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Estrogen action and cytoplasmic signaling pathways. Part II: the role of growth factors and phosphorylation in estrogen signaling

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Abstract

In recent years, distinct signaling pathways involving specific complexes of cytoplasmic proteins have been shown to orchestrate estrogen action. These pathways might supplement or augment genomic effects of estrogen that are attributable to transcriptional activation by liganded receptor. Signals might be transduced through phosphorylation of the estrogen receptors (ERs), or indirectly through effects upon transcriptional coactivators or cell receptors. Estrogen signaling is coupled to growth factor signaling with feedback mechanisms directly impacting function of growth factor receptors. These signaling pathways regulate important physiological processes, such as cell growth and apoptosis. Here, we focus on cytoplasmic signaling pathways leading to activation of ERs.

> Recent developments have clarified the role of cytoplasmic proteins and signaling pathways leading to activation of the estrogen receptors (ERα and ERβ). Here, we derive common themes from recent reports of cytoplasmic signaling modules that influence estrogen action. Ligand-dependent or ligand-independent ER activation has been shown to be accomplished by compounds other than steroidal estrogens, such as cAMP, dopamine and growth factors through signaling pathways that involve cytoplasmic proteins or protein kinases (reviewed in Ref. [1]).

> We approach this complex topic from the perspective that estrogen signaling is more complicated than the canonical 'genomic' pathway. It is useful to conceptualize estrogen action *in vivo* not as a strictly linear signal, but rather as acting through collateral, possibly divergent pathways. In contrast to experimental model systems that are overly simplistic, *in vivo* estrogen might activate cytoplasmic pathways in some tissues before transcription, and these pathways might enhance genomic actions of estradiol or influence cell function before (or in the absence of) gene transcription. Because the designation 'nonclassical' or 'nongenomic' is imprecise, we focus on specific cytoplasmic signaling pathways.

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The genomic pathway

The effects of estrogen in hormone-responsive tissues are caused by changes in gene expression modulated by the ERs that are members of the nuclear hormone receptor superfamily of transcription factors. The ERs are modular transcription factors containing two transcription activation functions, $_{AF-1}$ (see Glossary) located in the N-terminal A/B domain, and AF-2 located within the C-terminal ligand-binding domain. Although AF-2 function is dependent upon ligand binding, AF-1 functions independent of ligand binding but synergizes with AF-2 in the promotion of ligand-dependent transcription activation by the receptor (reviewed in Ref. [2]). Ligand binding by ERs produces a conformational change in their AF-2 domains that allows them to bind transcriptional coactivators and recruit them to responsive promoters. Numerous cofactors have been implicated in ER action, including p160 family members, p300/cBP, PCAF, DRIP/ARC/TRAP complexes, and other factors (reviewed in Ref. [3]). Recruitment of coactivator complexes results in histone acetylation by cofactors, chromatin remodeling and recruitment of RNA polymerase II to the promoter.

Estrogen and the ERK1/2, p44/42 MAPK pathway

Interest in the role of $_{EGF}$ in estrogen action arose from the findings that EGF mimicked estrogen action, and that antibodies to EGF inhibited estrogen-induced growth [4]. Coupling between EGF and estrogen was shown to involve ERα, the nuclear localization of which was promoted by EGF [5]. Activation of the EGF-R resulted in phosphorylation of Ser118 of ER α by p44/42 MAPK, an ERK [6,7] (Fig. 1). The physiological relevance of this pathway was underscored by the absence of estrogen-mediated EGF effects in α ERKO mice [8].

Tremblay *et al.* showed that phosphorylation of Ser124 in the AF-1 domain of mouse $ER\beta$ (analogous to Ser118 in human ER α) increases receptor interaction with sRC-1 [9]. The effect is blocked by PD98059 (which binds $M/NEK1$ and blocks activation of ERKs), but is not inhibited by the P_{KC} inhibitor, staurosporine [9]. The effect was independent of AF-2 [6,7,9]. This model is not without controversy, however, because phosphorylation of Ser118 in ERα was not found to be essential for coactivator recruitment by AF-1 of ER α [10]. In fact, E_2 led to phosphorylation of Ser118 by a mechanism that did not involve activation of p44MAPK in MCF-7 cells [11]. Ser118 might therefore be phosphorylated by kinases other than ERKs under different conditions or in different tissues. In support of this, Ser118 was phosphorylated upon ligand-dependent association of ERα with TFIIH by CDK7, which is a component of CAK, a cyclin-dependent kinase in the transcription complex [12].

Not only does phosphorylation by MAPK activate $ER\alpha/\beta$, but E_2 also influences activation of MAPK. EGF-mediated DNA synthesis was attenuated by the antiestrogen ICI164 384 [5]. *In vivo*, estrogen-induced activation of MAPK and PI3K (but not PKA or PKC) led to mitogenesis in MCF-7 cells, which was blocked by PI3K inhibitors and PD98059 [13]. A link between cytoplasmic signaling and cell-cycle regulation was suggested by the demonstration that a transcriptionally inactive ER induced mitogenesis upon estrogen treatment of NIH3T3 cells [14]. PD98059 inhibited MAPK phosphorylation in both αERKO and wild-type cortical explant cultures, but ICI182 780 blocked ERK phosphorylation only

mobilization of intracellular calcium stores in MCF-7 cells through a pathway that does not involve Raf-1 or $_{\mathbb{P}3}$ [16]. The authors suggest that E₂ activation of MAPK involves calcium as second messenger; thus, there is a feed-forward system with E_2 activating MAPK, and MAPK activating ER [16].

 $E₂$ was reported to stimulate the _{TGF} -β3 promoter in the presence of EGF [17], an effect that was blocked by PD98059 or a PKC inhibitor (GF109203), but not by the PKA inhibitor H89; however, the effect did not require phosphorylation of Ser118, because the ERαmutant S118A behaved similar to the wild-type receptor [17]. Gene regulation *in vivo* might therefore be more complex than phosphorylation of Ser118 alone [17].

Some transcriptional coactivators recruited by liganded ERs are also targets of the p44/42 MAPK pathway. Rowan *et al.* showed that SRC-1 expressed in COS-1 cells is phosphorylated and demonstrated the phosphorylation *in vitro* of SRC-1 by MAPK [18]. AIB-1 was also phosphorylated *in vitro* by MAPK and coexpression of constitutively active MEK1 enhanced the ability of a GAL4_{DBD}-AIB-1 chimera to activate a reporter [19]. The effect was attributed to increased recruitment of $p300$ and a consequent increase in HAT activity. Janknecht *et al.* demonstrated phosphorylation of CBP *in vitro* by MAPK and showed that overexpression of p44MAPK and constitutively active Raf-1 enhanced transcriptional activation by a GAL4DBD–CBP chimera [20]. Additional studies showed MAPK-dependent activation of CBP by NGF in PC12 cells and of CBP and p300 by phenylephrine in cardiac cells [21,22]. Mechanistic consequence was addressed by Ait-Si-Ali *et al.* [23], who showed that phosphorylation of CBP by p44MAPK stimulates its HAT activity.

Association of ERα with Src–_{SH}₂

A multiprotein complex of ERKs and nuclear receptors and SH2/SH3 regions of tyrosine kinase receptors has been described. Migliaccio *et al.* [24] found that antiestrogens blocked progestin-stimulated proliferation in breast cancer cells, and suggested that ERα links the Src-p21Ras-ERK pathway to ligand-activated PRB. Ligand-dependent binding of PR to SH3 domains involved a polyproline region in the N-terminus of PR_B [25]. In the human prostate cell line (LNCaP) E_2 led to association of ER β with Src and activated the Src–Raf-1–ERK2 pathway through a tertiary complex with $_{AR}$, $ER\beta$ and Src. Binding of ER α to SH2 required phosphotyrosine residue 537 (443 in ERβ) [26]. A Src/Shc/ERK pathway involving ERα, ERβ and AR was demonstrated both in primary calvaria cell cultures and an osteocytic cell line [27]. At present the relationships of these complexes to the previously reported Ras– Raf–MAPK pathway remains uncertain, but is extremely intriguing, because it involves a direct link between ER and growth factor receptors (Fig. 2).

ER activation by p38 MAPK

ERα and ERβ might be activated by $p38$ MAPK (Fig. 1). MEKK1, but not RAF or MEKK2, increased ligand-dependent expression of an $ERE-dr$ reporter in Ishikawa cells and the effect was blocked by ICI182 780, but not PD98059 [28]. BRX augmented activation of ERβ

in the presence of ligand, an effect that was blocked by the p38MAPK inhibitor, SB202190, but not by PD98059 [29]. Brx binds to nuclear receptors, including ERα [30]. Activation of ER α by Brx required Cdc42 [30], a small GTP-binding protein upstream of β _{NK} and p38MAPK. Furthermore, MEKK6-EE, a constitutively active mutant known to activate p38MAPKs, increased ERβ activity [29]. Interestingly, Brx contains a region with sequence identity to a protein A-kinase anchoring protein [30,31], suggesting that Brx might target substrates to specific subcellular regions.

Estrogen and 4-hydroxy-tamoxifen induced apoptosis in stably transfected ER-positive HeLa cells, and the effect was blocked by SB203580, a p38MAPK inhibitor [32]. ER was required, because the effect was not seen in ER-negative HeLa and MCF-10F cells [32]. Src enhanced ERα AF-1 activity through two independent mechanisms, one involving Ser118 (Raf-1–MEK–ERK) and a distinct MEKK– N KK– N KK pathway that did not. PD98059 only partially inhibited Src activation, but a dominant negative Rac mutant (S17N) and a dominant negative inhibiting MEKK1 mutant (K432M) abolished it [33]. Curiously, in HeLa cells, p38MAPK and ERKs inhibited the response, an effect that could be caused by a lack of an accessory protein not supplied in this system. It might not be possible to distinguish between a p38 pathway and a JNK-dependent effect based solely on the information provided [33], because a p38 inhibitor, such as SB202190, was not used, and MEKK1 might indirectly influence p38 activation via JNKK1 [28]. With one exception [33], reports suggest that p38-dependent activation of ER is limited to ligand-dependent effects [28,29,30].

PKA and estrogen signaling

PKA, a cAMP-dependent protein kinase, has been shown to augment the function of ER [34,35]. Aronica *et al.* described ligand-independent activation of an ERE-containing reporter [36]. Estrogen treatment rapidly increased cAMP levels in uterus and breast cancer cells [36]. PKA catalytic subunits α or β were also shown to enhance activity of an estrogendependent reporter [37]. Because PKA alters the ER agonist or antagonist profiles of tamoxifen and other $SENR$ activation pathway is of potential clinical significance. Phosphopeptide mapping confirmed that estrogens, ICI164 384, PKA and PKC promoted phosphorylation of ERα on Ser104, 106 and 118 [38]. The pathway led to ligandindependent activation of endogenous promoters of estrogen-responsive genes [39]. Three mechanisms involving PKA have been described: direct phosphorylation of the ER, phosphorylation of cofactors and effects mediated through CREB.

Studies in HeLa cells showed that EGF and PKA activation were distinct pathways leading to ER α activation [40]. PKA-mediated ER activation was abolished by cholera toxin α T/IBMX activation and the PKA inhibitor H-89, but not by the PKC inhibitor BIMD [40]. By contrast, EGF-stimulated ER activation was not blocked by either inhibitor [40]. As expected, EGF did not require the $_{\text{LBD}}$ of ER, but activation of PKA did require AF-2. Consistent with the notion of separate pathways, mutation of ERα Ser118 did not abolish the cAMP-mediated activation [40]. Experiments in αT3 cells showed ligand-independent stimulation of ERαdependent promoter activity to be blocked by H-89 and antiestrogens, but not by PD98059 [39]. Lamb *et al.* reported ligand-independent stimulation of ERα activity involving cAMP

and cyclin D1 [40]. An association between ERα and cyclin D1 has been described [41]. Lamb *et al.* found that activation of cAMP enhanced formation of the cyclin D1–ERa complex and they proposed a role for cyclin D1 and estrogen in mammary development [40]. Because PKA activation of ERα is cell-type specific [37], and is promoter context dependent [42,43], it is likely that factors, in addition to cyclin D1, contribute to PKAmediated ER activation.

PKA and ER phosphorylation

Phosphorylation of ERα is known to enhance receptor function, yet it is unlikely that PKAmediated effects are limited to direct phosphorylation of ER. Chen *et al.* found that PKA phosphorylated Ser236 of ERα *in vitro*, resulting in inhibition of dimerization [44]. E₂ removed the inhibition, but ICI182 780 did not. The functional effects of Ser236 phosphorylation were tested in COS-1 cells, in which overall ligand-independent activation was modest [44]. For the ERE-G-CAT reporter, mutation of Ser236 to glutamate (HEGO236E) did not abolish PKA-augmented ligand-dependent activation, which suggests that ligand-dependent activation by PKA does not involve Ser236 [44]. Some reports suggest the AF-1 region of ERα is not essential to PKA-mediated activation, because mutants lacking the region were activated by cAMP [37,44], and the response was blocked by antiestrogens [36,37,43].

Phosphorylation of cofactors

Transcription cofactors recruited to promoters by liganded ERs are also targets for PKA. In COS-1 cells the ability of SRC-1 to enhance PR activity was increased by 8-Br-cAMP [45]. The effect was dependent on MAPK and was not caused by direct phosphorylation of SRC-1 by PKA [45]. CBP activity can also be enhanced by the PKA pathway [21,46–48]. Assessment of the ability of GAL4DBD–CBP chimeras to activate reporters showed that signaling from PKA to CBP did not require MAPK activity because it was not blocked by PD98059 [21] or enhanced by RasR12 [47]. The ability of a GAL4DBD–CBP chimera lacking HAT activity to activate a reporter was enhanced by 8-Br-cAMP [48]. By contrast, HAT activity of CBP was required for activation of the transcription factor Pit-1 by forskolin [46].

CREB-mediated estrogen effects

Recently, Lazennec *et al.* examined the mechanism of PKA synergy with E_2/ERa in chinese hamster ovary and SK-BR-3 breast cancer cells and found that the effect was cell-line dependent [49]. However, mutation of potential phosphorylation sites in ERα (Ser236, 305, 338 and 518) did not negate the synergy observed, suggesting that phosphorylation of ERα was not required [49]. The limiting factor was shown to be CREB [49]. *In vivo*, acute and long-term treatment with estrogen altered CREB protein levels in rat central nervous system tissues [50]. Furthermore, estrogen led to rapid phosphorylation of CREB in cortical tissues [51,52]. Studies of mouse dopaminergic neurons showed that E_2 – $_{BSA}$ stimulated neuron dendritic growth through a signaling pathway involving the rapid release of intracellular calcium stores requiring PKA activity that led to phosphorylation of CREB [53]. The effect

was not inhibited by ICI182 780 [53]. The differences between cell lines and tissues suggest that mechanisms of PKA-mediated estrogen signaling might vary greatly among tissues.

Activation of ER by PKC

PKC is not a single entity, but comprises 11 isoforms, and has been shown to promote activation of ERα [35,54] by a mechanism distinct from that of EGF or PKA [55,56]. Activation of PKC by tpa led to phosphorylation of Ser118 of ERα [54]. Lahooti *et al.* examined the pathway using murine ERα in the liganded and unliganded state and found that receptor activity was stimulated by PKC δ , a Ca²⁺-dependent protein kinase, but not by PKC α or ε [56]. The effect required Ser122 of murine ER (analogous to Ser118 in human ERα) [56]. In agreement with the observations of others, stimulation was cell-type and promoter specific [56].

Evidence suggests that several PKC signaling pathways exist. An IP3-PKCa pathway in HepG2 cells was sensitive to inhibition with ICI182 780 and the $_{PLC}$ inhibitor U73122 [57]. In α T3–1 cells, however, α_{NRH} treatment stimulated estrogen-independent activation of an ERE-containing reporter via a pathway sensitive to both PKC inhibitors and PD98059 [58]. The authors suggest a PKC-MAPK-linked signaling pathway in these cells. Membraneassociated PKC-linked signaling pathways might exist that do not involve ERα/β. Sylvia *et al.* [59] described a membrane-linked estrogen-induced growth pathway in rat chondrocytes whereby E_2 –BSA led to increases in PKC α within three min, an effect that was not inhibited by ICI182 780. This pathway required phosphatidylinositol-PLC, rather than phosphatidylcholine-PLC or phospholipase D, and was coupled to a G-protein sensitive to inhibition by GDPβS [60].

For PKC and ER (or E_2) a feedback system is present. PKC has been shown to affect binding of labeled estradiol in osteoblasts [61] and the mouse uterus [62]. In the latter, inhibition of PKC led to increased estradiol binding in the cytosol, presumably because of post-transcriptional or post-translational modification of the ER by PKC α or PKC β _{II} [62]. In endometrial cancers, PKCα expression is inversely correlated with the presence of ER as determined by western blot [63]. Similar observations were made in breast cancers [64]. In rat or rabbit corpora lutea or cultured rat granulosa cells, estrogen treatment led to a 2-3-fold upregulation of PKCδ expression [65]. By contrast, in MCF-7 cells, estrogen treatment led to reduction in PKCδ mRNA and protein, but to no changes in other PKC isoforms [66]. Both upregulation and downregulation occurred over five days, a time course much longer than that in rat chondrocytes, in which it occurred within 90 min. The upregulation of PKCδ is of interest, because PKCδ is involved in growth regulation of MCF-7 cells [67]. Based on these observations, multiple feedback pathways appear to link PKC and estrogen signaling.

Feedback of estrogen to IGF-IR and EGF-R

Not only do growth factors such as EGF and $_{IGF-I}$ promote activation of ERs, but signals from activated ERs are also transmitted to transmembrane receptor complexes for these factors. Estrogen elicited tyrosine phosphorylation of the EGF receptor [68] and caused association of ER α with α F-IR and phosphorylation of the receptor [69]. E₂ also enhanced IGF-IR signaling in MCF-7 cells, but not by an increase in IGF-IR protein expression [70]. E_2 led to

an increase in IRS-1 expression and the p85 regulatory unit of PI3K at 48 h [70]. In MCF-7 cells, this resulted in increased PI3K activity and Akt phosphorylation by estrogen in combination with IGF-I [70]. These findings are interesting given direct association of ERα with the p85 subunit of PI3K (reviewed in Ref. [71]), and are consistent with the theme of synergistic activity between tyrosine kinase receptors and estrogen. Notably, IGF-IR and ER colocalize in the same cells in the female rat brain [72], as well as in reproductive tissues.

Conclusions and perspectives

It is clear that growth factors transmit signals to the genome and coactivators through nuclear receptors, including ER. Findings from several laboratories indicate the presence of integrated feedback loops involving growth factors and estrogen signaling. Recent reports describe specific complexes involving multiple receptors with cytoplasmic domains of growth factor receptors, thus revealing at least one molecular mechanism responsible for the coupling of growth factors and estrogen action.

A key feature of cytoplasmic signaling pathways of estrogen action is the heterogeneity, cell-line, tissue- and factor-specific natures of the responses. Assembly or disassembly of distinct cytoplasmic modules can dictate cell- and tissue-specific responses with remarkable plasticity that might contribute to regulatory flexibility in response to estrogens. The biological context of the signal (estrogen) determines the nature of the response, because specific messages might be transmitted differently, depending upon the temporal existence of a specific cytoplasmic estrogen-signaling module. Specificity of action is accomplished through subcellular compartmentalization of cytoplasmic signaling modules, The proteins that serve as scaffolds for these modules are just beginning to be discovered.

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Glossary

TGF $β$ transforming growth factor $β$

Fig. 1.

Cytoplasmic signaling pathways influencing the action of ER. Activation of EGF-R by EGF leads to Ras-Raf-MEK-Erk1/2 (MAPK)-mediated phosphorylation of ERα at Ser118. Activation of p38MAPKs leads to activation of ERα/β. PKCδ has also been shown to affect ER activation. In some pathways, unidentified intermediates have been indicated by question marks. Not all signaling pathways are shown. Abbreviations: see Glossary.

Fig. 2.

A possible molecular mechanism coupling transmembrane receptors to ER signaling. ER binds directly to the SH2 region of EGF through the carboxyl region. AR associates with ER, and also interacts with the SH3 region of EGF-R through the N-terminus of AR. Binding of EGF might contribute to augmentation of estrogen action through this molecular mechanism. Abbreviations: see Glossary.