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# Extranuclear Signaling by Sex Steroid Receptors and Clinical Implications in Breast Cancer

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# Abstract

Estrogen and progesterone, play essential roles in the development and progression of breast cancer. Over 70% of breast cancers express estrogen receptors (ER) and progesterone receptors (PR), emphasizing the need for better understanding of ER and PR signaling. ER and PR are traditionally viewed as transcription factors and directly bind DNA to regulate gene networks. In addition to nuclear signaling, ER and PR mediate hormone-induced, rapid extranuclear signaling at the cell membrane or in the cytoplasm which initiates downstream signaling to regulate either rapid or extended cellular responses. Specialized membrane and cytoplasmic proteins may also initiate hormone-induced extranuclear signaling. Rapid extranuclear signaling converges with its nuclear counterpart to amplify ER/PR transcription and specify gene regulatory networks. This review summarizes current understanding and updates on ER and PR extranuclear signaling. Further investigation of ER/PR extranuclear signaling may lead to development of novel targeted therapeutics for breast cancer management.

## Keywords

Extranuclear signaling; nongenomic signaling; estrogen receptor; progesterone receptor; growth factor signaling; breast cancer

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# Introduction

#### 1. Nature of extranuclear steroid hormone receptors

**1.1. Classical estrogen receptor structure and activity**—Early concepts of estrogen action proposed the involvement of both nuclear and extranuclear estrogen receptors (ER) driving ER activity, however most studies have focused on nuclear ER and its transcriptional activity [1–5]. There are two major types of estrogen receptors, ERa and ER $\beta$ , each produced from different genes with similar but not identical structure [2, 6]. On binding estradiol-17 $\beta$  (E2), ERa undergoes a change in conformation of the ligand-binding domain (LBD) to form a novel hydrophobic surface to modulate binding of coactivators and corepressors [2, 7, 8]. As a phosphoprotein, ER undergoes phosphorylation at serine and tyrosine residues after activation by E2 binding, and this contributes to receptor activity and DNA binding [9–11]. E2-bound ERs function as transcription factors by binding as homodimers to estrogen response elements (ERE) [7, 8] (Fig. 1).

Interestingly, about one third of E2-induced genes lack functional EREs, and estrogens indirectly regulate transcription of these genes by modulating activity of other transcription factors such as activator protein-1, Elk-1, serum response factor, cyclic AMP-responsive element binding protein, nuclear factor-kappaB (NFrkB), and signal transducers and activators of transcription [7, 8, 12]. Blockade of ER signaling by interfering with E2 binding to ER is the basis for antiestrogen, tamoxifen, action. Tamoxifen is a partial agonist, termed a selective estrogen receptor modulator (SERM), that limits E2-stimulated proliferation in breast tumors [13]. Fulvestrant is an ER antagonist that downregulates cellular levels of ER and, unlike tamoxifen, has no agonist activity [14–18]. Further, inhibitors of aromatase (anastrazole, letrazole, exemestane) reduce E2 biosynthesis in target tissues and shows some advantages over tamoxifen as antitumor therapies [19–22]. Recent findings based on molecular analyses of ER in BC metastases show that ER mutations can emerge in patients treated with extended courses of endocrine therapy, in particular, endocrine resistance tumors occur mainly with mutations in critical residues in the LBD of ERa and  $\beta$  [23, 24]. The precise role of ER $\beta$  in ERa-positive and ERa-negative BC subtypes remains to be determined (Hamilton et al., 2015). ER $\beta$  is often reported to exert antiproliferative and pro-apoptotic effects in BC cells that co-express ERa [25, 26], while ERβ may promote proliferation and block apoptosis in BC cells that lack ERα expression [27-29].

**1.2. Background on Extranuclear ER structure and activity**—Estrogen and other steroid hormones are well known to accumulate and are retained in responsive cells [30] by specific interactions with cellular macromolecules, initially considered to be located outside the nucleus [31, 32]. Binding to the target-cell receptor was the proposed mechanism to initiate transfer of E2 to the nucleus where the ER/E2 complex promoted expression of phenotypic changes [31, 33, 34]. The genomic hypothesis of steroid hormone action has generally prevailed as the exclusive mechanism of hormone action due to the pioneering work by molecular and cellular biologists, whose studies extended details of these concepts to understand the late nuclear actions of steroid hormones in their target cells. Development of monoclonal antibodies for specific binding to nuclear ER consolidated this concept and

has led to important clinical benefits for patients afflicted with ER-expressing BCs since these ER-positive tumors can be treated more effectively with targeted antiestrogens [1-3]. However, the emphasis on nuclear steroid actions overshadowed parallel observations on early extranuclear receptor-mediated signaling. More than 40 years ago, Pietras and Szego [35, 36] and Szego and Davis [37] first described rapid estradiol-induced actions, including changes in calcium fluxes and cyclic AMP generation that occurred in seconds to minutes after exposure of target cells to this sex steroid. These authors further provided evidence of estrogen binding with high affinity and specificity to a receptor molecule partially purified from target cell membranes [38, 39]. In the case of some hormone responses, rapid extranuclear effects may be sufficient to elicit a cascade of intracellular signals to alter specific cellular functions. One such example is the localized release of nitric oxide which is secondary to an instantaneous surge of calcium that occurs in endothelial cell response to E2 [40–42]. These events result in rapid vasodilation, thus bypassing the long (hours) transduction pathway response due to nuclear signaling. The local effects of E2 on the rapid electrophysiological activities of neurons are another examples of extranuclear receptor effects [43-45].

In evolutionary terms, steroid hormone recognition at the surface membrane appears to have been a primary response pathway in more primitive cells and organisms [46–48]. These early reports suggested that conserved functions of steroids from primitive cells to humans resulted from actions in part at the cell membrane. The concept of specific membrane-associated binding sites for steroid hormones is now supported by many independent laboratories [49–54]. Such membrane proteins appear to constitute a fraction of total receptor molecules (~5%) in target cells. Ultrastructural studies reveal extranuclear immuno-reactivity for ERa at membrane sites along dendritic spines and axon terminals of neurons [55]. Moreover, E2 may promote growth of mammary tumor cells, potentially in concert with ERE-dependent transcription, by stimulating downstream p42/p44 MAP kinase pathways [52]. Ligand-independent activation of steroid hormone receptors also occurs via peptide signaling systems. For example, ER can be activated in the absence of E2 through phosphorylation of tyrosine residues by EGF-stimulated MAP kinase [56–58]. Any comprehensive model of steroid hormone action must account for these important cellular interactions.

In vertebrates, the distribution of most sex steroid receptors favors nuclear localization, thereby allowing receptor binding to promoters and steroid receptor response elements in DNA of target cells to regulate specific gene expression. Although it is often assumed that steroid hormones enter target cells by passive diffusion, biophysical studies show that steroid hormones are largely lipophilic compounds that partition within the hydrocarbon core of lipid bilayer membranes. Steroid hormones appear to enter target cells by a saturable, temperature-dependent membrane-mediated process [59, 60]. Analysis of [<sup>3</sup>H]-estradiol-17 $\beta$  uptake in target cells was reported using analytical cell fractionation at progressive time periods, beginning within 10 seconds of exposure [61]. In these studies, E2 interacted specifically with membrane proteins in uterine target cells and underwent rapid internalization, resulting in intracellular delivery of a portion of E2 and its associated receptor protein. Concomitant with a decline in plasma membrane and endosomal subfractions, a significant amount of labeled E2 was shown to occur in Golgi and lysosome-

mitochondrial compartments before later peaks shown in nuclear compartment. Similarly, membrane-associated binding sites for E2 may mediate rapid effects of E2 that contribute to proliferation of BCs [57]. After controlled homogenization and fractionation of human MCF-7 BC cells, the bulk of specific E2 binding was found in the nuclear fractions, but a significant portion of specific, high-affinity E2 binding-sites was also enriched in plasma membranes. These E2 binding-sites in BC cells co-purify with 5'-nucleotidase, a plasma membrane-marker enzyme and were free from major contamination by cytosol or nuclei. Electrophoresis of membrane fractions allowed detection of a primary 67-kD protein and a secondary 46-kD protein recognized by E2 and by a monoclonal antibody directed to the LBD of nuclear ER. Independent experiments also showed the presence of endogenous membrane and nuclear ERa in human BC cells that were identical by mass spectrometry analysis of subcellular proteins isolated from E2 affinity columns [62]. Related reports of steroid receptor localization in mitochondria requires further investigation, particularly concerning steroid regulation of metabolism in target tissues [51]. Using antibodies to nuclear ERa, Pappas et al. [63] identified a plasma membrane-associated protein by established immunohistochemistry (IHC) methods. Furthermore, E2 binding to a surface membrane protein was reduced significantly by prior treatment with antisense oligonucleotides to suppress ERa expression [64]. A number of independent reports confirmed ERa association with plasma membranes by use of controlled homogenization with quantitative subcellular fractionation [38]. Specific antibodies were directed to different domains of nuclear ERa in intact breast [56, 63, 65], NSCLC [66, 67], and pituitary tumor cells [68], as well as in nonmalignant vascular cells [40]. In addition, conformation of E2 binding of plasma membrane proteins was established through ERa knockout models in astrocytes [69]. Although ERs localize predominantly in tumor cell nuclei, a significant pool of ERs has been shown to localize in extranuclear sites in archival BC and NSCLC cells [41, 66, 70, 71]. Thus, important actions initiated by membrane-associated forms of ER may play a collaborative role with liganded-ERs in the nucleus to promote signaling for hormonemediated proliferation and survival of BCs.

The *ESR1* gene encodes for a major 66-kD transcript and a minor 46-kD isoform lacking portions of the NH2-terminal region of full-length ERa [22, 72]. The 46-kD ER also occurs in membranes of endothelial [22] and breast [73] cells, possibly forming part of a signaling complex. To assess the nature of membrane ER, nuclear ERa gene was transfected in ERnull Chinese hamster ovary cells, and this resulted in cellular expression of both membrane and nuclear ERs, and the transfected cells responded to E2 with rapid signal transduction [74]. Independent studies also showed that transfection of ESR1 and ESR2 genes resulted in expression of both nuclear and membrane-localized receptors, confirming that both forms originate from the same gene transcripts [52, 73, 74]. Similar studies were done with progesterone (PR) and androgen receptor (AR) demonstrating that non-nuclear forms of these proteins or splice variants originate from the same gene [75-78]. Studies based on knockdowns of ERa by small interfering RNA [68, 73] or knockouts of both ERa and ERB in vivo [62] offer additional support for the hypothesis that membrane and nuclear ERs share a common origin. Further, membrane ERs do not occur in ER-negative MCF-7 BC subclones that lack nuclear ER [73]. These cells, unlike ER-positive MCF-7 cells, do not show rapid estrogen-induced phosphorylation of steroid receptor coactivator AIB1 [73].

Importantly, studies using mass spectrometry provide evidence that peptides derived from ERa occur in membrane fractions prepared from BC and vascular endothelial cells [79]. Together, these data indicate that membrane-associated ER derives predominantly from the same gene as nuclear ERa.

There is evidence that endogenous  $ER\beta$  also localizes to plasma membranes in some tissues including BC [71]. ER $\beta$  was first reported in 1996 and is the second major receptor that mediates some actions of E2 in various organs [6, 80]. ERa and ER $\beta$  are encoded by different genes, yet ERB has 96% homology with ERa in the DNA-binding domain and 60% homology in the LBD. However, ER $\beta$  is not identified in standard assays for ERa. Many studies also indicate that truncated forms of ERa or alternative steroid-binding proteins are expressed in a variety of organs. ERa isoforms, 46-kD [22] or 36-kD [24, 81, 82] in size, have been reported at the cell membrane, especially in BC cell lines. ERa isoforms of 46- and 36-kD are splice variants [22, 83, 84] but are not generally as abundant as ERa-66 kD in cells expressing both receptor forms. In comparison to the full length ERa-66 kD isoform, the ERa-36 kD isoform lacks the AF-1 and AF-2 transcription activation domains. Yet, the truncated ERa-36 kD isoform possesses an altered LBD and an intact DNA-binding domain, consistent with the report that ERa-36 kD lacks intrinsic transcriptional activity but can mediate extranuclear signaling [24]. ERa-36 kD is activated by both E2 and antiestrogens and can be detected in ERa-66 kD-positive and -negative BCs [85]. ERa-36 kD has been shown to mediate E2 and antiestrogen signaling via the MAPK/ERK pathway and stimulate cell proliferation [84]. ERa-36 kD is expressed in triple-negative breast cancer (TNBC) cell lines and associates with EGFR [24]. In addition to TNBCs, ERa-36 kD is overexpressed in apocrine and adenoid cystic carcinomas, tumors for which therapeutic treatment options are currently not available [24, 86]. Limited studies indicate ERa-36 kD overexpression associates with increased breast tumor growth, tamoxifen resistance and metastasis [87, 88]. However, efforts to evaluate the clinical utility of ERa-36 kD without the availability of specific and validated antibodies are daunting.

**1.3. Alternative estrogen-responsive membrane proteins**—It is reported that an orphan 7-transmembrane G-protein-coupled receptor, GPR30, serves as an estrogen membrane receptor and mediates E2 signaling from plasma membrane [89, 90] or endoplasmic reticulum [91, 92]. In some cell types, especially BC cells, there may be some coordination between GPR30 or other G-protein-coupled receptors and membrane ERa as part of a larger membrane signaling complex, thereby transmitting rapid-E2 downstream signals [93–95]. The function for GPR30 was first identified from studies showing MAPK activation by E2. Responses were detected in BC cells expressing GPR30 but not in cells lacking GPR30 [89]. Interestingly, ER-antagonists such as tamoxifen and fulvestrant also bind GPR30 [95], consistent with earlier studies showing that these agents are agonists for GPR30 [89]. Independent results indicate that tamoxifen activates PI3K through GPR30 but not ERa, suggesting a possible role of GPR30 in tamoxifen-resistant BCs and/or in the increased incidence of endometrial cancers in women treated with tamoxifen [96, 97]. Further, reduction of GPR30 expression has been reported to prevent growth stimulation of TNBC cells (that lack ERa) by E2 [98], conversely, other reports suggest GPR30/GPER expression correlates with better prognosis in ERa-positive BC patients [99, 100].

Despite these observations, a number of important questions remain about the physiological function of GPR30. It is not clear if GPR30 has redundant or overlapping functions with ERa, ER $\beta$  or other ER isoforms [101], or fully independent responses. From available reports, GPR30 is expressed in BC cells that do not express ERa including TNBC cells. It is not known whether drugs that selectively target GPR30 versus ERa and ERB would be superior to drugs currently available for BC treatment. Although ERa and GPR30 may both occur in membranes of target cells, some investigators have not confirmed a functional linkage between membrane-localized ERa and GPR30 in BC and other cell types [5, 51]. Further, it remains to be confirmed whether estrogen binds with high affinity and specificity to GPR30, recently termed GPER1. Saturation binding by [H<sup>3</sup>]-E2 to the endogenous membrane GPR30 in mouse kidney and ERa-negative BC cell line SKBR3 showed relatively minute amounts of ligand specific binding [51, 95, 102]. In ERa-negative cells expressing GPR30/GPER1, studies have not confirmed specific binding by or signaling in response to E2 [62, 103]. Reasons for these experimental differences are unclear. Further, 2methoxyestradiol, a final end product of estradiol metabolism with minimal to no binding affinity for either ERa or ER $\beta$  [104] appears to bind GPR30 and induce cellular responses primarily via GPR30 [105]. Notably, independent laboratories have created GPR30knockout (KO) mice [106, 107] and report few phenotypic differences under basal conditions and no overlap with the phenotypes observed in ERa- or ERβ-knockout mice. At present, the endogenous primary ligand for GPR30 has yet to be identified. With recent reports of a very selective, non-steroidal GPR30 agonist [108] as well as a GPR30 antagonist [109], future studies of GPR30 function should be facilitated using these agents. Finally, another distinct membrane ER form under investigation termed Gq-mER has been shown to rapidly activate kinase pathways to stimulate downstream actions in central nervous system neurons [45].

1.4. Membrane association of estrogen receptors—Although more evidence is emerging on estrogen membrane receptors, how ER associates with membranes remains a challenging question. While ERa has many hydrophobic regions on Kyte-Doolittle plots, there appeared to be no readily apparent trans-membrane, glycosylphosphatidylinositolanchor, or PDZ domains to foster membrane association [72]. However, it is reported that attachment of palmitic acid to an internal cysteine residue in ER promotes receptor trafficking to the plasma membrane [110]. Palmitoylation of ERa at cysteine-451 in the mouse and cysteine-447 in humans is required for about 5% of the total cellular ERa pool to localize to plasma membrane [66, 78, 83, 110]. A highly conserved nine-amino-acid motif in the LBDs of human ERa and ERß seems necessary to mediate receptor palmitoylation and subsequent steroid signaling [78]. Several studies have clarified how subpopulations (~5% of total cellular receptors) of newly synthesized ER, as well as PR, are directed to plasma membrane via palmitoylation at specific cysteine residues. Further, membrane ER localization via palmitoylation is reported to be important for delaying ER degradation and for estradiol-induced ER occupancy at ERE sites, suggesting that ER membrane localization and extranuclear signaling may be a prerequisite for ER transcriptional activity [66, 78, 110, 111]. Palmitoylation at specific cysteine residues occurs from enzymatic actions of specific palmitoylacyltransferase proteins (PATs) DHHC-7 and -21, resulting in disulfide bond linkage of the palmitic fatty acid with ERa [112]. Importantly, all of these actions are

highly conserved for membrane localization of both PR and androgen receptors (AR) [78, 112, 113]. Further, ER $\beta$  undergoes palmitoylation and subsequent membrane localization in some cell types thereby enabling rapid signaling *in vitro* and *in vivo* [114].

To better understand the impact of membrane-localized ER signaling for cell function, unique bioengineered mouse models have shown that membrane ER is required for normal organ development and functions. For example, in membrane ER-only mouse (MOER) model, ER-null mice were engineered to express the LBD of ERa targeted exclusively to the membrane of all cells. E2 treatment of cells derived from various organs of MOER mice triggered rapid activation of various kinase pathways and rapid generation of cyclic nucleotides, indicating that the membrane ER pool is sufficient in mediating acute signal transduction of E2 [115, 116]. In contrast, another mouse model was developed with selective loss only of membrane-localized ER but retaining abundant nuclearlocalized receptors. Cells from these nuclear ER-only mice (NOER) failed to respond to E2 with the ERK and PI3K/AKT signaling as seen in wild-type ER mouse cells. Consequently, E2 stimulation of genes regulated by both rapid membrane signaling and nuclear ER transcription was significantly reduced. Of note, NOER mice lacking membrane ERa also exhibited several developmental abnormalities including infertility, abnormal pituitary hormone regulation, hypoplastic uteri, abnormal ovaries, stunted mammary gland ductal development and trabecular bone formation [115, 117]. These abnormalities were rescued in heterozygous NOER mice that were comparable to wild type mice in this model. Further work with such unique mouse models [118] and cell constructs will help to decipher the biological roles of membrane-associated and nuclear steroid hormone receptors [118].

To understand the nature of steroid receptor association with cell membranes, it is important to consider current concepts of supramolecular membrane organization (Fig. 2). The present view of the lateral organization of plasma membrane constituents was revised significantly from the original fluid-mosaic model wherein membrane proteins were considered to diffuse freely in a sea of lipids [119]. Rather, current data suggest the existence of membrane macro- and micro-domains that concentrate key signaling molecules for efficient coupling to effectors. The concept of a 'signaling platform' has been advanced to characterize a structure in which many different membrane-associated components are assembled in a coordinated fashion. Plasma membrane microdomains termed 'lipid rafts' arise from the phase behavior of lipid components. Raft association may concentrate receptors to interact with ligands and effectors on either side of the membrane, thus facilitating binding during signaling. Contrary to earlier notions, recent studies suggest that classical receptors for ER, PR and AR contain two transmembrane helices within their LBDs and harbor pore-lining regions in the plasma membrane that potentially regulate ion permeability [120, 121]. Na+, K+-ATPase, well-known to maintain an electrochemical gradient across the plasma membrane, similarly serves as a receptor for cardiotonic steroids [122]. In addition to the full-length ERa-66kD isoform, an N-terminus truncated ERa isoform (ERa-46kD) is reported to be a potential integral transmembrane protein that plays a key role in rapid endothelial responses to E2 [22].

While lipid rafts are moving cholesterol-rich domains in membranes that provide a matrix for signal transduction [73, 83, 123], caveolae and caveolae-like rafts are specialized forms

of lipid rafts containing the structural proteins caveolin or flotillin, respectively. Caveolae are known to concentrate and assemble components of several signal transduction pathways (Fig. 2). As lipid rafts, caveolae are rich in cholesterol and sphingolipids, but, unlike rafts, they are lined intracellularly with clusters of caveolin protein, a cholesterol-binding molecule that contributes to membrane lipid organization. The list of caveolae-associated molecules includes receptor tyrosine kinases, G-protein-coupled receptors, protein kinase C, components of the MAPK pathway, and endothelial nitric oxide synthase (eNOS). In one such example, subpopulations of ERs are localized to caveolae in endothelial cells. In caveolae of plasma membrane isolated from endothelial cells, E2 directly stimulates ERs coupled with eNOS in a functional signaling complex to regulate local calcium levels and blood vessel contractility [124]. In another example, subpopulations of ERs are also enriched in caveolae in human BC cells and are coupled with downstream signaling pathways that promote tumor progression [73].

The precise nature of membrane-associated steroid receptors remains elusive, primarily because full structural characterization of receptors isolated from membranes is incomplete. Nonetheless, ER from target cell plasma membranes appears to be a protein species with high-affinity and saturable binding specific for E2. Further, current findings as detailed above suggest that membrane receptors for steroid hormones are, in certain instances, transcriptional copies or variants of nuclear receptors, and, in other instances, are products apparently unrelated to these [63]. Steroid receptors in membranes may also occur in multimeric complexes with other transmembrane molecules coupled to specific signaling cascades (Fig. 2). Apart from membrane receptor localization with caveolins and flotillins [73, 78], association with coactivator modulator of extranuclear actions of ER (MNAR)/ proline-, glutamic acid-, and leucine-rich protein-1 (PELP1) [111] or transmembrane growth factor receptors as docking sites [125] are reported. It may well be that multiple forms of protein receptors for steroids coexist in plasma membranes, thus complicating efforts to isolate and characterize individual binding species in membranes. As suggested before, nature may have designed more than one way to open a given lock [46].

**1.5 Classical nuclear progesterone receptor (PR)**—The classical PR is a steroid hormone-activated receptor belonging to the nuclear receptor superfamily, type 1 [126]. PR is expressed as two major isoforms designated PR-A (94kDa) and PR-B (120kDa), both produced from the same gene on chromosome 11 band q13 [127]. Each isoform originates from two transcriptional (AUG) initiation sites from two distinct E2-inducible promoter regions [128, 129]. The difference between the two isoforms is the presence of the unique 164 amino acids upstream of the NH<sub>2</sub>-terminal region of PR-B, absent in PR-A. This unique-PR-B NH<sub>2</sub> -terminal allows for different conformational, isoform-specific proteins that contribute to different interacting-binding partners and co-factors mediating distinct properties of the PR-B isoform [130, 131]. In common, the structures of PR-A and PR-B contain a C-terminal hormone-binding domain, a DNA binding domain, a hinge region, a nuclear localization sequence (NLS), and a shared NH<sub>2</sub>-terminal region. Two transcriptional activation domains TAF1 and TAF2 are present in the shared PR isoform sequence: TAF1 is located in the N-terminal, and TAF2 is located in the hormone-binding domain at the C-terminal [126, 132]. A third TAF (TAF3) is present in the unique-PR-B region [126, 133].

Another important feature of PR is the presence of a proline-rich SH3 motif in the N-terminal of both PR-A and PR-B, important for rapid, cytoplasmic (non-genomic) signaling, including binding of c-Src tyrosine kinases [134].

Traditionally, PR is well recognized for its role as a ligand-dependent transcription factor [126]. PR is part of a multi-protein chaperone complex and upon ligand binding, PR disassociates from this complex and forms homo- and hetero-dimers and interacts with co-activators, which then bind to specific progesterone response elements (PRE) found within the promoter regions of gene targets [132]. Upon progesterone binding, PR isoforms are phosphorylated with equal affinity and translocate to the nucleus [135]. Historically, PR-B was understood to function predominantly as an activator of transcription, while the PR-A isoform was mainly defined as an inhibitor of the transcriptional activity of PR-B and potentially other nuclear receptors (NRs) including ERa, glucocorticoid receptor, mineralocorticoid receptor and androgen receptor [136]. Albeit, when each isoform was independently expressed [137] or when the ratio of PR-A and PR-B was altered [138, 139], PR-A and PR-B demonstrated distinct and overlapping transcriptional activator.

Ligand-activated PR mediates major physiological roles in female reproduction [133]. Although expression of both isoforms is present in target cells, PR-B is essential for breast reproductive functions, and ablation of PR-A has no detrimental effect on breast function as described in mice; conversely PR-A is essential for normal ovarian functions [140]. PR is a well-recognized ERa-mediated target, but progesterone-bound/PR has also been shown to modulate ERa gene expression programming at the chromatin level [141]. In addition, a third PR isoform has been described and designated PR-C. PR-C isoform is truncated in the DNA-binding domain and is demonstrated to inhibit the PR-B isoform [142]. However, physiological roles of PR-C isoform remains to be further characterized [143].

In addition to the well-documented function of PR as a nuclear transcription factor, extranuclear function of PR have also been reported [76]. Progesterone binding to extranuclear PR results in rapid activation of the Src/MAPK signaling pathway [76]. This extranuclear activity occurs within seconds to minutes of progesterone binding, in the presence or absence of ERa [144]. These extranuclear actions of the PR are described in detail in section 2.2. *Rapid PR extranuclear signaling mechanisms.* 

**1.6 Membrane progesterone receptors (mPRs) and PGRMC1**—Emerging in the progesterone signaling field is the increasing complexity of proposed progesterone receptors (PR) which extend beyond the classical intracellular PRs. Two types of putative membrane-specific progesterone receptors that are unrelated to classical PR-A and PR-B and with very different structures have been reported. These include the membrane progesterone receptor (mPR) with a molecular weight (MW) of ~40 kD and the progesterone receptor membrane component 1 (PGMRC1) with a MW of 26–28 kD [145].

**1.6.1 Membrane progesterone receptors (mPRs):** Although mPRs have similar functional characteristics to the G-coupled receptor superfamily, they belong to the progesterone/ progestin and adipoQ receptor (PAQR) class II family, which are a class of transmembrane

proteins [146]. mPRs consist of 7-transmembrane domains with the N-terminal domain localized on the extracellular side of the plasma membrane [147]. There are five mPR subtypes identified, three in mammalian cells (mPR $\alpha$ , mPR $\beta$ , and mPR $\gamma$ ), and two characterized only in yeast (mPR $\delta$  and mPR $\epsilon$ ) [148].

Zhu and Thomas were the first to identify ligand-bound mPRs, distinct from classical PR, that activate rapid signaling of G-proteins and MAP kinase pathways [148, 149]. Direct interactions between G-proteins and mPRs were detected in co-immunoprecipitation studies, and disruption of G-protein-mPR interactions reduced progestin binding to mPRs and loss of ligand-dependent G-protein activation [147]. Given some disparities in the binding affinities of different progestins to nuclear PR versus mPR, mPR-specific agonists were identified, and these can now be exploited to study mPR-specific functions [150].

Thomas and colleagues described the first non-nuclear mPR on fish oocyte membranes [151]. Subsequently, mPR homologues with characteristics of fish mPR were identified in different species including humans [150–152]. In humans, mPR homologues were found in normal tissues, including breast, reproductive, neuroendocrine and immune cells, and were also detected in BC cell lines [151–153].

Three of the mPRs subtypes, mPR $\alpha$ , mPR $\beta$  and mPR $\gamma$ , have been detected in breast and ovarian cancer cells and tissues with mPR $\alpha$  identified as the predominant subtype [154, 155]. To date, mPR is reported to occur in most BC cell lines and biopsies tested [149] and is associated with cell survival. Specific mPR-agonists or low doses of progesterone in PR-negative, mPR-positive BC cells have been shown to inhibit apoptosis, potentially via activation of MAPK p42/44 signaling pathways [155]. In aggressive TNBCs expressing the mPR $\alpha$  subtype, specific knockdown of mPR $\alpha$  expression is reported to block tumor growth and metastasis *in vivo* [156]. A recent study further showed that expression and tumor size, suggesting a significant role for mPR $\alpha$  in BC progression (Wu, Sun, Wang et al., 2016). More studies are needed to determine the contribution of mPR and mPR-subtypes on BC development and their potential role in resistance to endocrine therapy.

**1.6.2 Progesterone receptor membrane component 1 (PGRMC1):** A putative membranespecific progesterone receptor, unrelated and distinct from the classical nuclear PR, has recently been identified. Human PGRMC1 is a member of the membrane-associated progesterone receptor (MAPR) family of cytochrome  $b_5$  (cytb5)-related heme/steroidbinding proteins, and is located on chromosome X (Xq22-q24) [157, 158]. PGRMC1 is a transmembrane protein with binding sites for Src homology 2 (SH2) and Src homology 3(SH3) domain-containing proteins [158]. PGRMC1 forms a stable dimer when bound to heme [159, 160], and this complex is necessary for downstream signaling initiated through multiple binding partners including multiple cyP450 proteins, plasminogen activator inhibitor 1 mRNA binding protein (PAIR-BP1), insulin induced gene 1 (Insig), and sterol regulatory element-binding protein (SREBP) [161]. Although the name implies that PGRMC1 binds progesterone, there is an ongoing debate whether PGRMC1 is a direct progesterone receptor or alternatively a modulator of the activities of other PR receptors or possibly both [162]. Hence, the function of PGRMC1 remains unclear, but this

transmembrane protein may function as an adapter protein involved in protein interactions, intracellular signal transduction and/or membrane trafficking [158]. PGRMC1 is reported to be overexpressed in a number of cancers including lung, breast, ovarian, colon and thyroid [161, 163]. In lung and BC, PGRMC1 overexpression correlates with tumor growth, migration, anchorage-independent growth, and serum-independent proliferation [164, 165]. Specific inhibitors of PGRMC1 in lung and BCs elicit lower ERK activity and reduced cell viability by impeding cell cycle progression past the G1 checkpoint [164]. Dimerized PGRMC1 complexed with CYP450 has also been implicated in cancer cell chemoresistance, potentially by facilitating antitumor effects of chemotherapeutics such as doxorubicin via CYP450 [159, 160] Cahill and colleagues suggest that PGRMC1 is a regulatory-nexus protein, associated with cancer pathogenesis and a potential target for novel antitumor therapeutics [162].

#### 2. Extranuclear sex steroid receptor: mechanism of action and signaling

2.1. Rapid ER extranuclear signaling mechanisms—Our ideas about sex steroid signaling have markedly expanded over the past 50 years. As ERa and ER $\beta$  are both ligandactivated transcription factors and not typical membrane receptors, trafficking of subsets of these receptors to cell membrane requires post-translational modifications. As highlighted above, two necessary modifications are palmitoylation and association with scaffolding proteins such as caveolins. It appears that it is through these complexes that ERs, which cannot by themselves activate G proteins or other signaling nodes, are able to initiate downstream intracellular signaling [166]. Palmitoylation results in interaction of ER with caveolin-1 that may serve as the transporter of ER to caveolae rafts in the membrane. ER and some isoforms are highly enriched in freshly isolated caveolae rafts [73, 167, 168] and probably initiate signal transduction from this site. On E2 stimulation, ERa dissociates from caveolin-1 allowing the activation of rapid signals critical to promote cell proliferation in BC cells [169]. Current data indicate that the process of palmitoylation occurs only on ER monomers. Hence, E2-induced ER dimerization limits the number of receptors that undergo this posttranslational modification and traffic to plasma membrane [112]. E2 binding to ERa results in receptor dimerization and activation of rapid signaling that involves, in part, interactions with discrete Ga and G $\beta\gamma$  proteins in target cells [74, 167]. Notably, ER have also been detected in early endosomes, which may play a role in trafficking of these receptors to the nucleus and/or in regulating receptor turnover [61]. Caveolin-1 also serves as a scaffold to bind/tether several critical signaling molecules to membrane caveolae raft sites, thereby enhancing interactions of ER with G-protein subunits and tyrosine kinases that promote acute downstream signals in seconds. Signaling to transactivation of growth factor receptors in cancer cells is reported to result in activation of MAPK and PI3K/AKT pathways that impact cell fate, proliferation, migration and other processes for malignant progression. Such 'extranuclear' regulation of cell functions may well act in concert with 'nuclear' effects of nuclear ER in BC cells [170].

**2.2 Rapid PR extranuclear signaling mechanisms**—In addition to its role as a transcription factor to direct nuclear action by binding to specific DNA sequences, accumulating evidence shows that PR can function outside the nucleus. The term 'extranuclear' activity of PR was suggested to specifically describe these PR functions

outside the nucleus [171]. Human PR contains a polyproline (PPD) domain within the Nterminal domain (NTD) that directly interacts with and activates cell membrane/cytoplasmic signaling cascades through interaction with SH3 domains of c-Src, thereby leading to rapid activation of Src/Ras/Raf/MAPK and other downstream targets [134]. This rapid extranuclear activation of c-Src and MAPK is progesterone-dependent but independent of PR transcriptional activity. Increasing evidence, mainly in BC cell models, suggests that extranuclear actions of PR are an integral part of PR function [172]. Progesterone-bound PR shows dichotomous functions in promoting and inhibiting cell proliferation in most tissues depending on the stage of development, and PR extranuclear signaling could play a major role in this dichotomy. Most studies to date demonstrate that progestin-dependent activation of extranuclear signaling stimulates several key kinases and results in modifications of PR, co-activators or other transcription factors required for maximum PR activity [134, 144, 173, 174] (Fig 3). However, a recent study also indicates that extranuclear PR actions can interfere or block growth-promoting signaling pathways. Expression of PR-B in NSCLC cells [77], independent of progestins, was shown to inhibit cell proliferation via PR-PPD mediated inhibition of the EGFR-mediated MAPK signaling pathway [77].

In addition to classical nuclear PR, recent data identify a novel truncated PR (PR-M) cloned from human adipose and aortic cDNA [175]. PR-M was predicted to contain a unique 16-amino acid sequence at the N-terminus lacking the A/B and the DNA-binding domain and was found to localize to mitochondria [176]. Progestin treatment induced a PR-M-mediated increase in mitochondrial membrane potential and oxygen consumption [176], but a detailed mechanism for PR-M mediated stimulation of cell respiration by progestins is unclear. More studies are clearly needed to determine the biological significance of PR-M in PR extranuclear signaling.

#### 2.3. Cross-communication between extranuclear ER and growth factor

**receptors**—Growth factor receptors such as EGFR and HER2 often concentrate in specific lipid rafts or caveolar domains of plasma membrane together with other signal transduction molecules. As noted above, extranuclear ERs also localize in these membrane domains, thereby promoting activation and transactivation of EGFR and HER2 receptors [73, 177] and interactions with other signaling molecules including insulin-like growth factor receptor (IGFR) I, the p85 regulatory subunit of PI3K, G-proteins, Src, and Shc, a protein that may couple ER with growth factor receptors [83, 178]. Coregulators, MNAR/PELP1 or metastasis-associated protein 1, also sequester ER in the extranuclear compartment to increase membrane action [169]. Activation of these pathways by E2 relays downstream proliferative and survival signals via MAPK and AKT that are important for BC survival, proliferation, migration and invasion. Further, MAPK stimulated by EGFR or/and HER2 signaling can, in turn, phosphorylate nuclear ER and receptor coactivators such as AIB1/SRC-3 [51, 73, 179, 180]. These events can be triggered by E2 (ligand-dependent signaling) or receptor kinases in the absence of E2 (ligand-independent ER activation), with the latter process underlying many forms of endocrine resistance in BC [181, 182].

**2.4 Cross-communication between extranuclear-nuclear PR and growth factor signaling**—About 70% of BCs are driven by ER activation [183]. In addition, several

studies suggest that progestogen treatment significantly increases BC risk and promotes E2induced cell proliferation [184, 185]. However, a recent report indicates that specific PR agonists inhibit E2-mediated cell proliferation, suggesting that PR functions as a molecular regulator of ER transcriptional activity via crosstalk between ER and PR in BC cells [141]. Of interest, two domains of PR-B were previously shown to be required for interaction with ER and for progestin activation of Src/MAPK signaling pathways [173]. Whether PR extranuclear signaling may play a role in modulating ER transcriptional activities remains to be explored further. Our current understanding of molecular mechanisms that underlie the convergence of extranuclear PR signaling with nuclear signaling is depicted in Figure 3.

Of 175 archival ER+/PR+ BC tissues investigated, about 55% exhibited ER/PR complexed with c-Src at the plasma membrane, and high levels of ER/PR/Src complexes were associated with poor BC prognosis [186]. BC cells expressing PR-B show enhanced responses to E2 and insulin-like growth factor (IGF-1), and this effect was independent of progestin and reported to be via PR-B/ER interaction with proline-, glutamate- and leucinerich protein 1 (PELP-1) [187]. However, inhibition of PR-B expression or treatment with a PR-specific antagonist, onapristone, significantly inhibited cell growth and helped to partially restore tamoxifen sensitivity in resistant BC cells [187]. These studies suggest potential clinical benefit of ER/PR extranuclear signaling and indicate that ER/PR extranuclear signaling could play a crucial role in modulating ER/PR transcriptional activities. There is good evidence that progesterone treatment via PR-PPD rapidly activates c-Src and its downstream Ras/Raf/MAPK (ERK) signaling pathway and downstream transcription factor targets such as Elk-1 [134, 173]. Further, PR-B bearing mutations in the PR-PPD that abolish PR-SH3 interactions (PR-B SH3) fail to mediate rapid progestindependent activation of c-Src [134], and progestin fails to activate Cyclin-D1 (CCD1) gene transcription in BC cells expressing PR-B SH3 [144]. PR-PPD mediated activation of Src/ MAPK signaling can result in PR phosphorylation at serine 345, allowing PR interaction with Sp1 through a tethering mechanism, to activate p21 and EGFR target genes [174] (Fig 3). MAPK phosphorylation of PR at serine 294 often associates with deSUMOylation on PR lysine-388 and hyperactivated PR [188]. Serine-294 phosphorylation and lysine-388 deSUMOylation of PR has been shown to regulate a unique set of genes found in BC with high ERBB2 expression and linked with decreased survival in tamoxifen-treated BC patients [189] (Fig 3).

Rapid extranuclear MAPK activation via interaction of PR and ER is essential for steps in the progestin activation of MMTV and other progesterone target genes. Activated phospho-MAPK translocates to the nucleus and forms a PR/MAPK and mitogen- and stress-activated protein kinase1 (Msk1) complex. This phospho-PR/MAPK/Msk1 complex is selectively recruited to the MMTV promoter by binding to BAF (BRG-1 associated factor) and p300/ CRE-binding protein-associated factor (PCAF) and recruiting BAP and PCAF to MMTV hormone response element (HRE); thereby displacing a repressive histone complex containing HP1 $\gamma$ , that leads to activation of the MMTV promoter [190]. Selective inhibition of MAPK or its downstream signaling interferes with rapid progesterone-mediated chromatin remodeling and blocks transactivation of the MMTV promoter [191]. These results demonstrate the convergence of PR extranuclear and nuclear signaling pathways on

progesterone-regulated gene transcription and indicate the significance of extranuclear signaling on overall gene regulation (Fig 3)

In addition to the Ras/Raf/MAPK signaling pathway, previous studies indicate that other mitogenic protein kinases, including cyclin-dependent kinase (Cdk2) and Casein Kinase II (CK2), phosphorylate PR and change PR transcriptional activities [192], suggesting that PR may serve as a mitogenic sensor in cells [193]. Cyclin A/Cdk2 is an important S-phase cell-cycle dependent kinase that phosphorylates large numbers of PR phosphorylation sites (8–14 sites) [194], such as Ser-162, Ser-190, Ser-400, and functions as a PR coactivator [195]. CyclinA/Cdk2 phosphorylation of PR regulates important PR activities including nuclear translocation, transactivation of PR-regulated genes, and hormone-induced degradation [194]. This intricate PR regulation by Cdk2 suggests significant roles of PR in the regulation of the cell cycle and proliferation.

CK2 is another important kinase that regulates PR functions. PR phosphorylation by CK2 on Ser-81 requires PR interaction with DUSP6, a negative regulator of the MAPK signaling pathway. This indicates that DUSP6 may serve as a scaffold to permit interaction between PR and the kinase that phosphorylates PR at Ser-81 [196]. Interestingly, PR phosphorylation at Ser-81 associates with a gene profile for mammary stem cell maintenance/renewal that is often altered in BC [196, 197]. PR with Ser-81 phosphorylation further regulates a set of genes associated with interferon/inflammation and STAT-signaling [197], thereby suggesting a potential link between PR regulation, inflammation and cancer [198]. Together, these data provide evidence that progesterone regulates gene expression by stepwise modifications ranging from the regulation of receptor activity and receptor degradation that require activation of several kinases and signaling pathways initiated by extranuclear functions of PR and ultimately chromatin organization, thereby indicating a natural crosstalk between nuclear and extranuclear PR (Fig 3).

#### 3. Clinical implications of extranuclear steroid signaling in breast cancer

3.1. ER extranuclear signaling in breast cancer: ER assays to predict clinical outcome—At diagnosis, about 70% of BCs contain ERs and depend on E2 for growth. Since ER expression in BC is predictive of a clinical response to hormonal therapy, this has led to use of antiestrogens (tamoxifen, fulvestrant) and aromatase inhibitors in treating ERapositive BC [16, 17, 20, 199, 200]. However, a substantial proportion of patients with localized BC, and essentially all patients with metastatic disease, become resistant to endocrine therapies. For more than 40 years, the IHC assay for nuclear ERa expression has been the predominant biomarker in BC, largely because of its important role in predicting potential benefit from endocrine therapy. Such observations led to current use of antiestrogens and AIs in treating ER-positive (ER+) BC. In addition, nuclear ER expression has important prognostic implications, with ER-negative tumors exhibiting a more aggressive phenotype. Earlier assays for ER depended on extensive homogenization of fresh-frozen tumor and preparation of cytosol extracts by centrifugation to perform a ligandbinding assay (LBA) [15, 200]. Of note, artifacts resulting from tumor homogenization in the LBA led to misinterpretations on the subcellular distribution of ER in cells. Unbound ER was found almost exclusively in cytosol, while ligand-bound ER was recovered in nuclear

homogenate fractions. Later, it was recognized that these results were due to dissociation of unbound ER from the nucleus during homogenization of intact cells. This eventually led to the notion that ER was exclusively a nuclear-localized protein, a concept reinforced by localization of ER predominantly in tumor cell nuclei by use of monoclonal antibodies targeted to nuclear ERa [1, 4]. These antibodies were used to develop current assays for nuclear ERa. Further laboratory studies have revealed that the same experimental artifact that elicited dissociation of unbound ER from the nucleus during homogenization was also responsible for extracting another small pool of ERs from target cell membranes [36, 38, 39]. Investigations reviewed above confirm the presence of extranuclear membraneassociated ERa in BCs and further show that these ER help to stimulate BC gene transcripts and promote cancer progression [5, 41, 52, 83, 201]. Of note, activation of ER by a membrane-impermeant estrogen-dendrimer conjugate was shown to stimulate extranuclear responses in ER-expressing human BC cells, also contributing to regulation of BC cell number [53]. Of note, E2-induced growth of MCF-7 cells in vitro was blocked by treatment with the antibody to ERa and correlated closely with acute hormonal activation of MAPK and AKT kinase signaling [57]. In addition, E2-promoted growth of human BC xenografts in nude mice was significantly reduced by treatment *in vivo* with the ERa antibody. Thus, regulation of BC progression by E2 may be mediated by coordinated actions of ERs in nuclear and extranuclear compartments. These findings suggest that it may be important to measure both nuclear ERa and extranuclear membrane-associated ERa for greater diagnostic accuracy and to predict the response to BC treatment going forward.

Based on current IHC assays for nuclear ERa, only about half of advanced BCs with expression of ERa and/or PR respond to endocrine therapy, suggesting a need for improved assays designed to correlate with patient outcome [16]. The relatively poor specificity of nuclear ERa status in identifying tumors that will respond to hormone therapy has been documented in several studies and suggests that other factors are important [15, 16, 202, 203]. IHC results depend greatly on the type of antibody used, detection procedures, subcellular densities of ERs and problems due to long-term formalin fixation and antigen masking. Ongoing efforts to standardize IHC methods for detecting nuclear ERa, especially antibody validation and antigen retrieval methods, are helping to improve the reliability of this assay in archival BCs [204–208] [209]. These reports underscore the importance of fully elucidating the nature of the ER+ phenotype. To date, use of clinical IHC assays developed to detect nuclear ERa have shown relatively low levels of extranuclear ERa in standard formalin-fixed, paraffin-embedded BC specimens from the clinic [41, 210]. Although extranuclear ERa is detected in a number of cases, Welch et al. [210] suggest that the low incidence ranging from 0 to 3-4% using current assay methods is unlikely to be of routine clinical value. However, the criteria for positive extranuclear ER staining included immunoreactivity at least 25% or greater than that of nuclear ER staining. Using differing preanalytic fixation and processing methods and a different scoring system for intensity, Kim et al. [41] reported that extranuclear expression of either ER or PR was observed in 9.5% of clinical cases. Both studies used validated nuclear receptor antibodies. Given the abundance of data on extra-nuclear ER/PR expression in BC cell lines and preclinical models as detailed above, it will be important going forward to assess variables such as tissue fixation methods, antigen retrieval and processing protocols and tissue age that may

impact extranuclear/membrane ER/PR versus nuclear receptor detection. Approaches to detect and quantify ER/PR forms present at very low levels or at a low density in subcellular compartments in human specimens are not yet available.

Recent work suggests that ER+ progenitor cells in the breast arise from primitive stem cells in fetal development [211]. ER+ progenitors proliferate in response to E2 and also produce paracrine factors that influence proliferation of adjacent ER-negative cells. Further, BC progression may result from transformation of these stem/progenitor cells. Findings indicate that only a minute subset of tumor cells (about 1%–10% of the primary BC mass) have tumorigenic potential [212, 213]. These tumor-initiating cells, termed cancer stem/ progenitor cells (CSPC), are crucial in BC progression and are resistant to most chemotherapies. As such CSPC cells drive tumor formation, this model has important implications to understand BC progression, and to develop strategies for BC detection, prevention and treatment. Methods to detect in clinical specimens such minute subsets of CSPCs and potential extranuclear ER expression is another major challenge [214, 215].

The prognostic/predictive value of ER $\beta$  in BC is uncertain at this time, and conflicting reports are complicated by the identification of several variant forms of ER $\beta$  that are not detected by the current clinical IHC assays for ER $\alpha$  [15, 71, 216]. There is evidence that endogenous ER $\beta$  localizes to plasma membranes in clinical tissue specimens including BCs [71, 217]. Extranuclear ER $\beta$ 2 expression, alone or combined with nuclear staining, predicted significantly worse overall survival among archival BC specimens [71]. Patients with only extranuclear ER $\beta$ 2 had significantly worse outcome. In this series, nuclear and extranuclear ER $\beta$  expression differentially affected outcome, suggesting that measurement of these variables in BC could provide a more comprehensive picture of patient outcome to complement ER $\alpha$  assays. However, such data are limited by availability of specific, validated antibodies. Further, expression of ER $\beta$ 1 is of particular interest because it is the only ER $\beta$  isoform that has an intact LBD, thereby serving as a potential drug target in clinic [218]. Of note, extranuclear signaling by ER $\beta$  has also been reported to be a significant factor in progression of medulloblastoma [219].

Besides full-length ERa-66 kD, the ERa-46 kD isoform is also expressed in a majority of archival BCs tested [220–222]. Binding affinities of both unliganded and fully-activated receptor forms toward coregulator peptides revealed that respective potencies of ERa-46 kD and ERa-66kD differed significantly, contributing to differential transcriptional activity of target genes to E2 stimulation. Expression of ERa-46 kD in BCs does not consistently correlate with expression of full-length ERa, but overall ERa-46 kD was expressed in ~70% of BC samples analyzed [220]. In BC cells *in vitro* with expression of both ERa-46 kD and ERa-66 kD, ERa-46 kD can impact cell proliferation depending on ratios of ERa-66 kD/ ERa-46 kD [223, 224]. However, mice lacking the N-terminal A/B domain of ERa-66 kD were infertile due to lack of uterine epithelial responses to E2 normally mediated by full-length ERa [225]. Collectively, these findings indicate the importance of choice of antibodies used for BC diagnosis which may or may not detect ERa-46 kD or other ER forms, a factor that may have clinical relevance.

**3.2 Metabolic effects of extranuclear ER signaling**—Recent studies suggest that E2 regulates glucose metabolism in normal and malignant cells at least in part from membrane ER signaling [226]. A recent review details extensive effects of E2 in modulating normal organ metabolism [227]. Both ERa and ER $\beta$  play roles in normalizing insulin sensitivity and glucose homeostasis as revealed in ER-knockout models. Membrane ERa and ER $\beta$  are reported to have important roles in promoting insulin sensitivity and synthesis and secretion from pancreatic  $\beta$ -cells [228, 229]. Notably, high glucose and increased insulin signaling promote aggressive development of BC in mouse models [230], suggesting an important pathway by which ER stimulates BC development. Emerging reports further indicate that E2 signaling via AMPK modulates glycolytic pathways in BCs under reduced glucose conditions to promote the survival of BC cells *in vivo* [226].

Both ER $\alpha$  and ER $\beta$  are identified in the mitochondria of BC cells, primarily in the mitochondrial matrix [231, 232]. There is evidence in BCs that mitochondrial gene regulation occurs in part by means of E2 binding to ER in this intracellular organelle [233]. Additionally, mitochondrial ER $\beta$  is reported to mediate cytoprotection and survival of BC cells on E2 binding in mitochondria [62, 234].

3.3 ER extranuclear signaling and endocrine therapy in the clinic—Hormonal therapy was first used more than 120 years ago, marking the start of the current era of targeted antitumor treatment [235]. This approach is based on blocking activity of estrogens and their receptors. In the clinic, endocrine therapy is an important intervention for BCs that express ER and/or PR, and it has proven to be one of the most effective BC treatment strategies. Despite recent improvements in therapeutic options, development of endocrine resistance is one reason that BC is the 2nd most frequent cause of cancer death in women [199, 236, 237]. In most cases, ER is present in resistant tumors, and in many of these its activity continues to regulate BC growth [3]. Tamoxifen has been the most widely used hormone therapy for over 25 years, achieving a 39% reduction in BC recurrence and 31% reduction in mortality in nuclear ER $\alpha$ + early BC [16, 20]. Although effective, tamoxifen has important drawbacks: a limited period of activity before resistance develops and undesirable side-effects in normal tissues such as uterus due to its activity as a partial agonist [18]. Of note, cooperative interactions between growth factor receptors and ER signaling pathways are identified in BC and NSCLC, and growth factor-mediated pathways, notably those of EGFR, HER2, and insulin-like growth factor receptor I, are critical in development of some types of antiestrogen resistance in BCs [17, 177, 238]. About 15-20% of BCs have overexpression of HER2 receptors, and increased HER2 expression correlates with poor clinical outcome and resistance to endocrine therapy [177, 238-242]. Such tamoxifen resistance is reported to associate with formation of HER2/ERa membrane-associated complexes that lead to non-genomic activation of both AKT and RPS6KA2, which in turn provides these BC cells with a survival advantage [243]. Similarly, overexpression of EGFR in about 50% of BCs correlates with endocrine resistance [244-247]. These data offered a rationale to target both ER and HER2 in ERa+/HER2-positive BCs in the clinic to overcome this type of endocrine resistance (Johnston et al., 2009).

As long as ER is present in tumors, growth may be stimulated by E2, partial agonists or ligand-independent action [17, 177, 238]. Introduction of aromatase inhibitors (AIs) for

postmenopausal patients, either initially or after tamoxifen, has yielded better outcomes than the prior standard of 5 years tamoxifen [20]. However, in patients with advanced BC, only about 1/3 of nuclear ERa-positive BCs respond to AIs, and resistance can evolve due to ER activation by ER hypersensitivity, ligand-independent ER activation by activated growth factor receptor signaling [17, 177, 238] or emergence of receptor mutations. Unfortunately, in about 14% of metastatic ER+ BCs from patients with multiple prior endocrine therapies, there is evidence for acquisition of functionally-aberrant ESR1 with point mutations that often occur in the LBD of ERa [23, 248, 249]. Such mutated ESR1 variants continue to respond at least partially to ER antagonists such as fulvestrant which has a unique capability to downregulate and eliminate ER by induction of the ubiquitin-proteosome pathway[18, 250, 251] but higher drug doses are required to achieve wild-type ER levels of inhibition. These data underscore the need to find more potent ER antagonists and to implement use of hormonal treatments in combination with other synergistic agents such as cyclin-dependent kinase (CDK 4/6) inhibitors [252, 253]. There is currently no data on the status of extranuclear ER forms in metastatic tumors from the clinic with ESR1 mutations. However, there is evidence that hormonal antagonists such as fulvestrant, that block nuclear ER actions, also effectively block rapid E2-induced signaling in BC cells with wild-type ER [57, 73, 83], but therapies to specifically target membrane-associated ERs are yet to be developed.

While many studies on extranuclear steroid signaling have focused on estrogens, similar data confirm that extranuclear signaling by other sex steroids may be equally important in the clinic [50]. An experimental model whereby a biologically relevant and rogen-mediated process is regulated completely independent of transcription was well-characterized in Xenopus laevis oocyte maturation [254]. Androgen-induced Xenopus oocyte maturation is mediated by classical androgen receptors (AR), as both the AR antagonist flutamide and AR knockdown by siRNA inhibit androgen-triggered maturation. Based on IHC and biochemical work, classical AR is expressed throughout the cell, with approximately 5% found in plasma membrane. These membrane-localized ARs regulate androgen-mediated maturation, as testosterone coupled to bovine serum albumin triggers oocyte maturation as well as free steroid. Of interest, prostate cancer is a well-characterized androgen-dependent tissue and is initially responsive to androgen deprivation (castration). However, over time, prostate cancer cells ultimately become castration-resistant due in part to AR amplification/ mutation that allows responses to lower levels of androgens. In castration-resistant prostate cancer, the synthetic AR antagonist enzalutamide exhibits potent affinity to bind AR and blocks translocation of extranuclear AR to the nucleus to suppress binding of the ligandbound receptor complex to DNA [255]. In the clinic, FDA-approved enzalutamide has had a major impact in improving the outcome of patients with castrate-resistant prostate cancer. Unexpectedly, enzalutamide is being assessed in clinical trials as a treatment for triplenegative breast cancers (TNBC) that lack expression of ERa, PR and HER2 [256].

TNBC occurs in 10–15% of patients, yet this disease subtype accounts for about half of all BC deaths. As TNBCs lack clinical expression of ER $\alpha$ , PR and HER2, they are not considered to be a target for hormonal therapy [257]. Although initially responsive to chemotherapy, TNBCs tend to relapse and metastasize early and have worse prognosis than other subtypes. Of note, several reports indicate that ER $\beta$  is expressed in TNBC cells [258,

259] and may play a role in TNBC progression [260–263]. ER $\beta$  forms occur in tumor cell nuclei, but also at extranuclear sites [71, 264]. ER $\beta$  can activate transcription by nuclear or indirectly by extranuclear pathways by interaction with coactivators/co-regulators such as PELP1 and SRAP [265] that in turn modulate signaling cascades to impact gene expression and TNBC progression [70, 71, 264, 266, 267]. As a consequence of such preclinical work, clinical studies to target ER $\beta$  are being planned in patients afflicted with metastatic TNBC [268].

#### 3.4. ER extranuclear signaling in the tumor microenvironment and

**immunotherapy**—Several steroid hormones regulate immune responses that contribute to significant changes in immune function during inflammatory and autoimmune diseases [269]. The regulation of immune responses by steroid hormones occurs at multiple levels including cell development, proliferation, cytokine or antibody production and apoptosis. Regulation of proliferation and apoptosis is especially critical in development of appropriate T- and B-cell repertoires. Of note, estrogen-binding proteins are reported to occur on plasma membranes of immune cells, and ERs are also reported to be expressed in myeloid-derived suppressor cells that act to block immune recognition of malignant cells [269, 270]. E2 binding to ERs and downstream activation of such immune suppressor cells are inhibited by fulvestrant, leading to promotion of antitumor activity in immune-competent mice. This work suggests that new approaches to management of BC such as repurposing of antiestrogens for combination therapy with immune checkpoint inhibitors may be forthcoming [271, 272]. Further studies are needed to detail membrane steroid receptor expression and signal transduction pathways in immune cells and in potential targets of the host immune system, particularly TNBCs [273].

#### 3.5. ER and PR extranuclear signaling and resistance to endocrine therapy-

Expression of PR and/or PR isoforms is a valuable marker for tumor aggressiveness and disease progression in breast, endometrium and lung cancers [274–276]. However, there are multiple hypotheses about how aberrations, loss and/or gain in PR expression is involved in cancer initiation, progression, metastasis and resistance to endocrine therapies. In normal mammary gland, PR+ cells are often found in non-proliferating cells [277]. However, the percentage of proliferating PR+/ER+ cells increases in hyperplasia and ductal carcinoma in situ (DCIS) states, and ER+/PR+ BC cells exhibit increased cell proliferation [278, 279]. Yet, loss of PR expression in BC is associated with endocrine resistance, BC aggressiveness and poor BC patient prognosis [280]. BC patients with PR-A rich tumors have worse disease-free survival as compared to those with PR-B rich tumors [281]. Selective loss of PR-B is also associated with poorly differentiated and progestin-resistant endometrium cancer [282]. Further, low PR expression predicts poor clinical outcome in NSCLC patients [276]; and treatment with PR antagonist (RU486) improves the survival of mice with spontaneous lung cancer [276]. Therefore, PR seems to mediate both proliferative and antiproliferative functions depending on cellular and tissue contexts. How the presence of PR or PR isoforms affects cell growth and differentiation in these tissues and cancers remains poorly understood.

Accumulating evidence suggests that aberrant ERa/PR signaling in both the nucleus and the cytoplasm plays a critical role in altering endocrine pathways that lead to endocrine resistance in BC [283]. The presence or absence of ER and its ligand significantly affects the outcome of progesterone action in the nucleus [284], and ERa/PR extra-nuclear signaling pathways in the cytoplasm and membrane, are strongly implicated in hormone resistance [134, 144, 285–287]

There is considerable cross-communication between ERa and PR pathways, and both hormone pathways have been shown to co-regulate/cooperate in extranuclear signaling. E2 is reported to preferentially promote activation of the PR-B isoform [288]. In cells expressing ERa and the PR-B isoform, treatment with progestin, a synthetic progestogen, can attenuate antiestrogen-mediated growth arrest [144]. When ERa and PR are uncoupled, as in ERa-negative cells, rapid PR signaling can be exerted through PR-B interaction via the proline domain interacting with SH3 domains of the c-Src family of tyrosine kinases [76, 134, 144, 171, 289]. In addition, rapid extranuclear progestin activation has been postulated to play a role in gene regulation by mediating PR phosphorylation, thus acting as a feed-forward mechanism for PR nuclear transcriptional activity [290].

Experimental studies indicate that rapid, extranuclear PR signaling is mediated by the PR-B isoform which exerts rapid hormone signaling in the cytoplasm that interacts with several cellular signaling pathways. These pathways include those regulated by epithelial growth factor (EGF) [291], PI3K/AKT [292], IGF [293], Src receptor tyrosine kinases and RAS/ p42/44 MAPK [76, 132–134, 144, 171, 289, 294]. Together, these pathways facilitate BC cell survival, cytoskeletal remodeling, proliferation and invasion. These acute signaling events are too rapid for classical transcriptional effects. As noted, extranuclear signaling of E2/ERa/Src occurs within minutes and is involved in transmitting signals that increase cell division or decrease apoptosis mediated by MAPK, and PI3K/AKT signaling pathways [295]. A functional Src interaction appears critical for extranuclear ERa function, and excessive extranuclear ERa/Src signaling associates with endocrine resistance [296]. Treatment with Src inhibitors is effective in blocking endocrine-resistant tumor progression [296], thus defining a potential role for pre-nuclear hormone receptor activity as a mechanism for endocrine-resistant BC phenotypes.

Additional reports suggest that a switch from ER/PR-mediated paracrine signaling for modulation of growth factor signaling pathways (e.g. EGF, HER2, Wnt, Src) to autocrine signaling contribute to BC progression [172] and potentially to the generation of endocrine-resistant tumors. How this information may be applied in endocrine therapy in the clinic remains unclear. However, treatments targeted to block rapid, hormone-dependent autocrine proliferation mechanisms may potentially have benefit in endocrine-resistant tumors.

**3.6. Extranuclear PR signaling in NF-\kappaB/p53 signaling**—Epidemiological studies indicate that full-term pregnancy early in reproductive life provides a long-lasting and strong protection against BC [297], thus offering a clue that early, limited sex steroid stimulation may play a role in reducing later BC risk. This protective effect is reproduced by a 3-week exposure to low pregnancy-level doses of both estrogen and progesterone [298], but neither hormone alone is sufficient to induce the protective effect. Further, the absence of p53 tumor

suppressor gene function was found to abrogate the protective effect of hormones against carcinogen-induced mammary carcinogenesis in mouse models. Limited data suggest that progesterone/PR and p53 may play a cooperative role in regulating chromosome stability [299]. Extranuclear PR signaling regulates p53 expression in human umbilical vein endothelial cells (HUVEC) [300], as progesterone treatment of HUVECs activates c-Src/Ras/Raf/MAPK/NFκB signaling to activate p53 gene transcription and control of cell cycle progression [300, 301]. Progesterone/PR also regulate p53 expression in BC, but it remains to be determined if similar mechanisms of extranuclear/nuclear PR activation of p53 occur in BCs. Thus, hormone stimulation at a critical period in mammary development appears to result in hormone-responsive cells with long-term changes in cellular regulatory loops governing proliferation and potentially responses to DNA damage.

#### 3.7. Extranuclear ER/PR signaling - actin/cytoskeletal rearrangement, cell

motility and metastasis-Cancer cell motility is essential in the process of BC invasion and metastasis. Cell migration is controlled by remodeling of the actin cytoskeleton by the formation of focal adhesion complexes and loss of stress fibers. This cytoskeletal reorganization associates with formation of specialized cell membrane structures such as membrane ruffles, filopodia and lamellipodia [302, 303]. Extranuclear ER/PR signaling is reportedly involved in actin remodeling and cell migration [301, 304]. E2 treatment of T47D ER+/PR+ BC cells induces rapid changes in the actin cytoskeleton with formation of membrane ruffles and pseudopodia [305]. These effects are achieved by acute interactions of ERa with G protein Ga13 resulting in recruitment of small GTPase-RhoA and activation of its downstream target Rho-associated kinase-2 (ROCK-2) and phosphorylation of moesin, an actin regulatory protein [306]. Similar signaling cascades mediated by ligand-bound ERa are also reported in endometrial cells [307]. These studies reveal that E2-activated ERa stimulates BC cell migration via activation of c-Src, paxillin and FAK, signaling molecules that play important roles in regulating focal adhesion dynamics to facilitate cell migration. Recent studies in MCF-7 BC cells similarly show that E2-mediated extranuclear signaling promotes cytoskeleton remodeling and cell migration via PELP1 scaffolding protein, c-Src, and integrin-linked kinase signaling pathways, with expression of PELP1 positively correlated with the invasiveness of BCs [308, 309].

Extranuclear PR signaling also plays a significant role in regulating cytoskeletal remodeling. Hence, PR-null MDA-MB-231 BC cells transfected with PR and exposed to progesterone exhibit increased expression of focal adhesion proteins, paxillin and FAK, increased numbers and diameters of stress fibers and increased cell migration and invasion [310]. Treatment with progesterone or synthetic progestin medroxyprogesterone acetate (MPA) also stimulates cell migration and invasion of HUVEC cells and T47D cells, a process preceded by rapid remodeling of the actin cytoskeleton and formation of membrane ruffles and pseudopodia [311]. These changes are mediated in part by phosphorylation of moesin through interaction between PR-A and G protein  $G_{\alpha,13}$ . MPA treatment also promotes PR interaction with c-Src resulting in activation of a PI3K/RhoA/ROCK-2 signaling cascade [311, 312]. Progesterone treatment further promotes cancer cell migration in 3D-matrices by activating formation of focal adhesion complexes as indicated by Tyr-397 phosphorylation of FAK via the c-Src/PI3k-Akt/RhoA/ROCK2 cascade [313]. Extranuclear PR signaling

regulates BC cell migration by progesterone stimulation of the c-Src/AKT pathway, leading to activation of ribosomal S6 kinase 1 (RSK1) and increased phosphorylation of p27 at T198 with formation of a p27pT198/RhoA complex in cytosol. The p27pT198 RhoA complex prevents RhoA degradation and enhances cell migration; inhibition of RSK1 with RSK1 inhibitor BI-D1870 reversed these events. Together, these data suggest RSK1 activation plays a role in progesterone-stimulated BC cell migration [314], yet clinical-translation of these findings remains to be accomplished.

#### 3.8. Extranuclear signaling by environmental estrogen and breast cancer risk

-Increasing evidence suggests that exposure to environmental estrogens could be one of the major factors contributing to increased risks of BC [315]. Environmental estrogens such as bisphenol A (BPA), DES, organochloride pesticides and genistein have been shown to impact both nuclear and extranuclear ER signaling [316]. Rapid signaling from membrane ER in response to both E2 and xenoestrogen DES regulates the histone methyltransferase enhancer of Zeste homolog 2 (EZH2). Extranuclear ER signaling via PI3K/AKT kinase phosphorylates EZH2 at S21, reducing levels of trimethylation of lysine-27 on histone H3 in hormone-responsive cells, a process that can reprogram the profile of E2-responsive genes in target organs such as breast and uterus [317]. In addition, phytoestrogen genistein, but not BPA, induce extranuclear ER activation of PI3K/AKT kinase and phosphorylation of EZH2 leading to hormonal hypersensitivity and higher risk of uterine tumorigenesis in rats [318]. To date, these data indicate that rapid extranuclear ER signaling may provide a direct link between xenoestrogen-induced receptor signaling and modulation of epigenetic machinery during tissue development leading to increased hormone-related cancer risk. Development of improved screening methods to identify potentially harmful environmental agents and new interventions to manage such exposures are urgently needed [319]. One unfortunate clinical example is the long-term impact of fetal exposure to the synthetic estrogen diethylstilbestrol (DES). Maternal ingestion of DES was associated with later development of reproductive tract abnormalities including adenocarcinoma of the vagina in young female offspring [320, 321].

## Concluding remarks

Numerous reports suggest that specialized proteins associated with target cell membranes are the first contact point for sex steroid hormones. For E2, progesterone and androgen, such extranuclear receptor molecules are often reported to share homology with nuclear forms. These primary interactions trigger a cascade of specific downstream cellular responses. Thus, rapid sex steroid hormone-dependent signaling is postulated as a prerequisite in part for later receptor-induced transcriptional activity, suggesting an integrated model of steroid receptor signaling where steroid hormone-dependent early extranuclear effects contribute to late receptor-dependent nuclear actions. Ongoing research continues to address whether extranuclear-initiated and nuclear transcriptional activities of sex steroid hormones are integrated to yield one overall effect or represent parallel pathways with distinct effects. More detailed understanding of molecular mechanisms, such as palmitoylation of ER and PR that target these receptors for membrane localization, remains to be elaborated in both normal and transformed cells.

While it is now well accepted that ER and PR can perform both nuclear and extranuclear functions and localize to cell membrane and nuclear pools, studies also suggest that alternative E2/progesterone-responsive membrane proteins such as GPR30, mPRs, or PGRMC1 may mediate extranuclear signaling. Physiological significance of these alternative molecules and their potential cross-communication with nuclear receptors clearly require further exploration.

Based on available evidence, nuclear and extranuclear ER/PR signaling appears to cooperate in mediating BC progression. Validation of extranuclear steroid hormone receptor structures and their specific biologic functions in health and disease may well lead to development of a new generation of targeted therapeutics to manage sex steroid hormone-responsive cancers in the clinic.

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# Abbreviations

aromatase inhibitor
breast cancer
ductal carcinoma in situ
estradiol-17β
estrogen receptor-alpha
estrogen receptor-beta
estrogen receptor gene
ligand-binding domain
non-small cell lung cancer
progesterone receptor-A
progesterone receptor-B
triple-negative breast cancer

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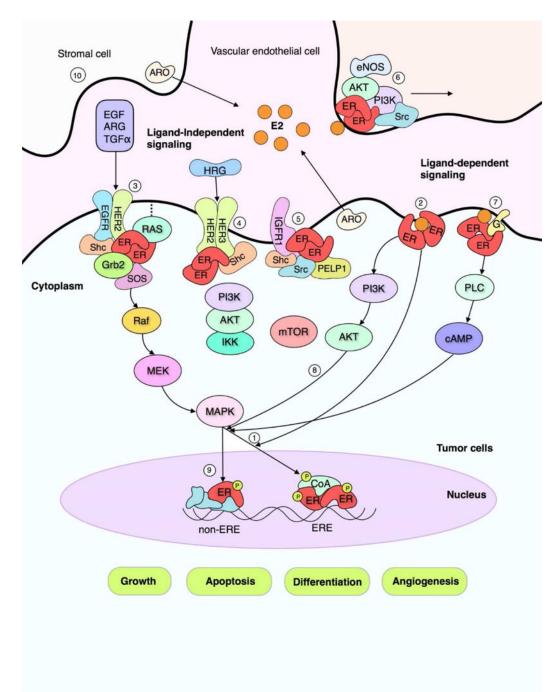
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• ER and PR play key roles in breast cancer development and progression.

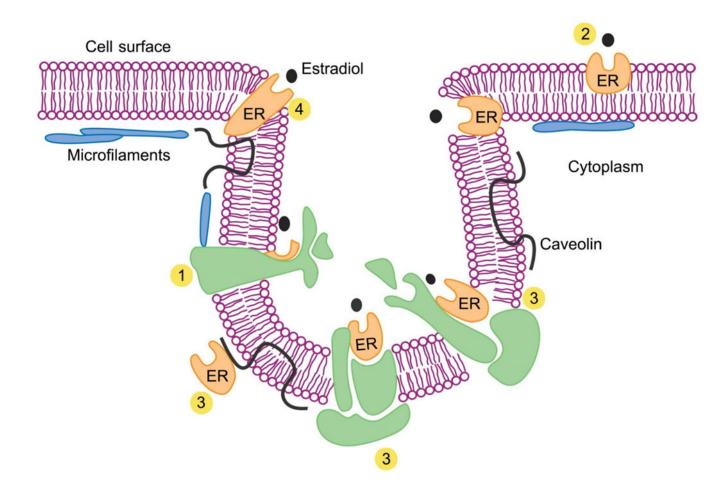
- Hormone-induced rapid extranuclear signaling affects signaling networks.
- Extranuclear signaling modifies ER/PR transcription and gene networks.
- History and updates on ER/PR extranuclear signaling are reviewed and discussed.



## Figure 1. Estrogen receptor signaling pathways in human breast tumors

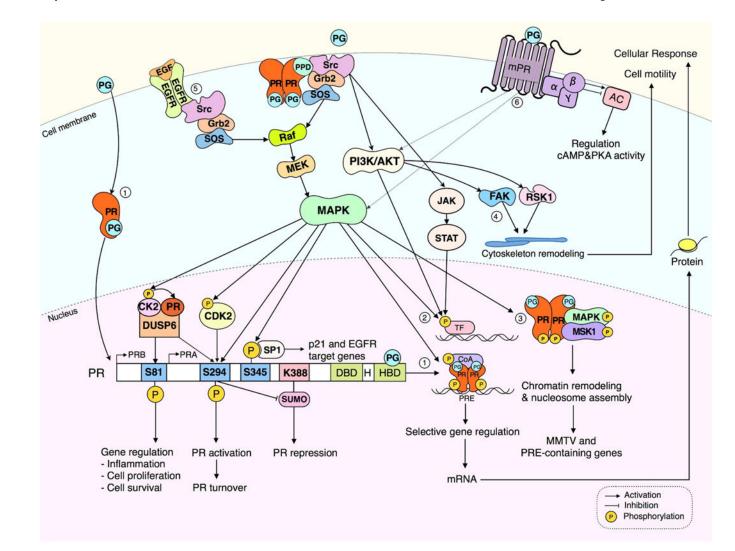
Proliferation and survival of BC cells is closely regulated by estradiol-17 $\beta$  (E2) and its receptors, ERa and ER $\beta$ , as well as growth factor receptors. In classic models of E2 action (1), E2 binds ER to promote dimerization and phosphorylation of ER. This allows direct binding of the E2-ER complex with steroid receptor coactivators and estrogen response elements (ERE) in DNA, leading to gene transcription to regulate growth, differentiation, apoptosis and angiogenesis. A subset of ERs occur in extranuclear sites, such as caveolae or lipid rafts in plasma membrane (2), and may interact with transmembrane growth factor

receptors such as EGFR (3), HER2 (4), insulin-like growth factor receptor I (IGFR1) (5) and other signaling molecules, including components of Ras-MAPK and phosphatidylinositol 3-kinase (PI3K)/AKT pathways, Src kinases, Janus-activated kinase/signal transducer and activator of transcription signaling, nitric oxide synthase (NOS) (6), and G-proteins (7). Membrane-associated ER may undergo posttranslational modification, such as palmitoylation, and/or associate with adaptor proteins, such as Shc, PELP1, or lipid raft proteins. ERs and growth factor receptors may form a structured complex for signal transduction to MAPK and/or PI3K/AKT kinase that interacts, in turn, with nuclear ER and steroid receptor coactivators (8). Signaling for cell growth involves phosphorylation (P) of nuclear ER and coactivators that can occur in ligand-dependent as well as ligand-independent modes. ERE-dependent and alternate transcription sites may be activated (9). Further, E2 is produced locally in BC and in host supporting cells by the action of aromatase (ARO) (10) which is regulated by both nuclear and extranuclear ER. Pathways modified from Pietras and Marquez [52]; refer to text for more details.



## Figure 2. Supramolecular organization of plasma membrane and occurrence of estrogen receptors (ER)

A model of the surface membrane from an estradiol (E2)-responsive cell in the region of a caveolar structure is shown. E2 may interact with one of several different forms of membrane-associated ERs. Full structural characterization of these ERs remain to be done. These molecules may be known membrane components, such as enzymes, G-proteins, ion channels, or receptors for nonsteroidal ligands, with unrecognized steroid binding sites (1); new isoforms of hormone receptors such as the truncated ERs that arise by alternative splicing (2); 'classical' receptors complexed with other membrane-associated proteins (3); or novel membrane proteins (4). Of note, splice variants of ER occur, and these give rise to proteins of different molecular size and possibly modified properties. Membrane insertion of receptors in primary transcript form would likely require one or more hydrophobic regions. ERa, for example, has several hydrophobic regions, but it is not known if these suffice for disposition as integral membrane proteins. Posttranslational modification of receptor protein leading to membrane targeting also occurs, including phosphorylation, glycosylation, nitrosylation and/or addition of lipid anchors or alterations such as palmitoylation or myristoylation. Evidence for palmitoylation of ER that fosters membrane association is documented in the text. Modified from Szego and Pietras [322].



## Figure 3. Convergence of PR nuclear and extranuclear signaling pathways to regulate biological responses

In the classic nuclear signaling pathway (1) progesterone activates progesterone receptor (PR) by binding and inducing conformational changes of the receptor causing in turn nuclear translocation, dimerization and binding to progesterone response elements (PRE) in the promoters or enhancer regions of PR target genes. Progestin treatment rapidly activates extranuclear signaling of a subpopulation of PR localized in the membrane or cytoplasm to transiently associate with c-Src through interaction with the PR polyproline domain (PPD) leading to activation of mitogen-activated protein kinase (MAPK), phosphatidylinositol 3-kinase (PI3K/Akt), or Janus kinase/signal transducers and activators of transcription (JAK/ STAT) signaling pathways. Activated MAPK further regulates PR transcriptional activity through phosphorylation of PR, co-activators (1) or other transcription factors (2). Rapid progestin activation of MAPK can directly phosphorylate PR or co-activators (1) or promote PR phosphorylations through CK2 and CDK2 (see text for details). Alternatively, activation of various cytoplasmic signaling pathways may phosphorylate and increase transcriptional activity of other transcription factors, independent of PREs (2). Rapid extranuclear activation of ER/PR complexes by progestins activates MAPK leading to a formation of phospho-PR/

MAPK/Msk1 (Mitogen and stress-activated protein kinase 1), chromatin remodeling and enhanced MMTV and PRE-containing gene transcription (3). PR extranuclear activation of Src/PI3K/Akt may stimulate focal adhesion kinase (FAK) or ribosomal S6 kinase 1 (RSK1) triggering actin cytoskeleton remodeling and promotion of cell motility (4). Alternatively, PR extranuclear signaling may cross-communicate with growth factor signaling such as epidermal growth factor (EGF) leading to activation of MAPK and downstream events (5). Membrane localized progestin receptor (mPR) unrelated to the classical PR mediates progestin extranuclear signaling through GPCR-like membrane proteins via modulation of adenylate cyclase (AC), cyclic adenosine monophosphate (cAMP) and protein kinase A (PKA). mPR has been reported to activate MAPK and PI3K/Akt signaling pathways. More work is needed to define the physiological significance of extranuclear mPR activation of these several cytoplasmic signaling pathways in BC cells.