells stably expressing the same PR mutant (PXXP) was greatly attenuated.⁶¹ In addition to c-Src and MAPKs, STATs are important effectors downstream of EGFR signaling. Progestins induce tyrosine phosphorylation and nuclear translocation of Stat5²⁵ and Stat3.³⁰ Proietti et al³⁰ demonstrate that Stat3 phosphorylation and activation by the nongenomic actions of PR is a critical event for breast cancer cell growth; T47D cell growth and tumor growth of progestin-induced mammary adenocarcinomas in BALB/c mice was dependent on PR activation of Jak1 and Jak2, c-Src and Stat3. Taken together, these data indicate that progesterone, via robust PR-B/c-Src signaling to MAPK in combination with PR-dependent transcriptional events, upregulates and activates EGFR signaling to induce cell proliferation. Dysregulation of either arm of this pathway may contribute to uncontrolled proliferation of breast cancer cells.

The extranuclear actions of PRs may contribute to deregulated breast cancer cell growth⁵⁹ and/or increased breast cancer risk,⁸ perhaps by linking steroid hormone action to the regulation of MAPK-regulated genes (i.e., transcription factor targets of MAPK). Similarly, the extranuclear actions of liganded ER α are thought to induce a state of “adaptive hypersensitivity” during endocrine therapy in which growth factor signaling pathways are co-opted by upregulated ER α .⁶² In this model of ER-dependent MAPK activation, liganded ER α associated with the cell membrane interacts with the adapter protein Shc and induces its phosphorylation, leading to recruitment of Grb-2 and Sos, followed by activation of Ras and the Raf-1/MEK/MAPK module. ER α activation of MAPK may explain why many tumors respond well to aromatase inhibitors, yet fail to respond to selective estrogen receptor modulators (SERMs) designed to inhibit ER transcriptional activity. SERMs can act as partial transcriptional agonists of phosphorylated receptors and may not block ER-dependent MAPK activation.⁶² In theory, PR-B or AR in SR-positive breast cancers could participate in MAPK-activating complexes, perhaps bypassing anti-estrogen therapies. Few groups have studied membrane-associated or cytoplasmic signaling complexes containing both ER α and

PR-B or AR.^{63,64} However, AR is frequently (70%) expressed in metastatic breast cancer,⁶⁵ and expression of functional AR defines a sub-set of ER/PR-negative breast cancers.⁶⁶ These studies suggest that it will be important to target SRs that may substitute for ER α in the activation of c-Src-dependent mitogenic signaling cascades.

PR Action and Breast Cancer Cell Growth, Apoptosis and Aggressiveness in Vitro

Among the most controversial issues regarding the role of progestins in breast cancers is their influence, or lack thereof, on tumor cell proliferation. Complicating the interpretation of the results utilizing in vitro breast cancer models of receptor function is the use, in addition to progesterone, of a myriad of different synthetic progestins with activities unrestricted to PR. For example, while the 19-nor progestins—norgestrel and gestodene—enhance MCF-7 cell proliferation, this effect is inhibitable by antiestrogens but not antiprogestins,^{67,68} indicating the lack of involvement of PR signaling. Indeed, cross-reactivity of synthetic progestins at pharmacologic doses with ER has been reported.^{69,70} One explanation of these confusing results is that progestin may interact with different PR isoforms to carry out inhibitory or proliferative functions. Sumida et al demonstrate the growth inhibitory effects of progestins with either PR isoform,⁷¹ and McGowan et al show that overexpression of PR-A sensitized breast cancer cells to progestin-mediated growth inhibition.⁷² In contrast, Moore et al report prolonged proliferative and survival effects of progestins on breast cancer cells.^{73,74}

Flow cytometric studies have also addressed questions of progestin-mediated proliferation by using a single physiological progestin pulse under transiently estrogen deprived conditions. These studies show biphasic effects of progestins in vitro, with cells accelerating through the first mitotic cell cycle then arresting in late G1 of the next cycle.^{27,75} Cycle arrest is associated with decreases in cyclins D1, D3 and E, loss of cyclin A and B and induction of the cell cycle inhibitors p21 and p27. Pulsing with progesterone did not restart proliferation; rather it delayed p21 depletion.²⁷ Similarly, Lin et al⁷⁶ report decreased cell proliferation in response to progesterone in conjunction with upregulation of p21, decreased cyclins A, B1 and D1 expression and downregulation of phosphorylated p42/44 MAPK. Thus, these studies suggest that progestins tend to be anti-proliferative in vitro in mono-layer cell cultures.

Equally confusing are conflicting reports of the effects of PR and progestins on apoptosis in vitro. Several studies report pro-apoptotic effects of progestins concomitant with decreases in expression levels of the anti-apoptotic genes bcl-2 and bcl-X_L.⁷⁷⁻⁸⁰ Antiprogestin/partial agonists, such as RU486, have also been shown to promote apoptosis,⁸¹ but dosage effects confound the interpretation of results.⁸² On the other hand, recent studies suggest that unliganded PR⁸³ and/or progestin-occupied PR⁸⁴ protect cells from damage and apoptosis induced by radiation⁸⁴ or chemotherapeutic agents, such as taxanes,⁸³ doxorubicin or 5-fluorouracil.⁸⁵ Moore et al⁷⁴ report progestin-induced protection of breast cancer cell death accompanied by upregulation of bcl-X_L, but loss of bcl-2. These contradictory in vitro data prevent a definitive conclusion regarding the apoptotic effects of progestins.

Similarly, the effects of progesterone on invasiveness of breast cancer cells in vitro are poorly understood. Many studies show that progestins increase cell invasiveness^{72,83,86} with PR-A exaggerating this phenotype. Sumida et al, however, report that treatment with progestins reduce cell invasiveness.⁷¹

Notably, studies using human breast cancer cell line models (T47D or MCF-7) grown in soft-agar (i.e., as 3-D colonies) clearly demonstrate a proliferative role for synthetic progestins (R5020) or progesterone in response to PR-dependent transcriptional upregulation of Wnt-1.⁸⁷ These results suggest that breast epithelial cells may require a specific architecture (i.e., polarity) for the mitogenic and other “appropriate” gene expression effects of progestins to occur. This architecture is not modeled on plastic surfaces in vitro (i.e., mono-layer cultures). Differences in cell behavior when grown using plastic as mono-layer cultures vs. 3D models have clearly contributed to the controversial area of PR action as a breast cancer cell mitogen. Therefore, we recommend

that future investigations into the effects of progestins on tumor cell behavior utilize 3-D models or in vivo models of PR-positive breast cancer.

Expression Profiling in Vitro

Results from expression profiling of breast cancer cells in vitro are consistent with the results from experimental mouse models, which suggest that the two PR isoforms subservise different functions. In mice—where the PR-A to PR-B ratio is 3:1 compared to humans where it is 1:1—ablation of one or the other PR isoform leads to divergent effects on the mammary gland. PR-A knockout (leaving only PR-B) leads to normal early development,¹⁶ while PR-B knockout (leaving only PR-A) leads to reduced pregnancy-associated lobuloalveolar development and reduced side-branching.¹⁵ On the other hand, overexpression of PR-B causes precocious ductal arrest and inappropriate ductal development,⁸⁸ while overexpression of PR-A causes mammary epithelial cell hyperplasia, excessive ductal branching and a disorganized basement membrane.⁸⁹ To explain these isoform-specific differences, gene profiling studies have been performed in vitro using human breast cancer cells expressing PR-A or PR-B. The first such study used 6 hrs of progesterone treatment in an attempt to identify direct PR target genes.^{14,90} Of 94 genes identified, 65 were regulated only by PR-B, 4 only by PR-A and 25 by both PR isoforms. This regulatory pattern was confirmed in subsequent studies using breast cancer cells with inducible PR-A vs. PR-B treated 6 hrs with progesterone.⁸³ The latter studies also demonstrate that unliganded PR can regulate transcription; CDK2 mediates ligand-independent activation of PR-B via Ser400 phosphorylation (44).

More recent studies used progesterone-treated breast cancer cells that express both PR isoforms.⁹¹⁻⁹³ Analysis of the protein pathways indicate that progesterone suppresses genes involved in proliferation and metastasis,⁹¹ supporting an anti-proliferative role for this hormone. However, a remarkable number of the genes upregulated by progestins encode proteins involved in signal transduction and cell adhesion,^{83,14} lending some support to the concept that progestins/PR may contribute to the dysregulation of pathways important for breast cancer progression that are perhaps not well modeled in vitro. Additionally, the above studies address gene regulation in response to unliganded or liganded PRs (i.e., single hormone exposure). We propose that PR isoforms act as sensors for signal transduction pathways (discussed above) and thus promoter selectivity is predicted to be highly sensitive to phosphorylation events. Further studies will be needed to address alterations in the signature of PR regulated genes in the context of the high kinase activities characteristic of aggressive breast cancer.

Progestins and Antiprogestins in Breast Tumor Models

Antiprogestins

For a time, therapeutic interest in antiprogestins led to many more studies on these drugs than on the biology of progestins themselves in breast tumors. Several rodent and human tumor models have been used to study the efficacy of antiprogestins for endocrine therapy. These include carcinogen- (DMBA or MNU) induced mammary tumors in the rat, serially transplantable MXT (+) mouse and human T61 mammary tumors and MCF7 human tumor xenografts. Tumors in each of these models are ER+ and PR+. Several different antiprogestins, including mifepristone (RU 38.486; Roussel), the Schering compounds onapristone (ZK 98.299 and ZK 112.993) and the ORG compounds (31710 and 31806) effectively inhibit tumor growth 40 to >90%, depending on the drug, dose and model.⁹⁴⁻¹⁰³ Antiprogestins were at least as effective as tamoxifen as a single-line therapy. Combination treatment of established tumors in both the rodent and human tumor models with an antiprogestin and an antiestrogen (tamoxifen or ICI164384) had an additive effect on inhibition of tumor growth.^{95,100} These studies led to speculation that antiprogestins would be useful for endocrine therapies and fueled the notion that progestins induced proliferation. Indeed, several small clinical studies investigated the potential of mifepristone and onapristone as first- or third-line therapies (reviewed in J.G. Klijn et al¹⁰⁴). However, because of apparent liver toxicity

(onapristone), discrepancies among results and the abortifacient properties of these hormones, the testing of antiprogestins for breast cancer therapy has generally been discontinued.

Only two of the above studies examined progestins alone. Megestrol acetate or MPA had no effect on MXT mouse tumors or slightly inhibited DMBA rat tumor growth.^{94,101} This suggests that antiprogestins do not directly antagonize progesterone-mediated tumor growth, even though PR expression was required for inhibition.¹⁰² It is possible that they exert a PR-dependent antiestrogenic effect through ER transrepression¹⁰⁵ or that they suppress effects of unliganded PR.⁸³

Progestins

Human tumor models utilize immune-compromised mice as hosts for “xenografted” breast cancer cell lines. Several ER+ and PR+ human breast cancer cell lines (MCF7, ZR-75, T47D) are grown as solid tumors in this manner.¹⁰⁶ Tumors derived from each of these cell lines are estrogen-dependent and require continuous estradiol administration for growth. They have been widely used as models for studying estrogen-suppression based therapies, such as antiestrogens and aromatase inhibitors.^{107,108} Only a few studies, however, have assessed effects of progestins in these models. Neither MCF7 nor T47D cells grow in response to progesterone in ovariectomized female mice¹⁰⁹⁻¹¹¹ in the absence or presence of estradiol.

In our experience, progesterone or MPA had negligible, nonsignificant growth inhibitory effects in ovariectomized mice bearing T47D xenografts in an estrogenized background.¹¹¹ These data suggest that in hormone-dependent models of human breast cancer, progestins are neither mitogenic nor effective at suppressing estrogen-dependent growth. ER^{neg} and PR^{neg} MDA-231 human breast cancer cells form hormone-independent tumors *in vivo*. If PR was expressed in these cells, progesterone treatment reduced tumor formation.⁸⁶

There is one example of progestin-dependent murine mammary tumor growth. Long-term (10-12 months) chronic treatment of female BALB/c mice with MPA leads to the formation ER+ and PR+ mammary tumors.^{112,113} They are maintained by serial transplantation and have a growth requirement for progestins (either progesterone or MPA) rather than estrogens.¹¹⁴ After serial passage, some tumors acquire progestin independence. Both progestin-dependent and -independent tumors can be inhibited by antiprogestins and antisense oligonucleotides to PR.^{115,116} Whereas most clinical human tumors are ER+ and PR+ and respond to antiestrogen therapies, it is possible that some tumors that originate during long-term HRT or in association with pregnancy may have developed in response to progestins. The BALB/c mice would serve as potential models for these rare tumors.

Progesterone Regulation of BRCA1

Carriers of mutations in the breast cancer susceptibility genes BRCA1 and BRCA2 exhibit a 10-fold higher risk of developing tumors in hormonally responsive tissues, such as the breast and ovaries (cumulative risk of 85-90% by age 70) compared to the general population.¹¹⁷ BRCA1 mutant breast tumors have a poor nuclear grade, high frequency of p53 mutations and are more often ER- and PR-negative compared to sporadic cancers. Because oophorectomy of premenopausal women reduces breast cancer risk substantially (>40%),¹¹⁸ linkages between steroid hormones and BRCA1 tumor types have been sought since their discovery.¹¹⁹⁻¹²¹ Fibroblasts from *Brca*^{-/-} knockout mice that are also p53^{-/-} exhibit ligand-independent activation of ER and PR-dependent transcription;¹²² see also Rosen et al.¹¹⁷ Haploinsufficiency of BRCA1 may be a deleterious state that initiates alterations in steroid hormone receptor expression and tumor mitogenic response.¹²³ Poole et al¹²⁴ report the accumulation of lateral branching and extensive alveologenesis in the mammary glands of nulliparous BRCA1/p53-deficient mice. PR, but not ER, were overexpressed due to a defect in their proteasome-dependent degradation. Notably, treatment of these mice with the PR antagonist mifepristone (RU486) blocked mammary tumorigenesis. These provocative studies suggest that antiprogestin therapy may help prevent the development of breast cancer in individuals with BRCA1 mutations.

General Steroid Receptors and Breast Cancer

A recent study¹²⁵ described the steroid receptor assay results of 54,865 patients with stage I–IIIA breast cancers. Their biopsy or mastectomy specimens were sent to two central laboratories that performed identical assays, monitored with tightly controlled quality control procedures. The authors report that ~82% of breast cancers were ER+ and of these ~71% were also PR+. Thus among all breast cancers, ~58% expressed both ER and PR. It is now well established that independent of treatment modalities, women with steroid receptor positive tumors live longer than their receptor negative counterparts. Large studies with long-term follow-up, such as those from San Antonio or the NSABP, indicate a 10% survival advantage for patients with receptor positive disease.¹²⁶ Positive hormone receptor status is an independent predictor of outcome and augurs a more favorable prognosis even after controlling for patient age, disease stage, tumor grade, histology, race/ethnicity and US geographical distribution.

Progesterone Receptors and Tamoxifen Responsiveness

The independent role of ER-positivity as a marker of good prognosis and responsiveness to endocrine therapies has been appreciated since the early 1970s. Resistance of a subset of ER+ tumors to endocrine therapies may be due to aberrant estrogen signaling in ER+ tumors that lack PR.¹²⁷ Indeed, compared to ER+ and PR– tumors, pretreatment PR-positivity in ER+ tumors is associated with improved outcome prediction as shown by 5 year disease survival rates¹²⁸ and by improved response to such adjuvant endocrine therapies as tamoxifen.^{128–133} However, not all studies have demonstrated a value for PR, due perhaps to assay variability.¹³⁴ The presence of both ER and PR in metastatic disease has also been shown to predict improved response to tamoxifen treatment.^{131,135}

Clinical Significance of PR-A vs. PR-B: Two Subsets of ER+, PR+ Tumors?

We first showed that human breast cancer cells express two forms of PR, the PR-A and PR-B isoforms.¹³⁶ Despite having a similar primary amino acid structure over the majority of their length, these receptors regulate entirely different gene subsets.^{83,14} The clinical implications of this remain under investigation. Studies using monoclonal antibodies show that PR-A and PR-B colocalize in the same cells in normal endometrium¹³⁷ and breast cancers,¹³⁸ further adding to the complexity of analyzing expression ratios of the two isoforms by IHC. By immunoblotting, their ratio changes during malignant progression, with approximately equimolar levels of PR-A and PR-B in normal human tissues, but aberrant PR-A:PR-B ratios in breast cancers. An immunoblotting study by Graham et al¹³⁹ of 202 PR+ breast cancers showed a median PR-A:PR-B ratio of ~1.3 (close to equimolar), but with outliers ranging between 0.04 (essentially PR-B+) to ~180 (essentially PR-A+) in a significant number of tumors. These authors concluded that when ratios are aberrant, the PR-A isoform tends to be in excess,^{140,141} and tumors tend to be less differentiated.¹⁴¹ We¹⁴² studied the association between PR-A:PR-B ratios and clinical outcome in 297 ER+, axillary node-positive patients, using MAb 1294 for immunoblotting. Eighteen percent of tumors had more than a 2-fold excess of PR-B over PR-A; 10% had more than a 2-fold excess of PR-A over PR-B. We concluded that high PR-A levels were due to loss of PR-B. Our studies also included clinical data showing that tamoxifen-treated patients with high PR-A:PR-B ratios were 2.76 times more likely to relapse. Thus, clinical studies that have addressed the issue of PR isoforms agree that an excess of PR-A is harmful. We suggest that patients with PR-A rich tumors may represent an ER+/PR+ subgroup with intrinsic insensitivity to tamoxifen and perhaps to other selective ER modulators. Growth factor signaling is tightly linked to tamoxifen resistance. Notably, Ser294 phosphorylated PR-B is hypersensitive to low progesterone concentrations and thus degrades very rapidly relative to PR-A, which is hypo-phosphorylated at this site (discussed above); hyperactive but unstable PR-B relative to PR-A may contribute to increased PR-A/PR-B ratios in a subset of breast cancers. In this setting, targeting PR-B and relevant kinases would seem appropriate, but remains untested clinically.

Concluding Remarks

Studies aimed at defining a proliferative role for progestins in breast cancer models remain controversial, but have perhaps been hindered by observations made with liganded receptors in the absence of controlled inhibition or activation of alternate signaling pathways. In the context of multiple signaling inputs, PR clearly coordinates receptor responses to growth factors and steroid hormones. The newly discovered ability of SRs to activate kinase pathways classically defined as key regulators of cell growth underscores the concept that activation of signal transduction pathways is an integral feature of SR action. This aspect of SR function is likely to play an important role in cancer progression and the development of resistance to endocrine therapies.⁶² Targeting the relevant protein kinases (c-Src, MAPKs and CDKs) as an integral feature of SR (PR, ER) action should provide significant improvements over the use of traditional SR-blocking strategies for advanced or progressive breast cancers.

Acknowledgements

We thank Michael Franklin for editorial comments and Dean Edwards (Baylor College of Medicine) and Natalie Ahn (University of Colorado, Boulder) for generous gifts of reagents. Studies contributed by Carol Lange's laboratory were supported by NIH grants DK53825/CA123763 and CA116790. These studies were also supported by grants to Britta Jacobsen (Department of Defense 06-1-0503), Carol Sartorius (Susan G. Komen Foundation: BCTR0402682) and grants to Kathryn Horwitz from the NIH (CA26869), the Avon Foundation, the Breast Cancer Research Foundation and the National Foundation for Cancer Research.

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