



A New Role for Estrogen Receptor α in Cell Proliferation and Cancer: Activating the Anticipatory Unfolded Protein Response

Mara Livezey¹, Ji Eun Kim¹ and David J. Shapiro^{1,2*}

¹ Department of Biochemistry, University of Illinois, Urbana, IL, United States, ² Center for Cancer Research, University of Illinois, Urbana, IL, United States

Cells react to a variety of stresses, including accumulation of unfolded or misfolded protein, by activating the endoplasmic reticulum (EnR) stress sensor, the unfolded protein response (UPR). The UPR is highly conserved and plays a key role in the maintenance of protein folding quality control and homeostasis. In contrast to the classical reactive mode of UPR activation, recent studies describe a hormone-activated anticipatory UPR. In this pathway, mitogenic hormones, such as estrogen (E_2), epidermal growth factor, and vascular endothelial growth factor rapidly activate the UPR in anticipation of a future need for increased protein folding capacity upon cell proliferation. Here, we focus on this recently unveiled pathway of E_2 -estrogen receptor α (ER α) action. Notably, rapid activation of the anticipatory UPR pathway is essential for subsequent activation of the E_2 -ER α regulated transcription program. Moreover, activation of the UPR at diagnosis is a powerful prognostic marker in ER α positive breast cancer. Furthermore, in cells containing ER α mutations that confer estrogen independence and are common in metastatic breast cancer, the UPR is constitutively activated and linked to antiestrogen resistance. Lethal ER α -dependent hyperactivation of the anticipatory UPR represents a promising therapeutic approach exploited by a new class of small molecule ER α biomodulator.

Keywords: estrogen, estrogen receptor α , rapid extranuclear signaling, unfolded protein response, calcium, breast cancer, cancer therapy

OPEN ACCESS

Edited by:

Antonio Brunetti,
Università degli studi Magna Græcia
di Catanzaro, Italy

Reviewed by:

Riccardo Sgarra,
University of Trieste, Italy
Eva Surmacz,
Temple University, United States

*Correspondence:

David J. Shapiro
djshapir@life.illinois.edu

Specialty section:

This article was submitted
to Cancer Endocrinology,
a section of the journal
Frontiers in Endocrinology

Received: 09 April 2018

Accepted: 31 May 2018

Published: 15 June 2018

Citation:

Livezey M, Kim JE and Shapiro DJ
(2018) A New Role for Estrogen
Receptor α in Cell Proliferation and
Cancer: Activating the Anticipatory
Unfolded Protein Response.
Front. Endocrinol. 9:325.
doi: 10.3389/fendo.2018.00325

INTRODUCTION

The endoplasmic reticulum (EnR) plays a key role in the synthesis, folding, and transport of proteins and is important in lipid synthesis (1, 2). Maintenance of protein folding and lipid homeostasis is critical for cell proliferation and viability. The unfolded protein response (UPR) is an EnR stress-response pathway that senses and responds to diverse stimuli, including changes in EnR luminal calcium, redox status, nutrient availability, lipid bilayer composition, and accumulation of unfolded or misfolded protein (3, 4). The UPR consists of three arms, IRE1 α , ATF6 α , and PERK that together decrease the flux of new protein into the EnR, while simultaneously increasing production of molecular chaperones to help fold unfolded or misfolded proteins. IRE1 α and PERK are activated upon oligomerization and autophosphorylation. ATF6 α is activated and transported to the Golgi apparatus, where it

Abbreviations: EnR, endoplasmic reticulum; TYS or TDG, T47D cells expressing either ER α Y537S or ER α D538G.

is cleaved by S1P and S2P proteases, although the mechanism of activation in the EnR is still unclear. There is increasing evidence that all three arms of the UPR can be activated in more than one way. For example, some of the earliest work suggested that the molecular chaperone, BiP, blocked oligomerization, and activation of IRE1 α and PERK through direct binding to their luminal domains (2). Upon accumulation of unfolded or misfolded proteins, BiP would be competed away, allowing activation of these UPR arms. Similarly, it is thought that upon depletion of EnR calcium, calcium-dependent molecular chaperones, such as BiP, fall off IRE1 α , and PERK, and other unfolded or misfolded proteins. This would allow IRE1 α and PERK to oligomerize and activate the UPR (5). Additional experiments and elucidation of the crystal structure of the luminal domain of IRE1 α showed that independent of BiP binding, IRE1 α can directly bind and be activated by peptides *via* an MHC-like structural domain (2). Interestingly, recent work has also suggested that IRE1 α and PERK may sense and be activated by changes in lipid content of the EnR membrane, independent of accumulation of unfolded protein, or depletion of calcium in the EnR (6).

Activation of the non-canonical RNase IRE1 α (inositol-requiring enzyme 1 α) results in alternative splicing of the transcription factor XBP1, leading to the production of spliced-XBP1 (sp-XBP1) and upregulation of molecular chaperones (7). ATF6 α (activating transcription factor 6 α) is translocated to the Golgi apparatus where it is cleaved by proteases to produce the transcription factor p50-ATF6 α that also upregulates chaperone production (8). Finally, activated PERK (protein kinase RNA-like EnR kinase) phosphorylates eIF2 α , resulting in transient inhibition of most protein synthesis, while promoting translation and production of selective proteins, including ATF4, CHOP, p58^{IPK}, and GADD34 (9). When UPR stress is mild, the chaperone p58^{IPK} binds to and inhibits PERK, and GADD34 dephosphorylates eIF2 α . Working together, p58^{IPK} and GADD34 reverse PERK activation and restore protein synthesis.

Classically, the UPR is activated in response to EnR stress. Several years ago, it was shown that progenitors of immunoglobulin-producing B cells activate the UPR before initiating antibody production. This pathway, which is activated in the absence of unfolded protein, was named the anticipatory UPR by Walter and Ron (2), but it was not studied extensively. We, and others, recently showed that diverse steroid and peptide hormones, including estrogen (17 β -estradiol, E₂), progesterone (P₄), epidermal growth factor, and vascular endothelial growth factor, and probably the insect hormone ecdysone (Ec), activate an anticipatory UPR pathway to prepare cells for the increased protein folding that accompanies cell proliferation (10–14). Notably, the steps between hormone receptor complexes and activation of the three arms of the UPR have largely been identified (10–12).

The proliferative and anti-apoptotic advantage of overexpressing or activating hormone receptors, such as EGF receptor (EGFR) or estrogen receptor α (ER α), has long been appreciated in cancer biology (15–18). However, hormone activation of the anticipatory UPR has only recently become a focus of cancer research and exploited as a therapeutic target. This review focuses on the role of E₂-ER α activation of the anticipatory UPR and a promising preclinical drug candidate, BHPI, which leverages this

novel action of ER α in order to block proliferation of and kill most ER α positive breast cancer cells.

ACTIVATION OF THE ANTICIPATORY UPR BY MITOGENIC HORMONES

Steroid and peptide hormones exert their effects through binding and modulating their specific receptors (18, 19). Using E₂-ER α as an example, when hormone receptors bind to their ligand, they dimerize and are recruited to specific DNA response elements (**Figure 1**). E₂-ER α then modulates the activity of thousands of genes either through direct binding to DNA, or by tethering of E₂-ER α to other transcription factors (20–22). The genomic actions of E₂-ER α are important for the pro-proliferation properties of E₂ in ER α positive breast cancer cells and while rapidly initiated, play out over hours or days.

In addition to classical genomic actions, E₂-ER α exerts rapid extranuclear actions important for activating signal transduction pathways (**Figure 1**). These pathways are important for diverse actions of E₂-ER α , crosstalk with the genomic program, and are rapidly activated and often play out over minutes to hours (23, 24). Of these pathways, activation of the anticipatory UPR is the most recently described (**Figure 2**) (12). Upon binding of E₂ to ER α at the plasma membrane, there is rapid activation of phospholipase C γ , resulting in cleavage of its substrate PIP₂ to IP₃ (inositol triphosphate) and DAG (diacylglycerol). The IP₃ then binds to and opens IP₃ receptor (IP₃R) calcium channels in the membrane of the EnR, allowing efflux of calcium out of the lumen of the EnR into the cell body. The modest decrease in EnR calcium caused by E₂ treatment of ER α positive cancer cells weakly activates the UPR, resulting in upregulation of molecular chaperones along with minimal and very transient inhibition of protein synthesis. By knockdown of UPR components, or blocking calcium release from the EnR, we showed that increased intracellular calcium from activation of the anticipatory UPR is critical for subsequent E₂-ER α -mediated modulation of gene expression and cell proliferation (12).

Consistent with E₂-ER α activation of the anticipatory UPR, T47D breast cancer cells modified with CRISPR-Cas9 to replace wild-type ER α with the constitutively active mutations ER α Y537S (TYS cells) and ER α D538G (TDG cells) upregulate the UPR in the absence of estrogen (13). Strikingly, YYS and TDG cells express higher levels of the molecular chaperones BiP and p58^{IPK} and YYS cells have increased levels of sp-XBP1. Surprisingly and possibly due to robust upregulation of progesterone receptor in these cell lines, P₄ further elevates some of these downstream products of mild UPR activation, including sp-XBP1, and in YYS cells, p58^{IPK}. Of note, the synthetic progestin R5020 increases resistance of YYS and TDG cells to OHT (4-hydroxy tamoxifen; the active form of tamoxifen) and fulvestrant/ICI 182,780 (ICI). In anchorage-independent three-dimensional (3D) assays, R5020-treated YYS and TDG cells exhibited robust proliferation in high concentrations of OHT and fulvestrant/ICI (13).

In diverse cancers, mild UPR activation is protective and helps cancers to proliferate, induce angiogenesis, and overcome hypoxia and toxic stress from chemotherapies (25, 26). Using microarray and outcome data from approximately 1,000 patient

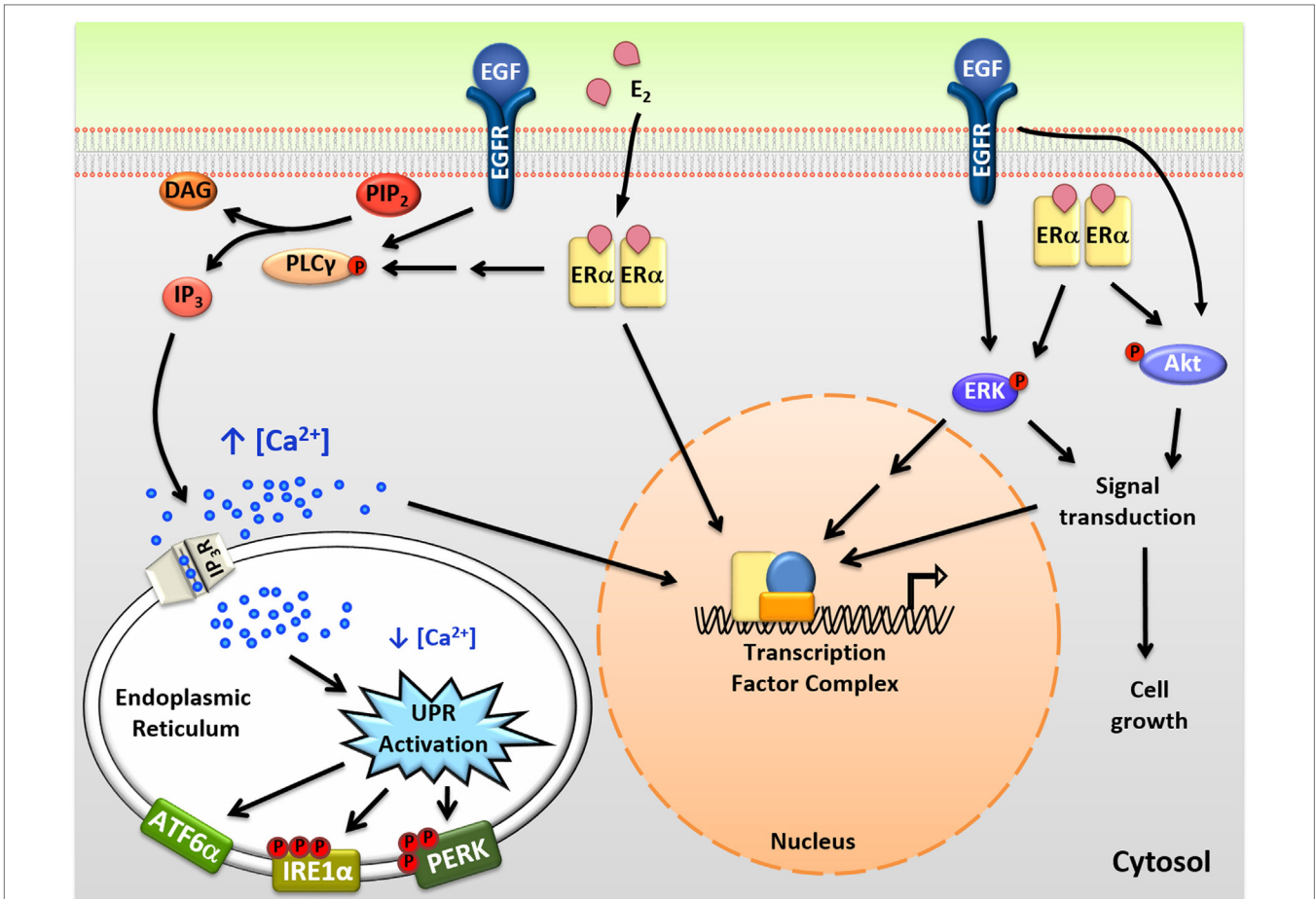


FIGURE 1 | Intracellular actions of mitogenic hormones. Estrogen (E_2) and epidermal growth factor (EGF) act on their respective receptors, estrogen receptor α (ER α), and EGF receptor (EGFR), to initiate crosstalk between extracellular signaling pathways and their genomic programs. ER α indirectly and EGFR directly activate phospholipase C γ (PLC γ), resulting in cleavage of PIP $_2$ to DAG (diacylglycerol) and IP $_3$ (inositol triphosphate). IP $_3$ then binds to IP $_3$ receptors (IP $_3$ Rs) in the endoplasmic reticulum (EnR) membrane, causing moderate efflux of calcium from the lumen of the EnR into the cell body. This calcium signal activates all three arms of the unfolded protein response (UPR) and acts as an authorizing signal for E_2 -ER α and EGF-EGFR modulation of gene expression and cell proliferation. In parallel, E_2 -ER α and EGF-EGFR modulate additional extracellular signal transduction pathways, including activation of ERK and Akt signaling. Activation of these pathways is also important for subsequent cell proliferation and crosstalks with the E_2 -ER α and EGF-EGFR genomic programs.

breast cancers, we demonstrated the significance of this pathway in ER α positive breast cancer. Increased expression of a UPR gene index consisting of UPR components and UPR-induced chaperones strongly correlated with reduced time to recurrence, subsequent resistance to tamoxifen, and reduced survival (12). The close correlation between the extent of activation of the UPR gene index and activation of E_2 -ER α regulated genes is consistent with ER α playing a major role in the elevated expression of the UPR gene index (10). Moreover, in triple negative breast cancer in which ER α is absent, the IRE1/XBP1 axis plays a central role in tumorigenicity and progression, and the extent of activation of an XBP1 gene index is correlated with reduced patient survival (27, 28). Taken together, this suggests that ER α -mediated activation of the anticipatory UPR likely plays an important role in early survival of breast cancers. At later times when therapeutic stress is added to nutritional deprivation and hypoxia, activation of the classical reactive UPR makes an important contribution to tumor survival (26, 29–31).

Because of the protective nature of the UPR in cancer, drugs that target components of the UPR are in preclinical development, in clinical trials, and have been approved (27, 28, 32). Most commonly, these drugs inhibit key components of the UPR, such as PERK, IRE1 α RNase, or the downstream chaperone BiP/GRP78. Unfortunately, due to lack of drug specificity, these drugs may have toxic effects in tissues with a large secretory burden, such as pancreas.

UPR HYPERACTIVATION AS A TOOL TO SELECTIVELY TARGET ER α POSITIVE BREAST CANCER

The standard of care for ER α positive breast cancer is endocrine therapy, including aromatase inhibitors that block E_2 production, and tamoxifen and fulvestrant/ICI that compete with E_2 for binding to ER α . Unfortunately, many tumors that were initially

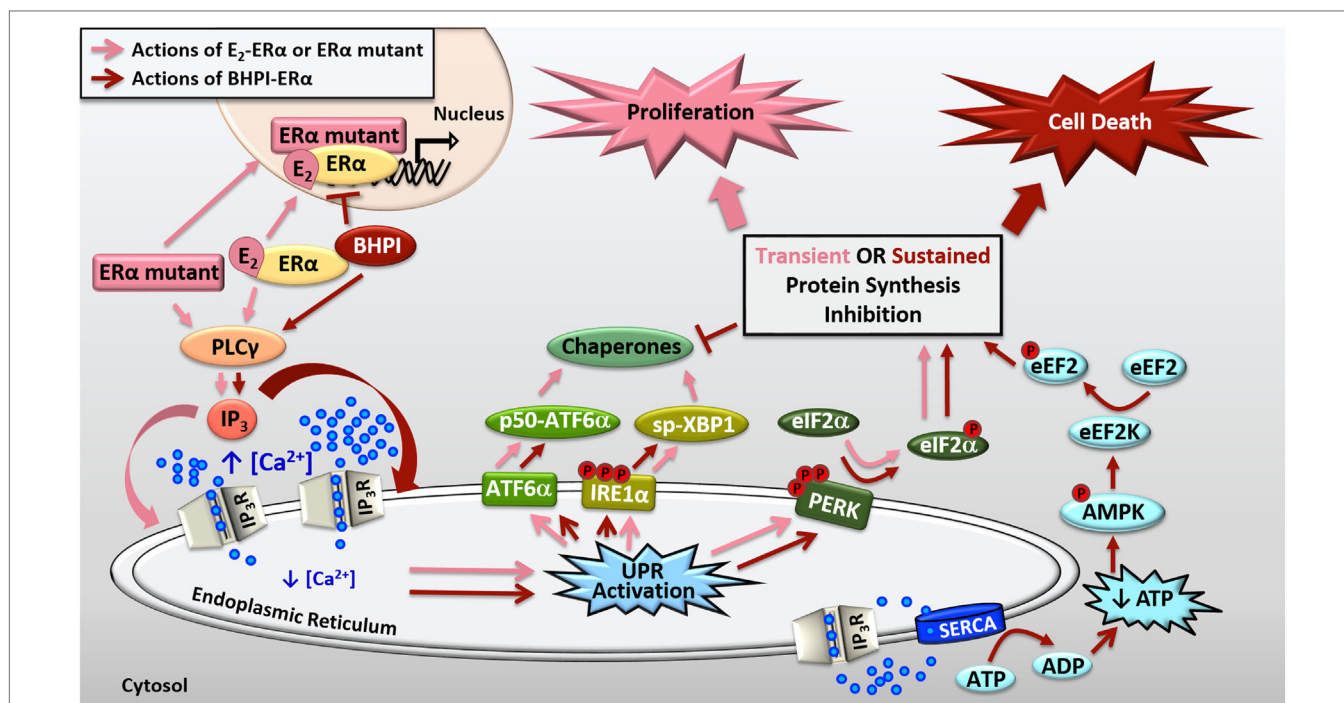


FIGURE 2 | Activation of the anticipatory unfolded protein response by estrogen receptor α (ER α). E₂-ER α and constitutively active ER α mutants activate a mild and protective anticipatory unfolded protein response (UPR) and the non-competitive biomodulator BHPI binds ER α and induces hyperactivation of this pathway leading to cell death. ER α indirectly activates phospholipase C γ (PLC γ), resulting in cleavage of PIP₂ to DAG (diacylglycerol) and IP₃ (inositol triphosphate). E₂-ER α and constitutively active ER α mutants cause moderate IP₃ production, whereas BHPI causes significantly more production of IP₃. The IP₃ then binds to IP₃ receptors (IP₃Rs) in the endoplasmic reticulum (EnR) membrane, causing efflux of calcium from the lumen of the EnR into the cell body. E₂-ER α and constitutively active ER α mutants cause moderate and transient release of calcium, resulting in weak and transient activation of all three arms of the UPR. Weak UPR activation results in very mild and transient inhibition of protein synthesis, production of molecular chaperones, and is critical for subsequent cell proliferation. BHPI-ER α induced hyperactivation of the UPR causes robust and sustained release of calcium from the EnR. This leads to robust PERK activation and rapid, sustained, and near-quantitative inhibition of protein synthesis. Although BHPI causes upregulation of chaperone mRNA, no protein is made, and the UPR-activating signal is never resolved. In an effort to re-establish cellular calcium homeostasis, ATP-dependent SERCA pumps in the EnR actively transport calcium back into the lumen of the EnR but since IP₃Rs remain open, an ATP-depleting futile cycle ensues. Decreased cellular ATP and increased AMP activate AMPK, which along with calcium, activates Ca²⁺/calmodulin-dependent kinase, eukaryotic elongation factor 2 kinase (CAMKIII/eEF2K). eEF2K then phosphorylates eEF2, causing inhibition of protein synthesis at elongation. Ultimately, BHPI-ER α induced hyperactivation of the anticipatory UPR causes death of ER α positive endometrial and breast cancer cells.

responsive recur after years of treatment. Moreover, there is selection and outgrowth of endocrine therapy resistant tumors expressing ER α mutations in about one-third of patients with advanced metastatic breast cancer, most commonly ER α Y537S and ER α D538G (33–35). Structural and biophysical studies suggest that estrogen receptors containing these mutations are stabilized in the active conformation and have lower affinity for antiestrogens, such as OHT (36). Additionally, a growing body of clinical evidence suggests that mutations in this ligand binding domain hotspot confer partial resistance to endocrine therapies (33–35, 37). Strikingly, patients whose tumors express ER α Y537S or ER α D538G have on average 12 and 6 months shorter survival, respectively, than patients whose tumors express wild-type ER α (38). We have shown that TYS and TDG cells containing these estrogen receptor mutations exhibit constitutively active ER α , allowing E₂-independent proliferation and gene expression, and partial resistance to the endocrine therapies OHT and fulvestrant/ICI (13). Additionally, compared to wild-type ER α in T47D cells, we observed resistance to fulvestrant/ICI-mediated

degradation of the mutant ER α s in TYS and TDG cells. Because of the resistance to endocrine therapies observed in cancers expressing ER α Y537S and ER α D538G, development of better selective estrogen receptor modulators and degraders (SERMs and SERDs) has been a focus in targeting these cancers (39–42).

Recently, we described a novel small molecule biomodulator, BHPI, that selectively targets ER α positive cancer cells (13, 43, 44). BHPI [3,3-bis(4-hydroxyphenyl)-7-methyl-1,3-dihydro-2H-indol-2-one] is a bis-phenylated oxindol. In a limited structure-activity-relationship study, addition of methyl groups to both phenyl rings significantly disrupted activity of BHPI (43). We demonstrated specificity by testing over 30 ER α positive and negative cell lines and showed that BHPI only inhibits proliferation of or kills cells that express ER α (43). Additionally, in the isogenic human breast cell lines MCF10A (ER α negative) and MCF10A_{ER IN9} (ER α positive), we showed that BHPI was only effective in the cells expressing ER α , and was ineffective when ER α was knocked back down in MCF10A_{ER IN9} cells. Demonstrating that BHPI physically interacts with ER α , BHPI

shifts the tryptophan emission spectrum of ER α , and protects peptides in the ER α ligand binding domain from protease digestion. Furthermore, BHPI inhibits recruitment of ER α to E₂-ER α regulated promoters (**Figure 2**). However, BHPI is not a competitive inhibitor, as it does not compete with radiolabeled E₂ for the ligand binding pocket and is equally effective in the presence and absence of estrogen (43). Rather than inhibiting a component of the UPR, BHPI takes advantage of the already moderately elevated UPR in cancer cells by hyperactivating the anticipatory UPR (**Figure 2**). BHPI, therefore, hijacks the normal protective actions of ER α activation of the UPR in order to push UPR activation into the lethal range. This is the first small molecule to modulate the action of a hormone receptor in this way.

We showed that BHPI blocks proliferation of ovarian cancer cells and kills most ER α positive breast and endometrial cancer cells (13, 43, 45). Compared to E₂, BHPI causes increased production of IP₃ in cancer cell lines (43, 45). This increased production of IP₃ hyperactivates the UPR through sustained opening of IP₃R calcium channels in the EnR, resulting in robust and sustained calcium release from the lumen of the EnR into the cell body (**Figure 2**). While E₂ causes mild and transient inhibition of protein synthesis, BHPI causes a rapid, sustained, and near-quantitative inhibition of protein synthesis in ER α positive breast and endometrial cancer cells. Surprisingly, BHPI also causes rapid depletion of intracellular ATP. Disruption of cytosolic calcium homeostasis can be toxic, specifically, high levels of calcium can lead to cell death (45–47). To restore intracellular calcium homeostasis after opening of IP₃Rs, ATP-dependent SERCA pumps calcium back into the EnR lumen, but because IP₃Rs remain open, the calcium leaks back out. This creates a futile cycle of calcium leakage and pumping that depletes intracellular ATP. Additionally, ATP depletion from prolonged SERCA pump activity results in increased levels of cellular AMP that activates the metabolic sensor AMPK. AMPK activation together with high levels of cytosolic calcium activates the Ca²⁺/calmodulin-dependent kinase, eukaryotic elongation factor 2 kinase (CAMKII/eEF2K). eEF2K then phosphorylates eEF2, which inhibits protein synthesis at a second site (**Figure 2**). Therefore, although BHPI increases the mRNA levels of chaperones, such as BiP and p58^{IPK}, no protein is made, leading to un-resolvable cytotoxic activation of the UPR. While other activators of the classical reactive UPR share similarities to BHPI's mechanism of action, such as disruption of EnR calcium homeostasis and inhibition of protein synthesis (1, 2), BHPI is unique in its ability to cause ATP depletion in cancer cells.

We recently described the efficacy of targeting breast cancer cells expressing ER α Y537S and ER α D538G with BHPI (13). In 3D culture, OHT and fulvestrant/ICI only partially inhibited growth of TYS and TDG cells and R5020 completely reversed antiestrogen inhibition of growth. In contrast, BHPI killed TYS and TDG cells in the presence or absence of R5020. Since BHPI is not a competitive inhibitor of ER α (43) and targets ER α positive cancer cells irrespective of their dependence on E₂ for proliferation, it is a promising preclinical drug candidate for the treatment of metastatic breast cancers expressing ER α Y537S and ER α D538G.

In ovarian cancer, a common mechanism for resistance to the taxane paclitaxel and other chemotherapy agents is overexpression of ATP-dependent efflux pumps, especially Multidrug Resistance Protein 1 (MDR1)/P-glycoprotein/ABCB1. Despite intensive efforts, effective and non-toxic MDR1 inhibitors have remained elusive. Due to its ability to deplete intracellular ATP, BHPI inhibited ATP-dependent MDR1-mediated drug efflux and restored sensitivity of multidrug-resistant breast and ovarian cancer cells to killing by therapeutically relevant concentrations of several anticancer drugs (44). Using multidrug resistant OVCAR-3 cells, BHPI was tested in an orthotopic ovarian cancer xenograft model. Although paclitaxel was ineffective against these tumors, BHPI alone strongly reduced tumor growth. Notably, tumors were undetectable in mice treated with BHPI plus paclitaxel. After the combination therapy, plasma levels of the widely used cancer biomarker, CA125, were at least several hundred fold lower than in mice with control tumors. Moreover, CA125 levels progressively declined to undetectable in all mice treated with the combination therapy (44).

CONCLUSION

Studies of the pro-proliferative effects of mitogenic hormones and their respective receptors have long focused on their actions on genomic programs and on extranuclear signal transduction pathways. Activation of the anticipatory UPR is an emerging, very rapid action of many mitogenic hormones that authorizes subsequent gene expression and cell proliferation. Important for targeting hormone receptor positive breast cancers is the finding that they exhibit elevated UPR activation. This UPR activation can be exploited by small molecules that hyperactivate the pathway, pushing UPR activation into the lethal range. As a first-in-class small molecule, BHPI is a model for investigating hyperactivation of the anticipatory UPR as a promising strategy for killing ER α positive breast cancer cells. A similar approach is also likely viable for breast cancers that overexpress other hormone receptors that activate the anticipatory UPR, such as progesterone receptor, or EGFR family members. However, agents that hyperactivate the anticipatory UPR through these receptors have yet to be identified. Thus, the anticipatory UPR is a key pathway for development of new anticancer drugs that can help overcome resistance to current therapies.

AUTHOR CONTRIBUTIONS

ML and DS contributed to writing and revising. JK contributed to revising the manuscript.

ACKNOWLEDGMENTS

We are grateful to L. Yu for the initial version of **Figure 1**. The research described in this review was supported by grants DOD BCRP W81XWH-13 and NIH RO1 DK071909 and the E. Howe Scholar award (to DS), and by an NSF Graduate Research Fellowship under DGE-1144245 (to ML).

REFERENCES

- Ron D, Walter P. Signal integration in the endoplasmic reticulum unfolded protein response. *Nat Rev Mol Cell Biol* (2007) 8:519–29. doi:10.1038/nrm2199
- Walter P, Ron D. The unfolded protein response: from stress pathway to homeostatic regulation. *Science* (2011) 334:1081–6. doi:10.1126/science.1209038
- Urrea H, Dufey E, Avril T, Chevet E, Hetz C. Endoplasmic reticulum stress and the hallmarks of cancer. *Trends Cancer* (2016) 2:252–62. doi:10.1016/j.trecan.2016.03.007
- Korennykh A, Walter P. Structural basis of the unfolded protein response. *Annu Rev Cell Dev Biol* (2012) 28:251–77. doi:10.1146/annurev-cellbio-101011-155826
- Preissler S, Chambers JE, Crespillo-Casado A, Avezov E, Miranda E, Perez J, et al. Physiological modulation of BiP activity by trans-protomer engagement of the interdomain linker. *Elife* (2015) 4:e08961. doi:10.7554/eLife.08961
- Volmer R, Ron D. Lipid-dependent regulation of the unfolded protein response. *Curr Opin Cell Biol* (2015) 33:67–73. doi:10.1016/jceb.2014.12.002
- Cox JS, Walter P. A novel mechanism for regulating activity of a transcription factor that controls the unfolded protein response. *Cell* (1996) 87:391–404. doi:10.1016/S0092-8674(00)81360-4
- Yoshida H, Haze K, Yanagi H, Yura T, Mori K. Identification of the cis-acting endoplasmic reticulum stress response element responsible for transcriptional induction of mammalian glucose-regulated proteins. Involvement of basic leucine zipper transcription factors. *J Biol Chem* (1998) 273:33741–9. doi:10.1074/jbc.273.50.33741
- Novoa I, Zeng H, Harding HP, Ron D. Feedback inhibition of the unfolded protein response by GADD34-mediated dephosphorylation of eIF2 α . *J Cell Biol* (2001) 153:1011–22. doi:10.1083/jcb.153.5.1011
- Karali E, Bellou S, Stellas D, Klinakis A, Murphy C, Fotsis T. VEGF signals through ATF6 and PERK to promote endothelial cell survival and angiogenesis in the absence of ER stress. *Mol Cell* (2014) 54:559–72. doi:10.1016/j.molcel.2014.03.022
- Yu L, Andruska N, Zheng X, Shapiro DJ. Anticipatory activation of the unfolded protein response by epidermal growth factor is required for immediate early gene expression and cell proliferation. *Mol Cell Endocrinol* (2016) 422:31–41. doi:10.1016/j.mce.2015.11.005
- Andruska N, Zheng X, Yang X, Helferich WG, Shapiro DJ. Anticipatory estrogen activation of the unfolded protein response is linked to cell proliferation and poor survival in estrogen receptor α -positive breast cancer. *Oncogene* (2015) 34:3760–9. doi:10.1038/ncr.2014.292
- Mao C, Livezey M, Kim JE, Shapiro DJ. Antiestrogen resistant cell lines expressing estrogen receptor α mutations upregulate the unfolded protein response and are killed by BHPI. *Sci Rep* (2016) 6:34753. doi:10.1038/srep34753
- Liu W, Cai M-J, Zheng C-C, Wang J-X, Zhao X-F. Phospholipase C γ 1 connects the cell membrane pathway to the nuclear receptor pathway in insect steroid hormone signaling. *J Biol Chem* (2014) 289:13026–41. doi:10.1074/jbc.M113.547018
- Normanno N, De Luca A, Bianco C, Strizzi L, Mancino M, Maiello MR, et al. Epidermal growth factor receptor (EGFR) signaling in cancer. *Gene* (2006) 366:2–16. doi:10.1016/j.gene.2005.10.018
- Deroo BJ, Korach KS. Estrogen receptors and human disease. *J Clin Invest* (2006) 116:561–70. doi:10.1172/JCI27987
- Evans RM. The steroid and thyroid hormone receptor superfamily. *Science* (1988) 240:889–95. doi:10.1126/science.3283939
- Huang P, Chandra V, Rastinejad F. Structural overview of the nuclear receptor superfamily: insights into physiology and therapeutics. *Annu Rev Physiol* (2010) 72:247–72. doi:10.1146/annurev-physiol-021909-135917
- Katzenellenbogen BS. Dynamics of steroid hormone receptor action. *Annu Rev Physiol* (1980) 42:17–35. doi:10.1146/annurev.ph.42.030180.000313
- York B, O'Malley BW. Steroid receptor coactivator (SRC) family: masters of systems biology. *J Biol Chem* (2010) 285:38743–50. doi:10.1074/jbc.R110.193367
- Carroll JS, Liu XS, Brodsky AS, Li W, Meyer CA, Szary AJ, et al. Chromosome-wide mapping of estrogen receptor binding reveals long-range regulation requiring the forkhead protein FoxA1. *Cell* (2005) 122:33–43. doi:10.1016/j.cell.2005.05.008
- Hah N, Danko CG, Core L, Waterfall JJ, Siepel A, Lis JT, et al. A rapid, extensive, and transient transcriptional response to estrogen signaling in breast cancer cells. *Cell* (2011) 145:622–34. doi:10.1016/j.cell.2011.03.042
- Levin ER. Plasma membrane estrogen receptors. *Trends Endocrinol Metab* (2009) 20:477–82. doi:10.1016/j.tem.2009.06.009
- Song RX-D, Santen RJ. Membrane initiated estrogen signaling in breast cancer. *Biol Reprod* (2006) 75:9–16. doi:10.1095/biolreprod.105.050070
- Ma Y, Hendershot LM. The role of the unfolded protein response in tumour development: friend or foe? *Nat Rev Cancer* (2004) 4:966–77. doi:10.1038/nrc1505
- Wang M, Kaufman RJ. The impact of the endoplasmic reticulum protein-folding environment on cancer development. *Nat Rev Cancer* (2014) 14:581–97. doi:10.1038/nrc3800
- Chen X, Iliopoulos D, Zhang Q, Tang Q, Greenblatt MB, Hatziaepostolou M, et al. XBP1 promotes triple-negative breast cancer by controlling the HIF1 α pathway. *Nature* (2014) 508:103–7. doi:10.1038/nature13119
- Zhao N, Cao J, Xu L, Tang Q, Dobrolecki LE, Lv X, et al. Pharmacological targeting of MYC-regulated IRE1/XBP1 pathway suppresses MYC-driven breast cancer. *J Clin Invest* (2018) 128:1283–99. doi:10.1172/JCI95873
- Wang M, Kaufman RJ. Protein misfolding in the endoplasmic reticulum as a conduit to human disease. *Nature* (2016) 529:326–35. doi:10.1038/nature17041
- Clarke R, Cook KL, Hu R, Facey COB, Tavassoly I, Schwartz JL, et al. Endoplasmic reticulum stress, the unfolded protein response, autophagy, and the integrated regulation of breast cancer cell fate. *Cancer Res* (2012) 72:1321–31. doi:10.1158/0008-5472.CAN-11-3213
- Rajapaksa G, Thomas C, Gustafsson J-Å. Estrogen signaling and unfolded protein response in breast cancer. *J Steroid Biochem Mol Biol* (2016) 163:45–50. doi:10.1016/j.jsbmb.2016.03.036
- Sykes EK, Mactier S, Christopherson RI. Melanoma and the unfolded protein response. *Cancers* (2016) 8:E30. doi:10.3390/cancers8030030
- Robinson DR, Wu Y-M, Vats P, Su F, Lonigro RJ, Cao X, et al. Activating ESR1 mutations in hormone-resistant metastatic breast cancer. *Nat Genet* (2013) 45:1446–51. doi:10.1038/ng.2823
- Toy W, Shen Y, Won H, Green B, Sakr RA, Will M, et al. ESR1 ligand-binding domain mutations in hormone-resistant breast cancer. *Nat Genet* (2013) 45:1439–45. doi:10.1038/ng.2822
- Spoerke JM, Gendreau S, Walter K, Qiu J, Wilson TR, Savage H, et al. Heterogeneity and clinical significance of ESR1 mutations in ER-positive metastatic breast cancer patients receiving fulvestrant. *Nat Commun* (2016) 7:11579. doi:10.1038/ncomms11579
- Fanning SW, Mayne CG, Dharmarajan V, Carlson KE, Martin TA, Novick SJ, et al. Estrogen receptor alpha somatic mutations Y537S and D538G confer breast cancer endocrine resistance by stabilizing the activating function-2 binding conformation. *Elife* (2016) 5:e12792. doi:10.7554/eLife.12792
- Merenbakh-Lamin K, Ben-Baruch N, Yeheskel A, Dvir A, Soussan-Gutman L, Jeselsohn R, et al. D538G mutation in estrogen receptor- α : a novel mechanism for acquired endocrine resistance in breast cancer. *Cancer Res* (2013) 73:6856–64. doi:10.1158/0008-5472.CAN-13-1197
- Chandarlapaty S, Chen D, He W, Sung P, Samoilu A, You D, et al. Prevalence of ESR1 mutations in cell-free DNA and outcomes in metastatic breast cancer: a secondary analysis of the BOLERO-2 clinical trial. *JAMA Oncol* (2016) 2:1310–5. doi:10.1001/jamaoncol.2016.1279
- Zhao Y, Laws MJ, Guillen VS, Ziegler Y, Min J, Sharma A, et al. Structurally novel antiestrogens elicit differential responses from constitutively active mutant estrogen receptors in breast cancer cells and tumors. *Cancer Res* (2017) 77:5602–13. doi:10.1158/0008-5472.CAN-17-1265
- Weir HM, Bradbury RH, Lawson M, Rabow AA, Buttari D, Callis RJ, et al. AZD9496: an oral estrogen receptor inhibitor that blocks the growth of ER-positive and ESR1-mutant breast tumors in preclinical models. *Cancer Res* (2016) 76:3307–18. doi:10.1158/0008-5472.CAN-15-2357
- Xiong R, Zhao J, Gutgesell LM, Wang Y, Lee S, Karumudi B, et al. Novel selective estrogen receptor downregulators (SERDs) developed against treatment-resistant breast cancer. *J Med Chem* (2017) 60:1325–42. doi:10.1021/acs.jmedchem.6b01355
- Bihani T, Patel HK, Arlt H, Tao N, Jiang H, Brown JL, et al. Elacestrant (RAD1901), a selective estrogen receptor degrader (SERD), has antitumor activity in multiple ER+breast cancer patient-derived xenograft models. *Clin Cancer Res* (2017) 23:4793–804. doi:10.1158/1078-0432.CCR-16-2561

43. Andruska ND, Zheng X, Yang X, Mao C, Cherian MM, Mahapatra L, et al. Estrogen receptor α inhibitor activates the unfolded protein response, blocks protein synthesis, and induces tumor regression. *Proc Natl Acad Sci U S A* (2015) 112:4737–42. doi:10.1073/pnas.1403685112
44. Zheng X, Andruska N, Lambrecht MJ, He S, Parissenti A, Hergenrother PJ, et al. Targeting multidrug-resistant ovarian cancer through estrogen receptor α dependent ATP depletion caused by hyperactivation of the unfolded protein response. *Oncotarget* (2018) 9:14741–53. doi:10.18632/oncotarget.10819
45. Berridge MJ, Bootman MD, Roderick HL. Calcium signalling: dynamics, homeostasis and remodelling. *Nat Rev Mol Cell Biol* (2003) 4:517–29. doi:10.1038/nrm1155
46. Criddle DN, Gerasimenko JV, Baumgartner HK, Jaffar M, Voronina S, Sutton R, et al. Calcium signalling and pancreatic cell death: apoptosis or necrosis? *Cell Death Differ* (2007) 14:1285–94. doi:10.1038/sj.cdd.4402150
47. Bruce JIE. Metabolic regulation of the PMCA: role in cell death and survival. *Cell Calcium* (2018) 69:28–36. doi:10.1016/j.ceca.2017.06.001

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Copyright © 2018 Livezey, Kim and Shapiro. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.