

## Membrane-Associated Estrogen Receptor Signaling Pathways in Human Cancers

Richard J. Pietras and Diana C. Márquez-Garbán

### Background

Hormonal therapy is a mainstay in the clinical management of patients with breast tumors that express estrogen receptor (ER). The antitumor efficacy of these endocrine treatments depends on tight regulation of breast tumor cell growth by estrogens and growth factors. However, as these cancers progress, they usually become resistant to antiestrogens, and most patients no longer respond to endocrine therapy. New options for antiestrogen treatment are urgently needed to enhance patient survival and quality of life. Improved therapies and treatment strategies may derive from the discovery that alternative pathways are involved in the modulation of estrogen signaling. Overexpression and activation of growth factor receptors often occurs in breast malignancies and tends to associate with the failure of antiestrogen therapy. Substantial data now show that both nuclear and extranuclear ERs interact in a cooperative fashion with transmembrane growth factor receptor signaling pathways in breast and other cancers that bear ERs, such as non-small cell lung tumors. This cross-communication between estrogen and growth factor receptors promotes downstream signaling for tumor cell proliferation, survival, and endocrine resistance. Defining the molecular interactions between estrogen and growth factor receptor signaling pathways that lead to tumor progression is essential to develop novel therapeutics that target this critical signal transduction axis.

Hormonal therapy was first used more than 100 years ago, marking the start of the current era of targeted antitumor treatment (1). This approach is based on blocking the activity of estrogens and their receptors, ER $\alpha$  and ER $\beta$  (2). The classic mechanism of hormone action involves estrogen binding to ER in the nucleus, thus promoting association with specific estrogen response elements in the promoters of target genes (3). However, ER also regulates the expression of many genes without direct binding to DNA. This occurs via protein-protein interactions with other transcription factors, such as activator protein-1, and with extranuclear signaling complexes that, in turn, modulate downstream gene expression (2, 4). A focus of this review will be extranuclear signaling that occurs in

seconds to minutes, with initiation at the plasma membrane. As first reported 30 years ago (5–7), these rapid effects are mediated by ER in or tethered to the membrane. Acute signaling by estrogen in target cells, such as breast tumors and non-small cell lung cancer (NSCLC), is now widely reported and involves nuclear and extranuclear ERs acting in concert with growth factor receptor pathways to promote downstream signaling for cell proliferation and survival (see refs. 8–14). Defining these molecular interactions is critical for developing novel antitumor therapeutics targeting specific receptor actions.

**Structure and activity of nuclear ER.** In early concepts of estrogen action, both nuclear and extranuclear ERs were postulated, but later studies focused largely on nuclear ER and transcription (3, 11, 15). ER $\alpha$  and ER $\beta$  are products of different genes and have similar, but not identical, structure (see Fig. 1). On binding estradiol, ER $\alpha$  undergoes a change in conformation of the ligand-binding domain to form a novel hydrophobic surface that modulates binding of coactivator and corepressor proteins (2, 16–18). As a phosphoprotein, ER undergoes phosphorylation at serine and tyrosine residues after activation by ligand binding, and this contributes to receptor activity and DNA binding (11, 19–21). Ligand-bound ERs function as transcription factors by binding directly as homodimers to estrogen response elements (2, 17). However, about one third of estrogen-induced genes lack functional estrogen response elements, 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  $\kappa$ B, and signal transducers and activators of transcription (4). Blockade of ER signaling by interfering with estrogen binding to ER is the basis of the hormonal agent tamoxifen, a partial agonist that limits estrogen-stimulated proliferation in breast. Fulvestrant is a novel ER antagonist that down-regulates cellular levels of ER and, unlike tamoxifen, has no agonist activity (22). Further, inhibitors of aromatase, anastrozole, letrozole, and exemestane, reduce estrogen biosynthesis in target tissues and show some advantages over tamoxifen as breast tumor therapies (23–25).

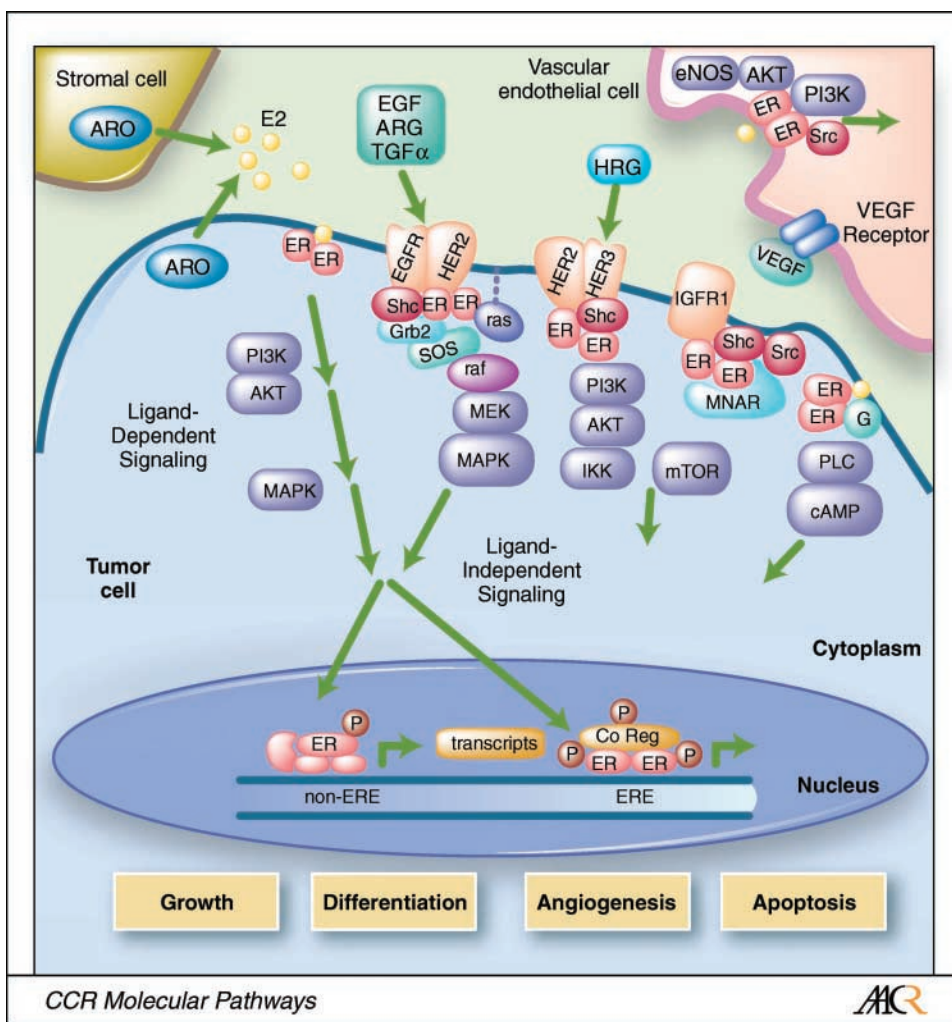
**Nature and activity of extranuclear ERs.** ER $\alpha$  gene codes for a major 66-kDa transcript and a minor 46-kDa isoform lacking portions of the NH<sub>2</sub>-terminal region of full-length ER $\alpha$  (26, 27). Although most ERs localize in tumor cell nuclei, a significant pool of ERs occurs in extranuclear sites in cell lines and archival breast cancer and NSCLC cells (14, 28). ER $\alpha$  associated with membrane is detected by the use of controlled homogenization procedures with quantitative subcellular fractionation to limit extraction artifacts (11, 29) and by antibodies directed to different domains of nuclear ER $\alpha$  in intact breast (30–32), NSCLC (13, 14), and pituitary tumor cells (33), as well as in nonmalignant vascular cells (34). The 46-kDa ER also occurs in membranes of endothelial (27) and breast (35)

**Authors' Affiliation:** Department of Medicine-Division of Hematology/Oncology, David Geffen School of Medicine, and Jonsson Comprehensive Cancer Center, University of California at Los Angeles, Los Angeles, California  
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**Requests for reprints:** Richard J. Pietras, Department of Medicine-Hematology/Oncology, Geffen School of Medicine, University of California at Los Angeles, 11-934 Factor Building, Los Angeles, CA 90095-1678. Phone: 310-825-9769; Fax: 310-825-6192; E-mail: rpietas@ucla.edu.

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**Fig. 1.** Interactions of estrogen and growth factor receptor signaling in human tumors. The proliferation and survival of human breast cancer and NSCLC cells is closely regulated by growth factor receptors as well as the activity of estrogens (*E2*) and their receptors, ER $\alpha$  and ER $\beta$  (2). ER $\alpha$  has six major functional domains including an NH $_2$ -terminal transactivation domain, an adjacent DNA-binding domain, and a portion involved in hormone-binding, receptor dimerization, and activity of a second transactivation region (16–18). In classic models of estrogen action, estrogen binds ER to promote dimerization and phosphorylation of the receptor (19). This allows direct binding of the ligand-ER complex with steroid receptor coactivators and estrogen response elements (*ERE*) in DNA, leading to changes in gene transcription that regulate growth, differentiation, apoptosis, and angiogenesis. These estrogen-dependent actions in the nucleus are modulated by ligand structure, receptor subtype (ER $\alpha$ , ER $\beta$ ) or isoform (splice variants), the gene promoter, and the balance of coactivator and corepressor proteins that modulate the transcriptional response (2, 17). In addition, there are alternate pathways of estrogen action that involve protein-protein interactions and do not require direct ER binding to DNA (4). A subset of ERs associate with extranuclear sites, caveolae, or lipid raft domains in plasma membrane, and there interact with transmembrane growth factor receptors such as EGFR, HER2, and insulin-like growth factor receptor I (*IGFR1*) and other signaling molecules, including components of the ras-MAPK and phosphatidylinositol 3-kinase (*PI3K*)/AKT pathways, Shc, Src kinases, Janus-activated kinase/signal transducer and activator of transcription signaling, nitric oxide synthase (*NOS*), and G-proteins (see refs. 11, 12, 49). Membrane-associated ER may undergo posttranslational modification, such as palmitoylation, and/or associate with adaptor proteins, such as Shc, MNAR, or lipid raft proteins. Growth factor and ERs may form a structured complex for signal transduction to MAPK and/or phosphatidylinositol 3-kinase/AKT kinase that interacts, in turn, with nuclear ER and steroid receptor coactivators. Signaling for cell growth involves phosphorylation (*P*) of nuclear ER and coactivator proteins, and such phosphorylation can occur in ligand-dependent as well as ligand-independent modes. Estrogen response element – dependent and alternate transcription sites may be activated. Further, estrogens are produced locally in breast cancer and NSCLC cells and in host supporting cells via the action of aromatase (*ARO*), and aromatase is regulated by both nuclear and extranuclear ERs and growth factor – mediated signaling. In addition, estrogens may regulate tumor-associated angiogenesis by direct interactions with vascular endothelial cells or by indirect stimulation of vascular endothelial growth factor (*VEGF*) secretion from tumors. Pathways derived from data previously published (21); refer to text for details.

cells where it may form part of a signaling complex. To assess the nature of membrane ER, nuclear ER $\alpha$  gene was transfected in ER-null Chinese hamster ovary cells, and this resulted in cellular expression of both membrane and nuclear ERs (36). Studies based on knockdowns of ER $\alpha$  by small interfering RNA (35) or ER antisense *in vitro* (37) and knockouts of both ER $\alpha$  and ER $\beta$  *in vivo* (38) support the hypothesis that membrane and nuclear ERs share a common origin. Further, membrane ERs do not occur in ER-negative MCF-7 breast tumor subclones that lack nuclear ER, and these cells, unlike ER-positive MCF-7

cells, do not show rapid estrogen-induced phosphorylation of steroid receptor coactivator AIB1 (35). Importantly, recent studies using mass spectroscopy provide evidence that peptides derived from ER $\alpha$  occur in membrane fractions prepared from breast tumor (39) and vascular endothelial cells (38). Together, these findings affirm that membrane-associated ER derives from the same gene as nuclear ER $\alpha$  (34–39). How ER associates with membranes is another challenging question. Although ER $\alpha$  has many hydrophobic regions on Kyte-Doolittle plots, there are no documented transmembrane,

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glycosylphosphatidylinositol-anchor, or PDZ domains to foster membrane association (26). However, localization and activity of ER at plasma membrane is stimulated by posttranslational palmitoylation of Cys<sup>447</sup> in the ligand-binding domain of ER $\alpha$  (14, 27, 40). Further, a highly conserved nine-amino-acid motif in the ligand binding domains of human ER $\alpha$  and ER $\beta$  seems to be necessary for mediation of receptor palmitoylation and subsequent steroid signaling (12). The membrane receptor may also form homodimers or couple with other proteins, such as caveolins, flotillins (12, 35), and coactivator protein modulator of nongenomic actions of the estrogen receptor (MNAR)/proline-, glutamic acid-, and leucine-rich protein-1 (PELP1; ref. 41). Transmembrane growth factor receptors may also provide docking sites (32, 42). Currently, most findings indicate that rapid and specific estrogen-induced membrane events are mediated predominantly by classic or variant ER forms (11, 12, 27, 34). However, other membrane estrogen-binding proteins, such as G protein-coupled receptor GPR30, or additional molecules may contribute to biological or promiscuous estrogen signaling in cells with or without ER expression (43, 44), but their role in regulating cell proliferation and survival is uncertain (38).

**Estrogen and growth factor receptor synergy in malignant progression.** Members of the *erbA* and *erbB* gene families cooperate in cell transformation (45, 46). Because ER is part of the *erbA* gene family, and epidermal growth factor receptor (EGFR)/HER-1 and HER2 are members of the *erbB* gene family, biological interactions between these receptors may be operative in human cancer progression. ER is an important prognostic factor in breast cancer that predicts response to endocrine therapy. However, resistance to endocrine therapy usually emerges, leading to tumor progression and poor patient survival. Cooperative interactions between growth factor receptor and ER signaling pathways have been identified in breast cancer and NSCLC, and growth factor-mediated pathways, notably those of EGFR, HER2, and insulin-like growth factor receptor I, are critical in development of antiestrogen resistance in breast tumors (47–49). About 20% to 25% of breast cancers have overexpression of HER2, and increased HER2 expression correlates with poor clinical outcome and resistance to endocrine therapy (47, 48). Similarly, overexpression of EGFR in ~50% of breast cancers also correlates with hormonal resistance (47, 50).

**Ligand-dependent and ligand-independent extranuclear ER signaling.** Growth factor receptors such as EGFR and HER2 often concentrate in specific domains of plasma membrane, termed caveolae or lipid rafts, together with other signal transduction molecules. Lipid rafts are moving cholesterol-rich platforms in membrane that provide a matrix for signal transduction (11, 12, 35, 51); caveolae and caveolae-like rafts are specialized forms of lipid rafts containing structural proteins, caveolin or flotillin, respectively (35, 51). Extranuclear ERs also localize in lipid rafts (11, 12, 35), thereby promoting activation and transactivation of EGFR and HER2 receptors (11, 32, 48) and interactions with other signaling molecules including insulin-like growth factor receptor I, the p85 regulatory subunit of phosphatidylinositol 3-kinase, G-proteins, Src, and Shc, a protein that may couple ER with growth factor receptors (11, 12, 42, 52). Coregulators, MNAR/PELP1 or metastasis-associated protein 1, also sequester ER in the extranuclear compartment to increase membrane action (41). Acti-

vation of these pathways by estrogen relays downstream proliferative and survival signals via mitogen-activated protein kinase (MAPK) and AKT. Further, MAPK stimulated by EGFR or/and HER2 signaling can, in turn, phosphorylate nuclear ER and receptor coactivators such as AIB1/SRC-3 (2, 35, 53, 54). These events can be triggered by estrogen (ligand-dependent signaling) or receptor kinases in the absence of estrogen (ligand-independent ER activation), with the latter process underlying many forms of endocrine resistance (52, 55).

### Clinical-Translational Advances

**Clinical significance of interactions between estrogen and growth factor signaling pathways.** A major problem in breast cancer management is the conversion from estrogen-sensitive to hormone-resistant disease after starting antiestrogen therapy (17, 47). Emergence of estrogen-independent tumors is due, in part, to enhanced interactions between growth factor receptors and ER during cancer progression that elicit ligand-independent ER activation, thus negating responses to antiestrogens (47, 55, 56). ER<sup>+</sup> and/or PR<sup>+</sup>, HER2-overexpressing tumors respond poorly to endocrine therapy (57–61), and a meta-analysis of seven clinical studies indicates that metastatic breast cancers with HER2 overexpression are tamoxifen resistant (62). In this setting, HER2 overexpression triggers signaling for increased ER phosphorylation (19, 48, 63) and loss of the inhibitory effect of tamoxifen on ER-mediated transcription (49, 61, 64–66). Recent clinical data on breast tumors also suggest that the ER<sup>+</sup>/PR<sup>-</sup> phenotype, in particular, associates with high expression of extranuclear ER and enhanced AKT signaling (28) and may identify tumors driven primarily by growth factor receptor signaling (47, 49, 67). It will be important to further examine the ER<sup>+</sup>/PR<sup>-</sup> phenotype to determine if it signals a greater likelihood to respond to growth factor receptor-targeted therapy in combination with hormonal treatment.

Tumor cells may also circumvent pharmacologic blockade of estrogen response pathways by enhanced expression of aromatase to increase estrogen biosynthesis. Aromatase induction occurs in HER2-overexpressing tumors (68, 69) and in those with active cross-talk between ER and growth factor-mediated signaling pathways (69). Laboratory models show that long-term estrogen deprivation that mimics low estrogen levels produced by aromatase inhibition eventually elicits adaptive changes in tumor cell behavior, with emergence of hypersensitivity to estrogen and increased aromatase activity (70, 71). In addition, estrogen hypersensitivity associates with acquired overexpression of growth factor receptors and enhanced estrogen binding to membrane-associated ER $\alpha$ , leading to interactions with Shc, insulin-like growth factor receptor I, and EGFR and activation of downstream signaling (42, 52, 71, 72). Together, these acquired changes allow tumors to be stimulated by much smaller amounts of estrogen, thus negating benefits of treatment with aromatase inhibitors. This complex modulation of estrogen sensitivity by tumors may support use of sequential endocrine therapy with agents having different mechanisms of action (22, 47).

**Approaches to overcome hormonal resistance.** Gaps in our understanding of mechanisms of endocrine action and resistance remain. However, based on the notion that multiple signaling pathways activate ER, combination therapies with

both endocrine and nonendocrine agents that block different pathways are rational and testable. Thus, a strategy to treat patients with ER<sup>+</sup>, HER2<sup>+</sup> tumors is dual blockade of HER2 and ER-dependent pathways that interact to promote growth. In preclinical work, combined treatment with HER2 receptor antibody, Herceptin, and tamoxifen enhanced antitumor effects in HER2-overexpressing cells with ER (48, 65, 73). Moreover, combination of Herceptin with fulvestrant is even more effective in blocking growth of tumors expressing HER2 and ER (65, 74). Fulvestrant down-regulates ER, resulting in reduced cell proliferation, and this action favors improved responses to growth factor-mediated ER activation (47). These dual treatment approaches are now being tested in the clinic (47, 75). Data from the TAnDEM trial in patients with metastatic breast cancer similarly indicate that treatment with Herceptin combined with anastrozole is superior to anastrozole alone (76), providing clinical confirmation for a working biological hypothesis.

There is also potential to use growth factor receptor tyrosine kinase inhibitors, alone or combined with antihormone agents, to treat endocrine-resistant breast cancer (47, 50, 73, 74). *In vitro* studies show that gefitinib, an EGFR kinase inhibitor, suppresses proliferation of breast cancer cells (77) and interferes with growth factor receptor and ER cross-communication (78, 79). Similarly, other inhibitors of downstream kinases, such as p42/p44 MAPK or p38 MAPK (47, 49), which are activated with tamoxifen resistance, may have future clinical utility.

In NSCLC, new research shows significant expression of ER forms that interact cooperatively with EGFR to stimulate tumor growth (13, 14). Preclinical studies reveal that either gefitinib or erlotinib, combined with fulvestrant, elicits marked inhibition of tumor progression. Results of a phase I clinical trial using fulvestrant together with gefitinib indicate that administration of this couplet is safe and shows antitumor efficacy in patients with advanced NSCLC (80), and a phase II trial with the combination of fulvestrant and erlotinib is under way (75).

EGFR and HER2 are responsible for some, but not all, states of endocrine resistance. Strategies to disrupt signal transduction by independent signaling pathways for cell proliferation and/or for blocking apoptosis are also being considered, including inhibition of insulin-like growth factor receptor I, Src, phosphatidylinositol 3-kinase, AKT, and mammalian target of rapamycin (see refs. 47, 52, 75, 81, 82).

Finally, the tumor itself is not the only therapeutic target. Cancer progression depends on formation of an adequate blood supply (83), and vascular endothelial growth factor promotes growth of breast cancer and NSCLC cells (83, 84). Vascular endothelial growth factor is increased by estradiol in breast cancers, and tumor angiogenesis is also driven by direct interaction of estradiol with vascular endothelial cells that contain ER and proliferate in response to estrogen (27, 85). Thus, it is reasonable to target estrogen signaling in both the tumor and the host vasculature, potentially offering more effective antitumor therapy than treating the tumor alone.

### The Path Forward

Our understanding of the mechanisms of estrogen action and the biological basis of resistance to hormonal therapies is advancing rapidly. However, it is now clear that there are multiple routes by which ER-mediated pathways may be activated, and this complexity enables tumors to use adaptive mechanisms for sustained growth after initiation of endocrine therapy. As we go forward, it will be important to determine the clinical significance of ER subtypes and isoforms (12, 47) and to investigate the role of tumor stem cells that bear ER and function as progenitor cells in the etiology of hormone-dependent cancers (55, 86). Currently, combination therapies directed to extranuclear and nuclear ER as well as to growth factor signaling pathways are being assessed in the clinic, and hopefully, these new strategies will offer improved antitumor efficacy for patients afflicted with hormone receptor-positive tumors.

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