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Estradiol-Induced Regression in T47D:A18/PKC α Tumors Requires the Estrogen Receptor And Interaction with the Extracellular Matrix

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Abstract

Several breast cancer tumor models respond to estradiol (E2) by undergoing apoptosis, a phenomenon known to occur in clinical breast cancer. Prior to the application of tamoxifen as an endocrine therapy, high dose E2 or diethylstilbesterol (DES) treatment was successfully utilized, albeit with unfavorable side effects. It is now recognized that such an approach may be a potential endocrine therapy option. We have explored the mechanism of E2-induced tumor regression in our T47D:A18/PKC α tumor model that exhibits autonomous growth, tamoxifen-resistance and E2-induced tumor regression. Fulvestrant, a selective estrogen receptor downregulator, prevents T47D:A18/PKC α E2-induced tumor growth inhibition and regression when given prior or subsequent to tumor establishment, respectively. Interestingly, E2-induced growth inhibition is only observed *in vivo* or when cells are grown in Matrigel but not in two-dimensional tissue culture, suggesting the requirement of the extracellular matrix (ECM). Tumor regression is accompanied by increased expression of the pro-apoptotic Fas/FasL proteins and downregulation of the pro-survival Akt pathway. Inhibition of colony formation in Matrigel by E2 is accompanied by increased expression of Fas and shRNA knockdown partially reverses colony formation inhibition. Classical ERE-regulated transcription of pS2, PR, TGF α , C3 and cathepsin D is independent of the inhibitory effects of E2. A membrane impermeable E2-BSA conjugate is capable of mediating growth inhibition, suggesting the involvement of a plasma membrane ER. We conclude that E2-induced T47D:A18/PKC α tumor regression requires participation of ER α , the ECM, Fas/FasL and Akt pathways, allowing the opportunity to explore new predictive markers and therapeutic targets.

Keywords

PKC α ; T47D; breast cancer; Fas; apoptosis; extracellular matrix

Introduction

Currently, tamoxifen is prescribed for the treatment of all stages of estrogen receptor alpha positive (ER α +) breast cancer and was the first drug approved as a chemopreventative agent for women at high-risk for developing the disease (1). Tamoxifen belongs to the class of drugs known as selective estrogen receptor modulators (SERMs), exhibiting target site-specific

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activities in the body (2). Resistance to tamoxifen often occurs as a result of a variety of cellular changes (3) but usually is not a result of loss or mutations within the ER (4-6). Evidence of cross-talk between both the nuclear and plasma membrane ER with growth factor receptors is well-documented in the literature (7) including involvement of IGF-1R(8), HER2/neu (9-11), EGFR (HER1) (12) and PC cell-derived growth factor (PCDGF/GP88) (13). Consequently, secondary signaling effectors are also modulated including Akt (14), PTEN(15), MAPK (16, 17) and AP-1(18-20). Increased levels of coactivators such as AIB1 (21) and decreased levels of the corepressor NCOR1 have also been implicated in mediating tamoxifen resistance (22).

Prior to the introduction of tamoxifen for the treatment of breast cancer, high-dose E2 and diethylstilbesterol (DES) were often used with response rates similar to those observed with tamoxifen (23,24). Tamoxifen became the drug of choice due to the lower incidence of side effects. A report comparing DES-treated and tamoxifen-treated patients with a 20 year follow-up indicated a survival advantage for DES-treated patients (25). Another small trial conducted in postmenopausal patients with advanced breast cancer exposed to multiple endocrine therapies reported 31% of patients achieved complete or partial response to DES therapy(26). Several cell and tumor models exhibiting growth inhibition and apoptosis with E2 have now emerged (27-34). A common mechanism shared by at least three of these models is the involvement of the Fas/FasL apoptotic pathway (28,33,34). Fas is a member of the death receptor family also known as the tumor necrosis factor (TNF) superfamily that mediates the extrinsic apoptotic pathway upon activation by Fas ligand (FasL) binding in many tissues, the most well-studied of which include the immune system and the breast (35,36).

We previously reported a T47D:A18 cell line stably transfected with protein kinase C alpha (PKC α) that is hormone-independent *in vitro*, and produces tumors in athymic mice that exhibit autonomous growth, are tamoxifen-resistant and are exquisitely sensitive to E2, resulting in tumor regression and apoptosis (31,37). We have also reported the potential predictive value of PKC α overexpression in tamoxifen-resistant breast cancer in the clinic (38). PKC is a family of serine-threonine protein kinases that is comprised of at least 12 isozymes that are subdivided into three subfamilies: conventional (α , β I, β II, γ), novel (δ , ϵ , η , θ and μ), and atypical (ζ and ν / λ) (39-41). The PKC family of isozymes mediates a multitude of physiological processes in a cell-type and tissue-specific manner. The inverse relationship of PKC activity and ER status in breast cancer was initially reported several years ago (42), and our laboratory discovered the inverse relationship of PKC α and ER α in human breast cancer cell lines (37) and endometrial cancers (43). Other laboratories have described the importance of PKC α and other PKC isozymes in MCF-7 breast cancer cells in invasion (44,45) and tamoxifen-resistance (41,46,47). Specifically Frankel et al. (47) documented PKC α overexpression in 9 antiestrogen-resistant cell lines and showed that stable overexpression of PKC α in MCF-7 cells resulted in reduced antiestrogen sensitivity. Recently, Assender et al. (46) reported that PKC α expression correlates with poor clinical response to endocrine therapy.

Our T47D:A18/PKC α tumor model is unique since PKC α overexpression may be a useful tumor marker to identify patients likely to respond to an E2-like treatment regimen. We previously reported that E2 causes tumor regression *in vivo*, but does not inhibit cells grown on two-dimensional (2D) plastic (31). In this study we explored the mechanism of E2-induced tumor regression in the T47D:A18/PKC α model by focusing on the contribution of the tumor microenvironment, involvement of the Fas/FasL pathway, and the role of ER α .

Results

The role of the Fas/FasL pathway in E2-induced regression

We previously demonstrated that regression of T47D:A18/PKC α tumors by E2 is accompanied by apoptosis (31). To determine the signaling pathways that may mediate the apoptotic process

induced by E2, T47D:A18/PKC α tumors were established in thirty untreated athymic mice. Following 9 weeks, mice were randomized to continue without treatment (NT group, 10 mice) or were implanted with a 1.0 cm E2 capsule (E2 group, 20 mice) and the experiment was continued for an additional 15 days. Tumor stabilization was observed in the E2 treated mice until day 7, followed by tumor regression (Fig. 1A). The size of the E2-treated tumors was significantly reduced compared with the NT tumors at day 7 and all time points up until day 15 ($P < 0.001$). Tumors were collected at various time points from NT group and post-E2 treated mice on days 1, 2, 4, 5, 7, 8, 9, 12 and 15 following capsule implantation (Fig. 1A). Apoptosis was assessed by TUNEL assay as described in “Materials and Methods” in tumors from both the NT and E2 groups on days 8, 12 and 15. The percentage of apoptotic cells was significantly higher at all three time points in the E2 group compared with the NT group (Fig. 1B). These results are in agreement with our previous study and suggest that apoptosis contributes to E2-induced tumor regression (31).

Previous reports have indicated the involvement of the Fas/FasL pathway in mediating apoptosis by E2 in other model systems (28,33,48). To determine the participation of Fas/FasL in mediating apoptotic cell death in T47DA18/PKC α tumors, Fas/FasL protein expression was compared in NT and E2-treated tumors (Fig. 1A). Initially, the E2-treated tumors express reduced Fas protein on Day 2 and Day 4 compared to the NT tumors. However, at later time points, Fas is significantly increased in the E2-treated tumors compared with the NT tumors (Fig. 1C). The same biphasic pattern of FasL regulation is observed with an initial decrease of FasL expression in E2-treated tumors relative to NT tumors at Day 2 and Day 4, followed by increased FasL expression in E2-treated tumors compared with NT tumors on Day 8 and Day 12. Therefore, upregulation of both Fas and FasL is temporally related with E2-induced T47D:A18/PKC α tumor regression and apoptosis (Figs. 1A, C). However since upregulation of Fas/FasL is not observed prior to tumor regression it is likely that Fas/FasL-induced apoptosis may not be the sole pathway responsible for E2-induced tumor regression.

The Akt/PKB pathway is downregulated in E2-regressing tumors

Since the Fas/FasL pathway is not likely to be the only mechanism whereby E2-induced tumor regression occurs, we examined the possible involvement of the pro-survival Akt/PKB pathway. We find that both total and p473-Akt are reduced in the E2-treated tumors compared with the untreated control at all time points (Fig. 1D). These results suggest that the pro-survival, anti-apoptotic Akt pathway is inactivated by E2 treatment and therefore contributes to E2-induced tumor regression. These results suggest that repression of the pro-survival, anti-apoptotic Akt pathway precede activation of the Fas/FasL apoptotic pathway and coordinately these two pathways mediate tumor regression.

E2-induced T47D:A18/PKC α growth inhibition requires contact with the ECM

We previously reported that E2 inhibits the growth of T47D:A18/PKC α cells *in vivo*, but has no growth inhibitory effect *in vitro* (31). To determine whether the ECM is sufficient to recapitulate the *in vivo* environment, the ability of T47D:A18/PKC α and T47D:A18/neo cells to form colonies in Matrigel was examined. T47D:A18/neo cells form statistically significantly more colonies in phenol-red containing Matrigel compared with T47D:A18/PKC α cells (Fig. 2A), suggesting that the presence of E2 in the Matrigel environment inhibits colony formation. When cells are plated in phenol red-free Matrigel (E2 depleted environment), T47D:A18/PKC α colony formation is inhibited by 50% in the presence of E2 compared with the absence of E2 (Fig 2B). In the absence of E2, T47D:A18/neo cells form few colonies whereas in the presence of E2, numerous colonies are formed (Fig. 2B). These results suggest that a component provided by Matrigel is sufficient to recapitulate the E2-induced growth inhibitory effects observed *in vivo*. To determine if growth factors present in Matrigel are required to produce E2-induced growth inhibition, colony formation was examined in growth factor

reduced (GFR)-Matrigel (Figs. 2C, D). E2 retains the ability to inhibit T47D:A18/PKC α colony formation in GFR-Matrigel, exhibiting similar results when cells are plated either in phenol red-containing (Figure 2C) or phenol red-free RPMI Matrigel (Fig. 2D). Since cells grown in 2D are also exposed to growth factors (31,37), it is unlikely that growth factors present in Matrigel are responsible for mediating the E2 inhibitory effect. However, we cannot rule out this possibility since growth factors are not eliminated, but simply reduced in GFR-Matrigel. It is likely that another Matrigel component is required for E2-induced growth inhibition. E2 does not inhibit colony formation when the cells are plated in soft agar (results not shown) nor have we been able to successfully establish T47D:A18 colonies on collagen or laminin coated plates.

E2 induces apoptosis and increases Fas expression in T47D:A18/PKC α cells grown in Matrigel

To determine whether the mechanism of E2-induced growth inhibition of T47D:A18/PKC α cells grown in Matrigel is similar to the E2-growth inhibitory effects observed *in vivo*, apoptosis and Fas/FasL protein expression was examined. The percentage of apoptotic cells in the T47D:A18/PKC α colonies increased with time following E2 treatment and was significantly increased compared to the untreated control colonies at 7 and 15 days post-E2 treatment (Fig. 3A).

Since both Fas and FasL protein expression exhibited a temporal increase with concomitant tumor regression (Fig. 1A, C), Fas/FasL expression was determined in T47D:A18/PKC α colonies treated with E2 growing in Matrigel. Fas expression was increased in the E2-treated T47D:A18/PKC α colonies at all three time points (Fig. 3B, C) and is consistent with Fas upregulation observed in tumors (Fig 1C). However there was no difference in the expression of FasL in E2-treated colonies versus the untreated control (Fig. 3B, D). Neither Fas nor FasL expression is regulated in T47D:A18/PKC α cells growing in 2D (results not shown), nor does E2 cause growth inhibition of cells on 2D plastic, suggesting that the tumor microenvironment and Matrigel provide the context whereby E2 causes tumor regression or colony growth inhibition respectively.

Knock-down of Fas by shRNA partially reverses E2-inhibited colony formation in Matrigel

To further clarify the role of the Fas/FasL pathway in E2-induced apoptosis in the T47D:A18/PKC α cell model, we applied RNA interference to stably knockdown Fas. Lentiviral transduction particles containing shFas constructs were used to transfect T47D:A18/PKC α cells and selected clones were screened by western blot to confirm decreased expression of Fas. Fas expression was reduced to 80% and 60% as determined by densitometry in two Fas stable T47D:A18/PKC α clones, #6-3 and #7-9, respectively (Fig. 4A). To determine whether reduced Fas expression was sufficient to abrogate Fas/FasL-mediated apoptosis, recombinant FasL peptide was used to initiate apoptosis in the shFas stable clones in 2D tissue culture. Whereas apoptosis was increased by 2-fold in both the parental T47D:A18/PKC α cells and the negative control, T47D:A18/PKC α /NTS cells (stable transfectant with non-targeting sequence), the two shFas clones were not responsive to FasL (Fig 4B). Therefore we concluded that sufficient knockdown of Fas was achieved in the selected shFas clones and were used to study the effect on the E2-inhibitory phenotype.

To study the effect of Fas knockdown on regulation of E2-inhibition in Matrigel, colony formation assays were performed comparing the parental T47D:A18/PKC α and T47D:A18/PKC α /NTS to the shFas clones, #6-3 and #7-9. The E2-treated T47D:A18/PKC α cells formed 50% fewer colonies compared to the untreated control and the T47D:A18/PKC α /NTS cells formed 60% fewer colonies following E2-treatment (Fig. 4C). However the shFas clones formed 30-35% fewer colonies compared to the untreated control groups. Therefore Fas

interference in these clones only partially reversed the E2-inhibitory phenotype, suggesting that the Fas/FasL pathway contributes to tumor regression mediated by apoptosis, but may not be the only pathway.

Estradiol-induced inhibition of T47D:A18/PKC α tumor growth requires the ER

To determine the role of the ER in E2-induced tumor regression, the ability of the selective estrogen receptor downregulator (SERD) fulvestrant, to block E2-induced tumor regression was tested. T47D:A18/PKC α cells were injected into fifty ovariectomized athymic mice. Forty mice were not treated (NT group), 10 mice were implanted with a 1.0 cm E2 capsule and given weekly injections with fulvestrant for all 13 weeks. After 7 weeks, 40 mice from the NT group were randomized to three treatment groups: continued NT (10 mice), E2 capsule (10 mice) and E2 capsule plus weekly injections with fulvestrant (20 mice) (Fig. 5A). Fulvestrant prevented E2-induced tumor growth inhibition and regression when given prior or subsequent to tumor establishment, respectively. Since the mechanism of action of fulvestrant involves the destruction of the ER (49), these results suggest that the ER is required for E2 to exert growth inhibitory effects. This finding is in agreement with another study reporting participation of the ER in an MCF-7 tumor model exhibiting E2-induced tumor regression (33).

To examine the mechanism whereby fulvestrant rescues E2-mediated tumor regression, 40 mice were bilaterally injected with T47D:A18/PKC α cells and tumors were left untreated for 5 weeks. At that time, mice were randomized into 4 treatment groups (10 mice/group); continued no treatment, E2 capsule, fulvestrant or E2 + fulvestrant. Tumors were excised from all treatment groups 2, 4 and 8 days post-treatment for assessment of apoptosis by the TUNEL assay and expression of Fas/FasL proteins by western blot. Tumor regression was observed in both the E2-treated and fulvestrant-treated mice, whereas tumors in the untreated and E2 + fulvestrant treated mice continued to grow (Fig. 5B). However in contrast to the previous mouse experiment (Fig. 5A), tumors in the E2+fulvestrant group grew faster than the untreated control group in this experiment. TUNEL assay revealed that the level of apoptosis was greatest in the E2-treated tumors following 4 and 8 days post-treatment, with a similar trend in the fulvestrant alone tumors, whereas the E2+fulvestrant group showed no increased apoptosis compared to the untreated control tumors (Fig 5C). Tumors treated with either E2 alone or fulvestrant alone exhibited upregulation of both Fas and FasL expression, however expression of Fas/FasL in the E2 treated tumors occurred earlier (4 days) compared with the fulvestrant treated tumors, where expression of Fas/FasL appeared later (8 days) (Figs. 6A, B). This suggests that the mechanism whereby E2 induces Fas/FasL protein expression is different than the mechanism initiated by fulvestrant treatment. Interestingly, treatment with a combination of E2+fulvestrant partially suppresses tumor Fas upregulation on Day 4, and completely reverses Fas expression on Day 8. The E2+fulvestrant combination completely reverses FasL expression similar to levels observed in the untreated control tumors. Taken together these results indicate that addition of the SERD fulvestrant can reverse E2-induced apoptosis in T47D:A18/PKC α tumors, partially due to reversal of Fas/FasL expression and perhaps via a mechanism requiring the ER.

Both E2 and fulvestrant are known to cause degradation of the ER, however these ligands act via distinct pathways. Whereas E2-mediated ER downregulation is a result of transcriptional activation, coactivator recruitment and subsequent proteosomal degradation (50-53), fulvestrant stimulates ER degradation directly via disruption of nucleocytoplasmic shuttling and the ubiquitin-proteasome pathway (49,54). Treatment with either E2 alone or fulvestrant alone at 2, 4 and 8 days results in ER α protein downregulation in T47D:A18/PKC α tumors (Fig 7). This result indicates that although E2 causes regression of these tumors, the downregulation of ER α in response to E2 is as expected. While fulvestrant treatment also results in ER α downregulation as predicted, the combination of E2+fulvestrant partially reverses

ER α downregulation at all time points examined. Therefore the differential regulation of ER α may be responsible for the prevention of apoptosis and restoration of tumor growth.

Classical ER α regulated gene expression

To investigate the role of ER α in E2-mediated tumor regression, expression of genes well known to be regulated by E2 was determined in T47D:A18/PKC α tumors derived from treatment groups harvested on Day 8 as shown in Fig. 5B. We chose to determine the expression of five classical ERE-mediated E2-responsive genes, *C3* (complement component 3), *PGR* (progesterone receptor), *CTSD* (cathepsin D), *TFF1* (trefoil factor 1 or pS2) and *TGFA* (transforming growth factor α) by real-time RT-PCR (Fig. 8). Although E2 treatment causes tumor regression and addition of fulvestrant prevents regression, 4 out of the 5 classically estrogen regulated genes including *C3*, *PGR*, *CTSD* and *TFF1* are upregulated in both the E2 and E2 + fulvestrant treated tumors compared to the untreated control tumors. Despite the fact that treatment with either E2 or fulvestrant causes tumor regression, these 4 genes exhibit opposite regulation. An exception to this is *TGFA*, the expression of which appears to be upregulated in tumors from all 3 treatment groups (E2, fulvestrant, or E2+fulvestrant). These results suggest that although E2 is causing complete regression of T47D:A18/PKC α tumors, E2 nonetheless can induce classical ERE-mediated gene expression in this tumor model. Therefore, perhaps E2-induced tumor regression is mediated by the ER via a non-classical mechanism.

The plasma membrane-associated ER is implicated in the E2-inhibitory effect

To investigate the possibility that the E2-induced growth inhibitory effects may be mediated by a plasma membrane associated ER, the membrane impermeable E2-BSA conjugate was used as a tool. To check whether free E2 was present in the E2-BSA conjugate, an ERE-luciferase construct was transfected into T47D:A18/neo cells. It was concluded that insignificant levels of free E2 were present in the E2-BSA conjugate since the E2-BSA treated cells showed luciferase activity similar to the vehicle treated control cells, whereas E2 treatment resulted in 17-fold induction of luciferase activity (Fig 9A).

The ability of the membrane-impermeable E2-BSA conjugate to inhibit T47D:A18/PKC α colony formation was examined in Matrigel. Treatment with the E2-BSA conjugate resulted in slightly more T47D:A18/neo colonies compared with the untreated control group when grown in Matrigel, suggesting that E2-BSA can stimulate modest colony formation. However 3.5-fold more colonies were present in the E2-treated group compared with the E2-BSA group (Fig. 9B). Both E2-BSA alone and E2 alone inhibited T47D:A18/PKC α colony formation compared to vehicle treated control cells (Fig. 9C). These results imply that perhaps the E2-inhibitory effects may be mediated via an ER associated with the plasma membrane.

Discussion

The possibility of using an E2-like compound in patients that exhibit tamoxifen resistant breast cancer is a concept that is re-emerging (55-58), and the ability to predict *a priori* patients that would benefit from such an approach is very attractive. Our laboratory reported the correlation of PKC α overexpression with disease recurrence following tamoxifen treatment (38). We have explored the signaling events leading to E2-induced tumor regression in our unique T47D:A18/PKC α tumor model (31,37). This is a relevant model of autonomous growth, tamoxifen-resistance and E2-induced tumor regression, characteristics often encountered in the clinical setting. In this study we report the requirement of the tumor microenvironment, specifically the ECM, for E2 to elicit inhibitory effects. The ER is likely required for E2-induced tumor regression since the SERD fulvestrant prevents these effects. We provide evidence that upregulation of the Fas/FasL apoptotic pathway occurs concurrent with E2-induced tumor

regression and this pathway appears to be modulated only *in vivo* or in Matrigel, but not in 2D tissue culture. These results suggest that communication between the ECM and the tumor plays an important role in eliciting the E2-induced growth inhibitory effects. Tumor regression is also accompanied by downregulation of the Akt pro-survival pathway. ERE-mediated induction of 5 known E2-regulated genes is not altered in this model indicative of intact classical ER signaling. The membrane impermeable E2-BSA conjugate elicits growth inhibitory effects; therefore a plasma membrane form of the ER is likely involved.

PKC isozyme expression has been studied in the MCF-7 cell lines by several investigators (44,45,47). To our knowledge, our T47D:A18/PKC α cell and tumor model is the only reported breast cancer model other than MCF-7 examining the effect of PKC isozyme expression on antiestrogen resistance. In this model, although we reported cross-upregulation of PKCs β and δ (37), we have determined that coordinate overexpression of PKC δ and PKC β is not sufficient to impart autonomous, tamoxifen-resistant and E2-inhibitory growth (60).

Several reports of MCF-7 tumor models describe E2-induced growth inhibition and regression. A cyclical response to E2 in MCF-7 tumors was first reported by Yao et al., (27) whereby short-term exposure to tamoxifen (1 year) yielded tamoxifen resistant and E2 responsive tumors whereas long-term tamoxifen exposure (5 years) tumors remained tamoxifen-resistant but also acquired an inhibitory response to E2 (27). We found elevated PKC α expression in both tumor types suggesting a correlation of PKC α overexpression with the hormone-responsive phenotype (31). Shim et al. (32) demonstrated that long-term estrogen deprived cells (LTED) form tumors that can be inhibited by E2 via activation of the apoptotic Fas pathway (28). The Fas pathway also mediates the E2-induced apoptosis in MCF-7 TAMLT tumors (33) and the MCF-7/Ral tumor model (34). However T47D:A18/PKC α tumor regression is likely mediated by both pro-apoptotic signals through Fas/FasL and inhibition of pro-survival signals by downregulation of the Akt pathway. Another example of E2-induced tumor regression was established by the stable transfection of MCF-7 cells with Akt-3 (61). Tumors derived from this cell line are hormone-independent, tamoxifen-stimulated and growth inhibited by E2. Whereas all of the MCF-7 tumor models show E2-induced cell growth inhibition both *in vitro* and *in vivo*, our T47D:A18/PKC α cell model is not inhibited by E2 under 2-D culture conditions; E2 only elicits growth inhibition either *in vivo* (31) or in Matrigel. Matrigel is a solubilized basement membrane preparation derived from the Englebreth-Holm-Swarm mouse sarcoma and the major components include laminin, collagen IV, heparan sulfate proteoglycans, entactin and nidogen. The advantage of utilizing 3D models for breast cancer research is well-described and shown to better mimic the tumor microenvironment (62). T47D:A18/PKC α colony formation is inhibited in Matrigel either in media containing serum and phenol-red (Fig. 2A) or with the addition of exogenous E2 (Fig. 2B). The ability of E2 to inhibit colony formation is not impaired when T47D:A18/PKC α cells are grown in GFR-Matrigel (supplemental data), suggesting that a component in Matrigel other than growth factors is responsible for mediating the inhibitory effect. However, the inhibition of colony formation by E2 does not completely mimic tumor regression *in vivo*. Both Fas and FasL are upregulated by E2 in tumors (Fig 1C), whereas Fas alone but not FasL is upregulated in Matrigel (Fig. 3). Another distinction from the *in vivo* model is the inability of fulvestrant to reverse E2-mediated inhibition of colony formation (results not shown). Taken together these results suggest that Matrigel cannot completely recapitulate the tumor microenvironment. Stromal cells, which are obviously absent from Matrigel, may contribute to E2-induced tumor regression and apoptosis. One possibility is that E2 induces antiangiogenic signals, although no changes in VEGF was observed in tumors following E2 treatment (results not shown). There is abundant evidence in the literature that PKC interacts with integrins (63-65), specifically a direct interaction between PKC α and β 1 integrins was reported (66). Integrin expression following E2 treatment both in tumors and colonies growing in Matrigel was not altered (results not shown), but this does not rule out the possibility that enhanced PKC α /integrin signaling

may lead to secretion of an antiangiogenic factor that mediates tumor regression *in vivo*. Another potential mediator of ECM interactions is syndecan-4, a transmembrane proteoglycan known to physically interact with PKC α (67,68). Our laboratory is currently investigating this potential pathway.

Our results indicate that E2-induced tumor regression may also require the participation of the ER itself. We find that E2-induced tumor regression can be prevented by co-administration of fulvestrant (Figs. 5A, B), a SERD currently approved as a second-line endocrine therapy that binds to and causes complete destruction of the ER (69). ER α expression is downregulated in T47D:A18/PKC α tumors in response to E2 or fulvestrant alone, but is retained in tumors treated with the E2+fulvestrant combination, most evident on Day 8 (Fig. 7). E2 is reported to cause rapid downregulation of the ER that is dependent upon coactivator recruitment and new protein synthesis whereas destruction of the ER by fulvestrant is independent of these processes (70). In our T47D:A18/PKC α tumor model, ER α regulation by the combination of E2+fulvestrant appears to be dissimilar from regulation by either E2 or fulvestrant treatment alone. Perhaps this differential ER α regulation is vital in the reversal of the growth inhibitory effects observed with E2. Similar to our model, MCF-7TAMLT E2-induced tumor regression is also abrogated by fulvestrant, implicating the participation of the ER (33).

We previously reported that ER function is reduced 10-fold in T47D:A18/PKC α cells as assessed by ERE-luciferase reporter assays (37). This suggested to us that classical ERE-mediated signaling may not be the mechanism whereby the ER mediates E2-induced tumor regression. However, we find that 5 classically ERE-regulated genes, pS2, PR, TGF α , C3 and cathepsin D are upregulated in tumors from E2-treated mice (Fig. 8). Except for TGF α , transcription of the other genes is not induced by fulvestrant. Interestingly, all 5 genes are induced by the combination of E2+fulvestrant to a similar level as E2 treatment alone. We conclude that transcriptional induction of these genes is independent of tumor response to E2, therefore classical signaling is not likely to mediate tumor growth inhibition. Since interaction of T47D:A18/PKC α cells with the ECM appears to be crucial for E2 to exhibit growth inhibitory effects, an attractive scenario is that tumor growth inhibition is mediated by the plasma membrane ER (71). The plasma membrane ER interacts with G proteins affecting downstream signaling cascades including ERK, PI3K and PKC (72-74). There is evidence that estradiol can directly bind and activate PKC α at the plasma membrane through non-genomic effects in endometrial cancer cells and rat colon (75,76). The fact that the membrane impermeable E2-BSA conjugate can inhibit colony formation in Matrigel equally well as E2 (Fig. 9), suggests that the plasma membrane ER is more likely to mediate the E2-growth inhibitory effect. Alternatively, we do not rule out the genomic non-classical ER signaling with other transcription factors such as AP-1 or Sp1(77,78). We are currently pursuing these possibilities.

We described here the characterization of a tamoxifen-resistant T47D:A18/PKC α tumor model that has several similarities and distinctions with the MCF-7 derived tumor models. This T47D:A18/PKC α tumor model is a unique tool to further study potential therapeutic targets for tamoxifen resistant breast cancer, especially since we and others have evidence that PKC α overexpression may be a predictive marker of tamoxifen resistance (38,46). It may be possible to identify patients that would benefit from treatment with E2 or E2-like compounds simply by screening for PKC α expression. Perhaps the success rate of the PKC α antisense oligonucleotide Affinitak (LY900003/ISIS 3521) (79) can be improved by identification of patients with PKC α overexpressing tumors. Alternatively, the plasma membrane ER may be a potential novel therapeutic target. Finally, our findings have important implications for the application of fulvestrant and aromatase inhibitors in patients that harbor PKC α -overexpressing tumors. Based on our findings, the efficacy of fulvestrant may potentially be compromised in patients with circulating estrogens. Furthermore, E2 deprivation created by

aromatase inhibitors may cause tumor growth. We are currently testing the latter hypothesis in a preclinical tumor model.

Materials and Methods

Cell Culture Conditions

Human breast cancer cell line T47D:A18 is a hormone-responsive clone that has been previously described (80), and was maintained in phenol red-containing RPMI-1640 supplemented with 10% fetal bovine serum (FBS). Stable transfectant cell lines T47D:A18/neo and T47D:A18/PKC α (37) were maintained in RPMI-1640 (phenol-red) supplemented with 10% FBS-containing G418 (500 μ g/ml). When indicated, prior to treatment with E2 or fulvestrant for protein isolation, western blot or Matrigel experiments, cell lines were placed in phenol red-free RPMI-1640 supplemented with 10% 3 \times dextran-coated charcoal-treated FBS (E2- depleted media) for 3 days.

Growth of Cells in Matrigel

T47D:A18/neo and T47D:A18/PKC α cells were maintained either in phenol-red free RPMI (E2-depleted) media for three days or in phenol red-containing RPMI supplemented with 10% FBS. Matrigel (Becton Dickinson; Lincoln Park, NJ) was thawed overnight at 4°C and E2 (10^{-9} M), E2-BSA (10^{-9} M) or vehicle (ethanol) was added to the Matrigel (phenol red-free, phenol red-containing, or growth factor reduced Matrigel). Six-well plates were coated with 800 μ l Matrigel/well, and incubated at 37°C for 30 min. Cells were suspended at 5×10^3 cells/ml in 1.3 ml of either phenol red-containing RPMI media or phenol red-free RPMI with E2 or E2-BSA (10^{-9} M) supplemented with 500 μ g/ml G418 and spread on pre-gelled Matrigel. Plates were incubated at 37°C for 3 weeks; media was replaced to the top of the Matrigel every 3 days. Five 1.0 cm² areas were counted under 5X power, and the average number of colonies \pm SEM was determined.

Growth of T47D:A18/PKC α Tumors *In Vivo*

T47D:A18/PKC α cells were injected bilaterally (1×10^7 cells/site) into the axillary mammary fat pads of ovariectomized nude 4-6 week old athymic mice (Harlan Sprague Dawley, Madison, WI). Mice were randomized into treatment groups consisting of at least 10 mice/group. E2 was administered via silastic capsules (1.0 cm) implanted subcutaneously between the scapulae. The 1.0 cm capsules produce a mean serum E2 level of 379.5 pg/ml (81) and were replaced every 8 weeks. Fulvestrant, ICI 182,780, (a generous gift from AstraZeneca Pharmaceuticals, Macclesfield, Cheshire, UK) was injected subcutaneously at a dose of 5 mg (0.1 ml peanut oil) per animal once per week. Tumor cross sectional area was determined weekly using Vernier calipers and calculated using the formula: length/2 X width/2 X π . Mean tumor area was plotted against time in weeks to monitor tumor growth. The mice were sacrificed by CO₂ inhalation and cervical dislocation, tumors were excised and either immediately fixed in 10% buffered formalin for TUNEL assay, stored in RNAlater buffer (Ambion, Inc., Austin, TX) or snap frozen in liquid nitrogen and stored at -80°C. The Animal Care and Use Committee of the University of Illinois at Chicago approved all of the procedures involving animals.

Tumor and Cell Protein Isolation and Western Blot

Tumors were ground in liquid nitrogen into a fine powder and resuspended in cell lysis buffer (20 mM Tris-HCl, pH 7.5; 150 mM NaCl; 1 mM Na₂EDTA; 1 mM EGTA; 1% Triton; 2.5 mM sodium pyrophosphate; 1 mM beta-glycerophosphate; 1 mM Na₃VO₄; 1 μ g/ml leupeptin; 1mM PMSF; 2% glycerol) and homogenized. The tumor lysate was centrifuged at 12,000 RPM for 15 min. at 4°C. Protein concentration was determined by the BCA method (BioRad Laboratories, Hercules, CA) and was stored at -80°C. Equal amounts of protein were run in a

standard western blot protocol using the following antibodies: Fas (B-10, Santa Cruz Biotech, Santa Cruz, CA), FasL (G2474, BD Pharmingen, San Jose, CA), phospho473-Akt and T-Akt (Cell Signaling Technology, Beverly, MA) ER α (G-20, Santa Cruz Biotech). All the antibodies were diluted in TBS-T (20mM Tris, 7.6; 137mM NaCl; 0.1% Tween20) containing 5% dry milk. Either the ECL Chemiluminescent Detection system (Amersham, Arlington Heights, IL) or the Supersignal West Dura western detection system (Pierce, Rockford, IL) was used to visualize the target band. Equal loading of total protein per lane was assessed by blotting with β -actin antibody (Sigma-Aldrich). Chemiluminescent signal was captured using a Chemi Doc™ Gel Documentation System (Bio-Rad Laboratories).

Assessment of Apoptosis

The identification of apoptotic cells *in situ* was determined using the ApopTag® TUNEL Apoptosis Detection kit (Chemicon International, Inc., Temecula, CA). This method detects DNA fragments by end-labeling 3'-hydroxyl DNA with digoxigenin using terminal deoxynucleotidyl transferase (TdT). An anti-digoxigenin antibody conjugated with peroxidase is bound and visualized with peroxidase substrate. Tissue sections of T47D:A18/PKC α tumors were 5- μ m thick and spread on silanized slides. Paraffin-embedded tissue sections were deparaffinized and progressively rehydrated. Sections were then pretreated with proteinase K from Sigma-Aldrich (20 μ g/ml for 20 min at room temperature). Endogenous peroxidase was blocked with 0.3% H₂O₂ in PBS. ApopTag® Plus Peroxidase In Situ Apoptosis Detection Kit (Chemicon) was then used according to the manufacturer's instructions. Staining of nuclei was expressed as the percentage of TUNEL-positive apoptotic cells in each tumor. TUNEL-positive cells were counted in 10 different high-power fields (X40) from 6 sections representing at least 3 independent tumors of each group. Repeated measures were obtained at 8, 12, and 15 days post-E2 capsule implantation.

Generation of T47D:A18/PKC α /Fas shRNA stable transfectant

T47D:A18/PKC α cells were transfected with MISSION™ shRNA lentiviral particles for Fas (Sigma-Aldrich, St. Louis, MO) following the manufacturer's instructions. T47D:A18/PKC α cells were suspended in culture medium (2 X 10⁴ cells/mL medium) and seeded in each well of a 12-well plate. After 30 hour incubation, media was removed and 1 mL fresh media with hexadimethrine bromide (8 μ g/mL) (Sigma-Aldrich) was added. Hexadimethrine bromide was used to enhance transduction according to the manufacturer's instructions. Lentiviral transduction particles for Fas were provided in a set of 4 clones. For each clone, 4 μ L of particles (4 X 10⁴ transducing units) were added into the wells. Media containing the lentiviral particles were removed and fresh media was added after 18 hr incubation at 37°C in a humidified atmosphere containing 5% CO₂. Drug selections with puromycin (InvivoGen, San Diego CA) at 1 μ g/mL and G418 (500 μ g/mL) were started on the following day. T47D:A18/PKC α cells were transfected with MISSION™ non-target shRNA control transduction particles (Sigma-Aldrich) and selected in the same way to serve as a negative control. Individual colonies were picked following three weeks of selection and screened for Fas expression by Western blot.

Real Time PCR analysis of estrogen responsive genes

For each treatment and control 3 independent tumor samples were available. RNA was extracted using the Rneasy Mini kit (Qiagen, Valencia, CA) and reverse transcribed with the Retroscript kit (Ambion). Taqman Gene Expression Assays were purchased from Applied Biosystems (Foster City, CA). The following assays were used: *HPRT1*, cat.# Hs99999909_m1, *C3*, cat.# Hs00163811_m1, *PGR*, cat.# Hs01556702_m1, *CTSD* cat.# Hs00157205_m1, *TFF1*, cat.# Hs00907239_m1, *TGFA*, cat.# Hs00608187_m1. PCR reactions were run in duplicate to check for reproducibility, and the Ct values were averaged. The PCR reaction was conducted in an Applied Biosystems 7900HT Real-Time PCR System.

The 20 μ l reaction mix consisted of 1X Taqman Universal PCR master mix, 1X gene expression assay and 10 ng cDNA template. PCR conditions were 50°C for 2 minutes and 95 °C for 10 minutes followed by 40 cycles of 95 °C for 15 seconds and 60 °C for 1 minute. For data analysis, the comparative Ct method was used, with *HPRT1* serving as housekeeping gene.

Transient transfection and luciferase assays

Prior to transient transfection, T47D:A18/neo and T47D:A18/PKC α cells were maintained for 3 days in phenol red free, E2-depleted RPMI-1640 media supplemented with G418. After stripping, cells were transiently transfected by electroporation as previously described, (37). Briefly, 8 X 10⁶ cells were harvested and resuspended in 0.5 mL of serum-free, phenol red free RPMI 1640 medium. ERE-tk-Luc plasmid containing the luciferase reporter gene controlled by the ERE (82) (5 μ g) and β -galactosidase (β -gal, 1 μ g) expressing plasmid pCMV β were added to the cell suspension and incubated for 5 min at room temperature. The cells were pulsed at 250 volts at 950 mF, resuspended in whole culture medium, and incubated at 37°C in a humidified CO₂ incubator overnight. On the following day, media containing E2 (10⁻⁹ M), E2-BSA conjugate (10⁻⁹ M) (Sigma-Aldrich) or vehicle (ethanol) were added.

Luciferase activities were measured using Luciferase Reporter Gene Assay System from Applied Biosystems (Bedford, MA). β -gal signals were measured with Galacto-Light Plus™ assay systems (Applied Biosystems, Bedford, MA). After 18 - 20 hours incubation, cells were washed with ice-cold PBS and lysed in the lysis buffer provided. The cell lysates were cleared by centrifugation and luciferase activity and β -gal signals were determined after adding corresponding substrates and read by a Monolight 2010 luminometer (Analytical Luminescence Laboratory, San Diego, CA). β -gal signals were applied to normalize the luciferase activity.

Statistical Analysis

Differences in mean tumor area between groups of three or more were measured using analysis of variance (ANOVA) test followed by Bonferroni Multiple Comparisons Test. Differences in three or more groups were measured using ANOVA followed by Tukey-Kramer Multiple Comparisons Post-Test. Unpaired Student's *t*-test was used to determine statistical significance between two groups and was used to analyze data obtained from the colony formation assays, TUNEL stain, and western blots. The GraphPad InStat Version 3.06 statistical software package was used (GraphPad Software, San Diego, CA) and SPSS 13, SPSS, Inc.(Chicago, IL). All statistical tests were two-sided.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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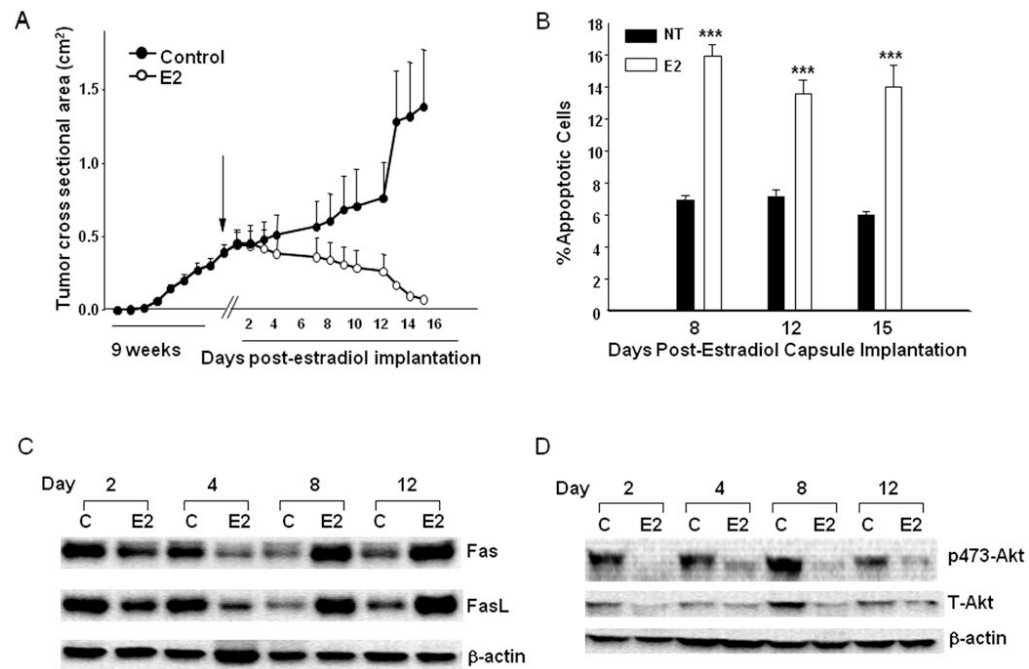


Figure 1. E2 induces tumor regression, apoptosis, upregulation of Fas/FasL and downregulation of p-Akt/T-Akt in T47D:A18/PKC α tumors

A.) Tumor growth of T47D:A18/PKC α *in vivo*. T47D:A18/PKC α cells were bilaterally injected into the mammary fat pads of thirty athymic mice. Mice were left untreated for 9 weeks until the mean tumor cross-sectional area reached 0.5 cm², and then randomized to two treatment groups: continued no treatment (Control, 10 mice) and 1.0 cm E2 capsule (Control + E2, 20 mice). Results are shown as mean tumor cross-sectional area \pm SE (upper limit only is shown for each point). Tumors were collected from both the Control and Control + E2 groups on 2, 4, 8, 12 and 15 days post-E2 capsule implantation. **B.)** Apoptotic effect of E2 on T47D:A18/PKC α tumors. Apoptosis in tumors was assessed by the TUNEL method from both Control and Control + E2 groups excised 8, 12 and 15 days post-E2 capsule implantation (corresponds to data shown in panel A). Data represent the percentage apoptotic cells in each treatment groups. Error bars represent SE. *** $P < .001$ compared with Control treatment group. Statistical test was two-sided. **C.)** Western blot analysis of Fas ligand (Fas L) and Fas protein expression in T47D:A18/PKC α tumors. **D.)** Western blot of T-Akt and P473-Akt protein expression in T47D:A18/PKC α tumors. Tumors were excised from both Control and Control + E2 groups on days 2, 4, 8 and 12 post-E2 capsule implantation. β -Actin was used as a loading control. Western blots are representative of two independent tumors, with three replicate experiments, all with similar results.

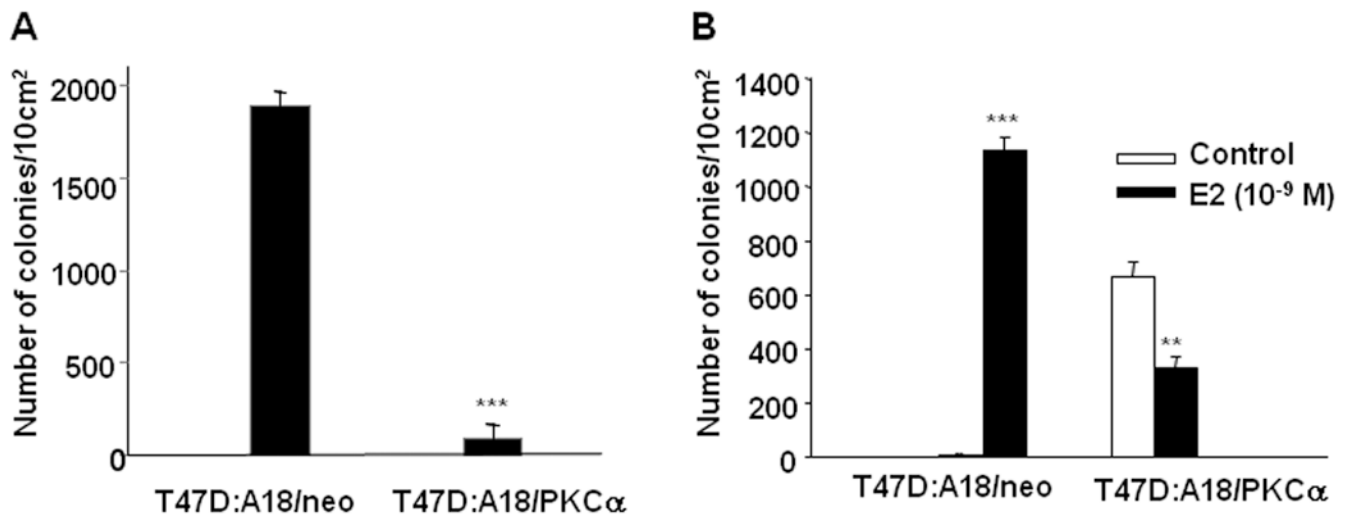


Figure 2. Growth inhibitory effects of E2 in T47D:A18/PKC α cells grown in Matrigel
 Cells were plated in Matrigel as described in “Materials and Methods”. **A.**) T47D:A18/neo and T47D:A18/PKC α cells were grown in phenol red-containing RPMI and seeded into phenol red-containing Matrigel. Data are the mean number of colonies per 10 cm², error bars represent the standard error (SE). ***Comparison of the number of colonies between each cell line is statistically different at $P < 0.001$. **B.**) T47D:A18/neo and T47D:A18/PKC α cells were grown in phenol red-free RPMI and seeded into phenol red-free Matrigel. Cells were either not treated (Control) or treated with E2 (10⁻⁹ M). ***Comparison of the number of T47D:A18/neo colonies in the Control versus E2-treated cells was statistically different at $P < 0.001$. *Comparison of the number of T47D:A18/PKC α colonies in the Control versus E2-treated cells was statistically different at $P < 0.01$.

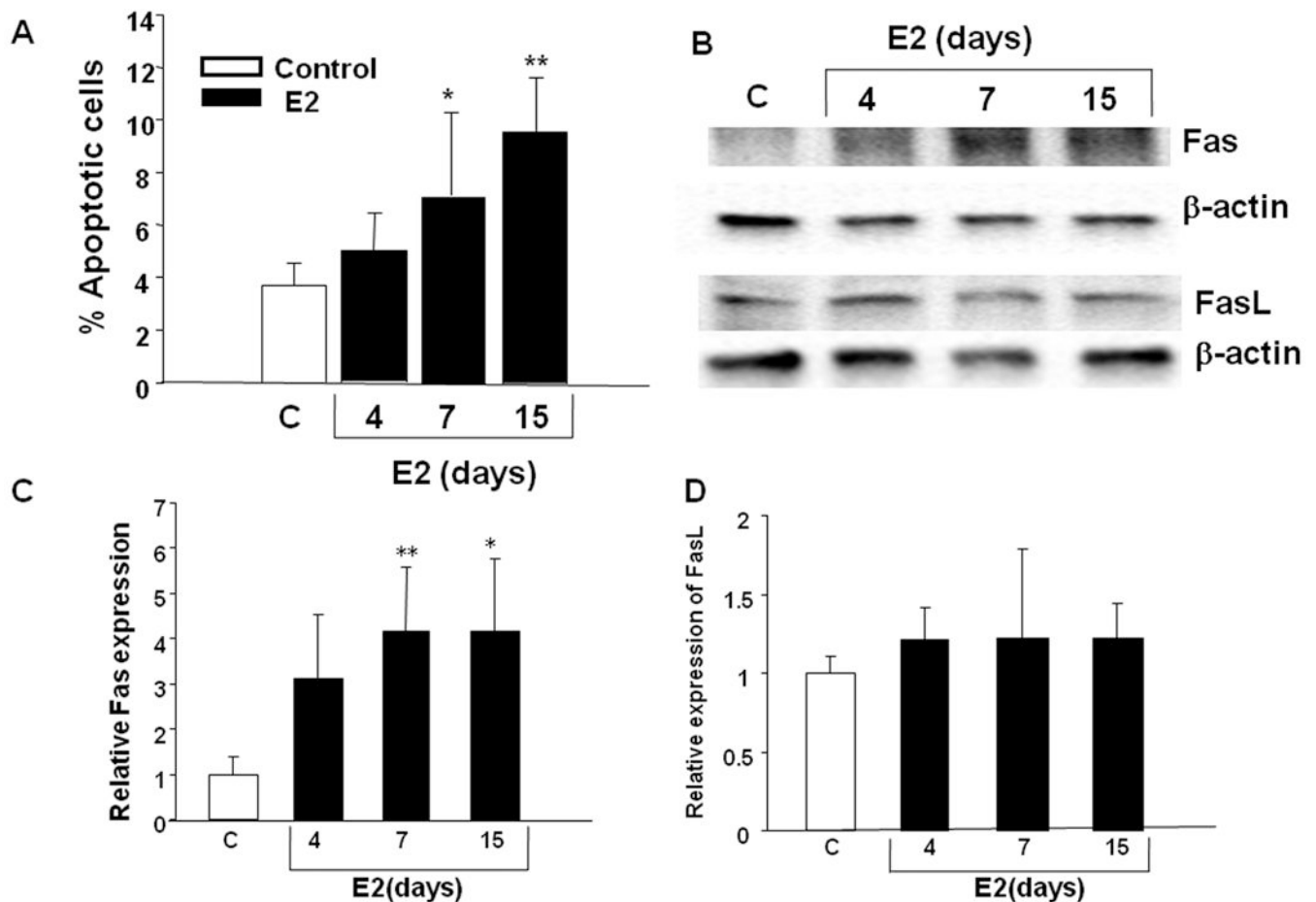


Figure 3. E2 induces apoptosis and expression of Fas in T47D:A18/PKC α cells grown in Matrigel
A.) T47D:A18/PKC α cells were grown in phenol red-free media, seeded on top of phenol red-free Matrigel and were maintained for 15 days. E2 treatment (10^{-9} M) was initiated at various times and on Day 15, colonies were released from Matrigel, fixed, sectioned and TUNEL stain was performed. Apoptosis was significantly increased in a time-dependent manner in E2-treated colonies compared to untreated colonies. *: Comparison of apoptosis in colonies treated with E2 for 7 days versus control (untreated) colonies was significantly different at $P < 0.05$; **comparison of apoptosis following E2 treatment for 15 days versus control, $P < 0.01$. The results are representative of two independent experiments. **B.)** T47D:A18/PKC α cells were grown in phenol red-free Matrigel and either left untreated (control) or treated with E2 (10^{-9} M) for 4, 7 or 15 days. Western blot showing Fas and FasL protein expression is representative of three independent experiments. **C, D.)** Quantitation of the bands in panel B using Quantity One software (Bio-Rad Laboratories.). *Fas expression is significantly different at $P < 0.05$ compared to control; **Fas expression is significantly different at $P < 0.01$ compared to control.

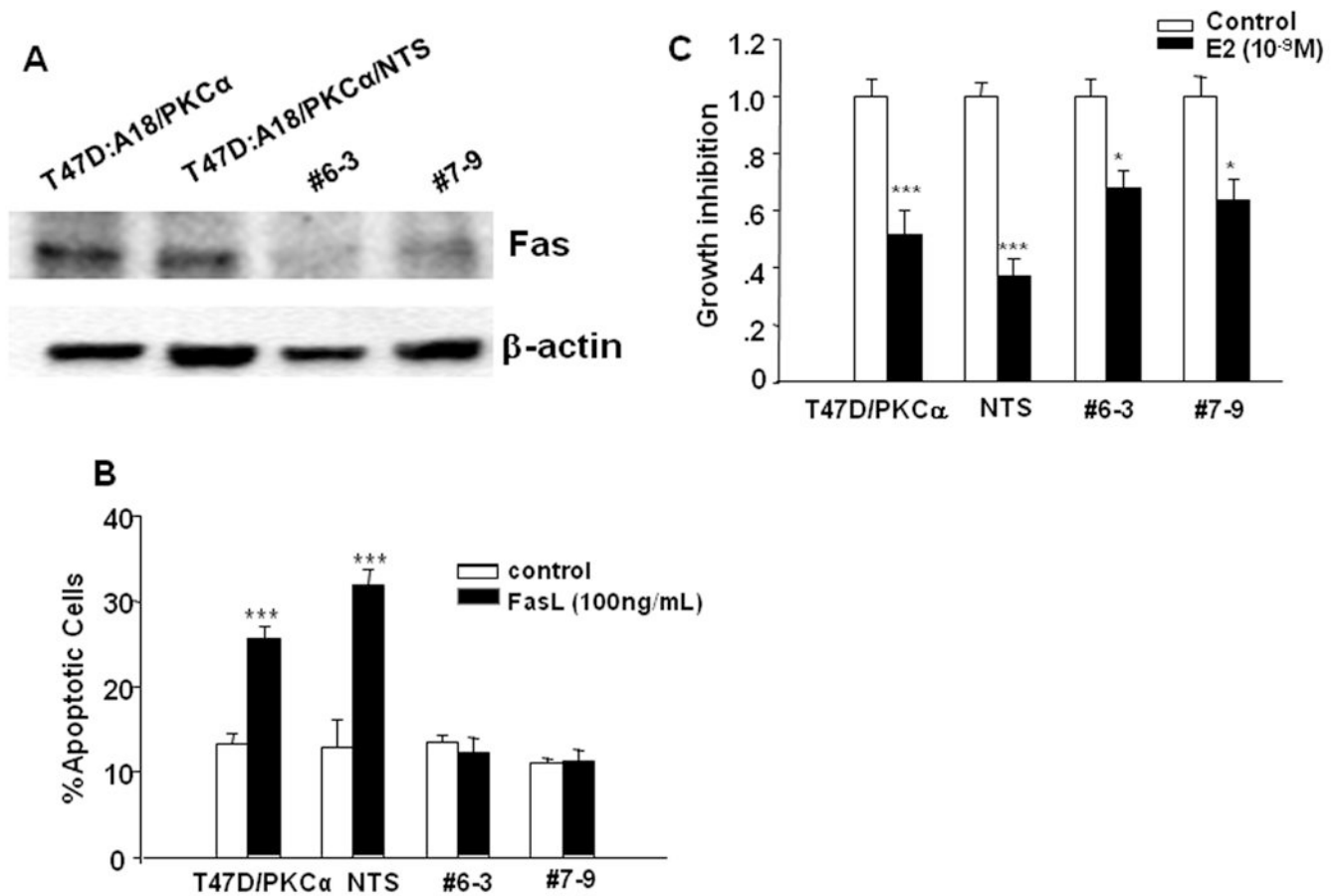


Figure 4. Knock-down of Fas by shRNA transfection reduces the E2-inhibitory effects

A.) T47D:A18/PKC α cells were transfected with lentiviral shFas transduction particles, and stable clones were selected and screened for Fas expression with western blot. NTS: T47D:A18/PKC α cells stably transfected with non-targeting sequences. **B.)** T47D:A18/PKC α , T47D:A18/PKC α /NTS, T47D:A18/PKC α /shFas#6-3 and #7-9 clones were treated with recombinant FasL peptide for 16 hours and apoptosis was detected with Annexin-V/PI double staining followed by flow cytometry. ***: Comparison of % apoptosis in FasL treated cells versus untreated (control) cells was significantly different at $P < 0.001$. **C.)** T47D:A18/PKC α , T47D:A18/PKC α /NTS, T47D:A18/PKC α /shFas#6-3 and #7-9 were grown in phenol-free Matrigel and treated with E2 (10^{-9} M) or vehicle for 20 days. The number of untreated colonies (control group) was set as one, and the growth inhibition was represented by the ratio of the number of E2-treated colonies to control group. *** and *: Comparison of the number of E2-treated colonies versus the untreated control group was significantly different at $P < 0.001$ and $P < 0.05$ correspondingly using t-test.

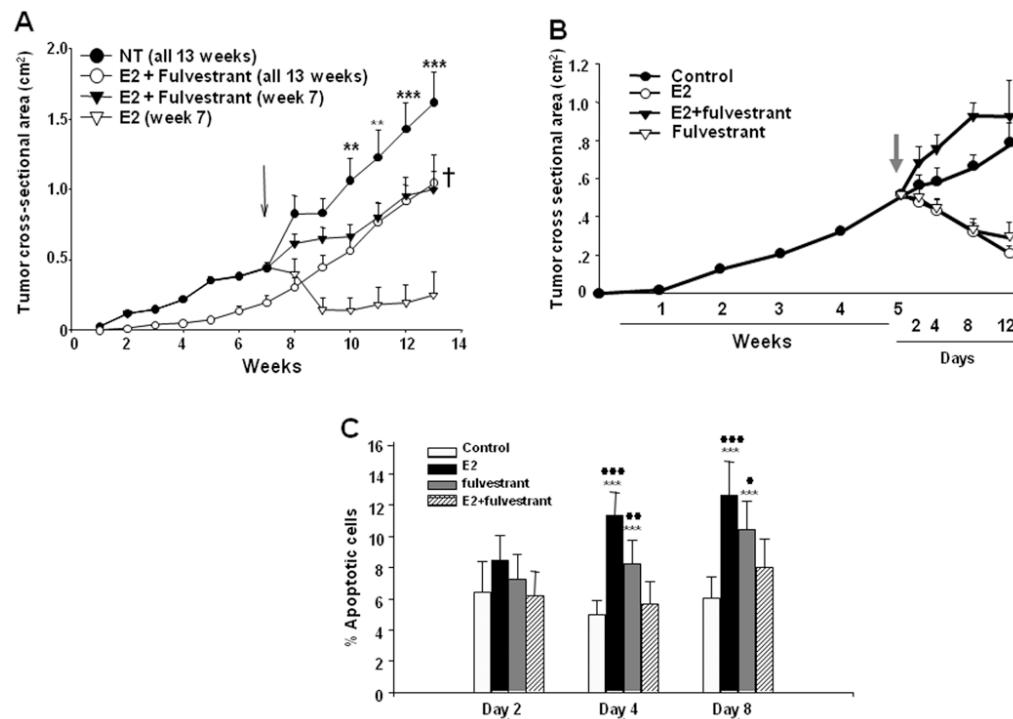


Figure 5. The presence of fulvestrant abolished E2-induced T47D:A18/PKC α tumor regression and apoptosis in athymic mice

A.) Cells were injected into the mammary fat pads of fifty athymic mice. Forty mice were not treated (NT group), 10 mice were implanted with a 1.0 cm E2 capsule and given weekly injections with fulvestrant (E2 + fulvestrant, 13 weeks). After 7 weeks (indicated by arrow), forty mice from the NT group were randomized to three treatment groups: continued NT (10 mice), E2 capsule (E2, week 7, 10 mice), E2 capsule and weekly injections with fulvestrant (E2 + fulvestrant, week 7, 20 mice). Results are shown as mean tumor size \pm SE (upper limit only is shown for each point). Tumors in the NT group were statistically significantly larger than tumors in the E2 treatment group at weeks 11-13. ** $P < 0.01$; *** $P < 0.001$. †Tumors in the E2 + fulvestrant, week 7 and E2 + fulvestrant, 13 weeks groups were statistically significantly larger than tumors in the E2 group at week 13 ($P < 0.01$) using the Bonferonni multiple comparisons test. **B.)** Cells were injected into the mammary fat pads of forty athymic mice. The mice were not treated and tumors allowed to grow until week 5. After 5 weeks (indicated by arrow), the mice were randomized to four treatment groups with 10 mice in each group: continued untreated (control), E2 1.0 cm capsule (E2), E2 1.0 cm capsule and weekly injections with fulvestrant (E2+fulvestrant), and fulvestrant injections alone (fulvestrant). Results are shown as mean tumor size \pm SE. **C.)** Analysis of apoptosis in T47D:A18/PKC α tumors treated with E2, fulvestrant, or E2 plus fulvestrant. TUNEL assays were performed on the T47D:A18/PKC α tumors in control, E2, fulvestrant or E2+fulvestrant treatment groups to detect apoptotic cells. The data was expressed as mean \pm SE based on measurements of 3 independent tumors. **: a significant difference between control and treatment groups with $P < 0.01$. **: $P < 0.001$. #: significant difference between treatments and E2+fulvestrant group with $P < 0.05$ (##: $P < 0.01$; ###: $P < 0.001$) (Tukey-Kramer Multiple Comparison post-hoc test).

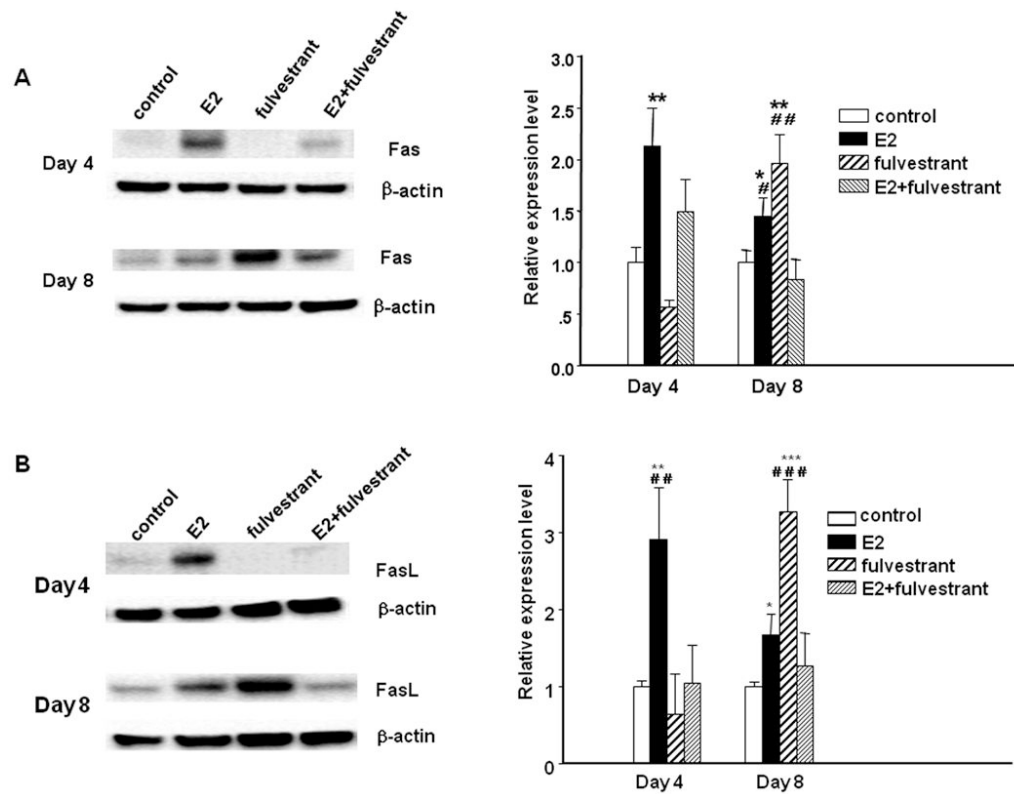


Figure 6. Western blot analysis of Fas and FasL protein expression in T47D:A18/PKC α tumors and cells

T47D:A18/PKC α tumors were excised from Control, E2, fulvestrant and E2 plus fulvestrant groups on day 4 and day 8 post-E2 capsule implantation (corresponds to Figure 5B) and protein was extracted. Western blot indicates Fas (A) and FasL (B) protein expression and is representative of four independent tumors, all with similar results. β -Actin was used as a loading control. *: $P < 0.05$; **: $P < 0.01$; ***: $P < 0.001$, significant difference between treatment and control using one-way ANOVA followed by Tukey-Kramer Multiple Comparison post-hoc test. #: significant difference when compared to E2 plus fulvestrant treatment group.

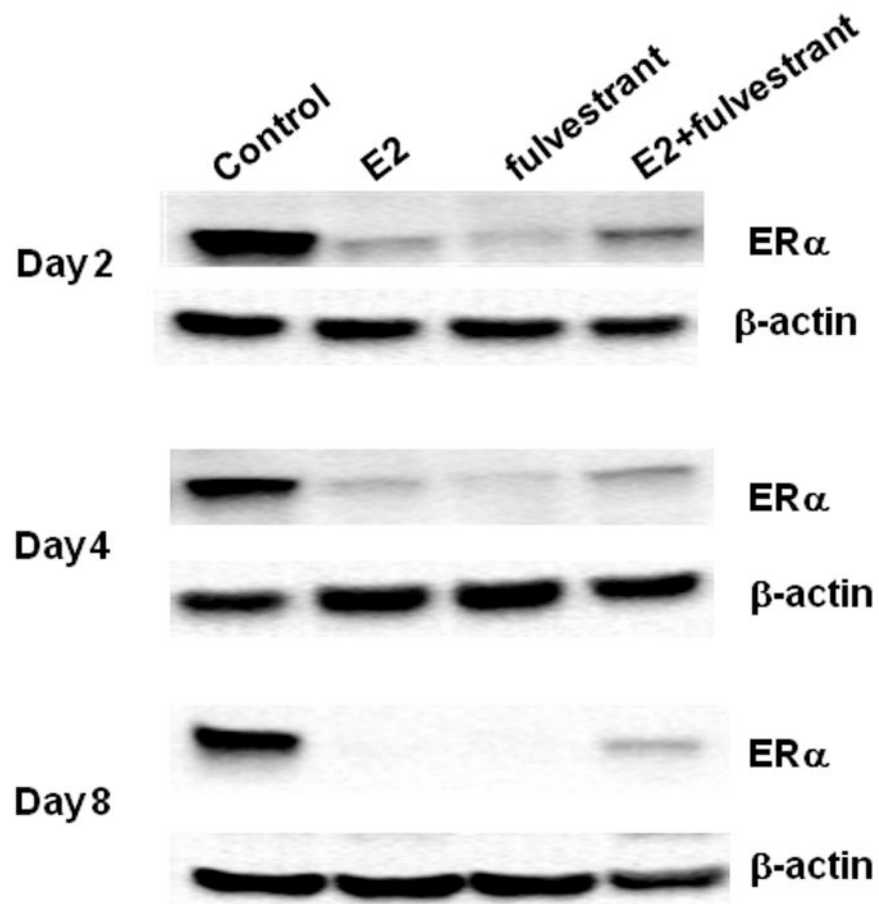


Figure 7. Western blot analysis of ER protein expression in T47D:A18/PKC α tumors
T47D:A18/PKC α tumors were excised from Control, E2, fulvestrant and E2 plus fulvestrant groups on days 2, 4 and 8 post-E2 capsule implantation (corresponds to Figure 5B) and protein was extracted. Western blot indicates ER protein expression and is representative of three independent tumors, all with similar results. β -Actin was used as a loading control.

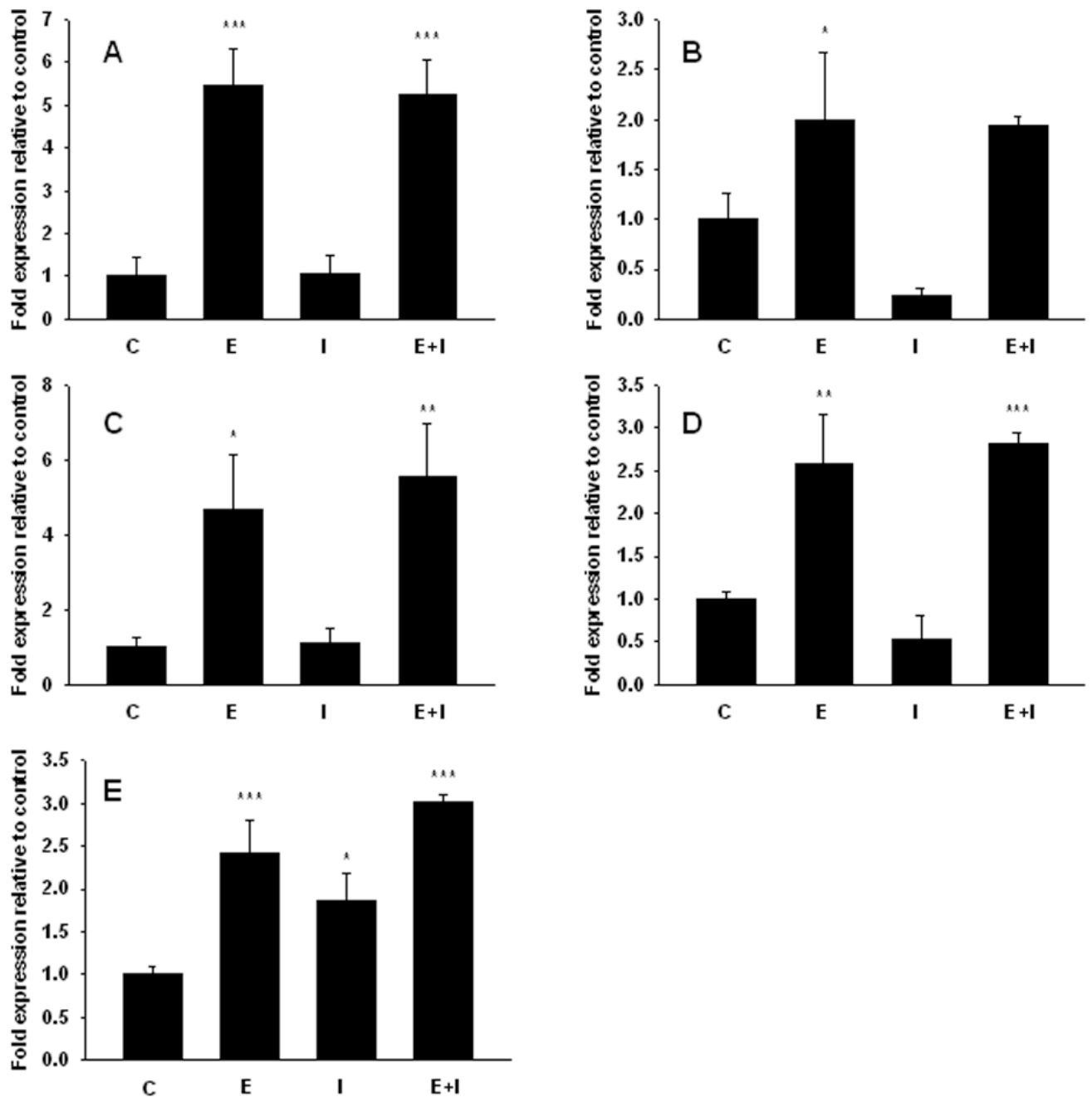


Figure 8. Regulation of ER-responsive gene transcription

mRNA levels of estrogen responsive genes in PKC α overexpressing tumors 8 days following treatment. **A)** *C3*. **B)** *PGR*. **C)** *CTSD*. **D)** *TFF1*. **E)** *TGFA*. Each group contains 3 independent tumors. Expression levels were compared using Tukey's HSD test followed by one-way ANOVA. Stars indicate groups significantly different from control. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$. C, control, E, 17 β -estradiol, I, fulvestrant, E+I, 17 β -estradiol+fulvestrant.

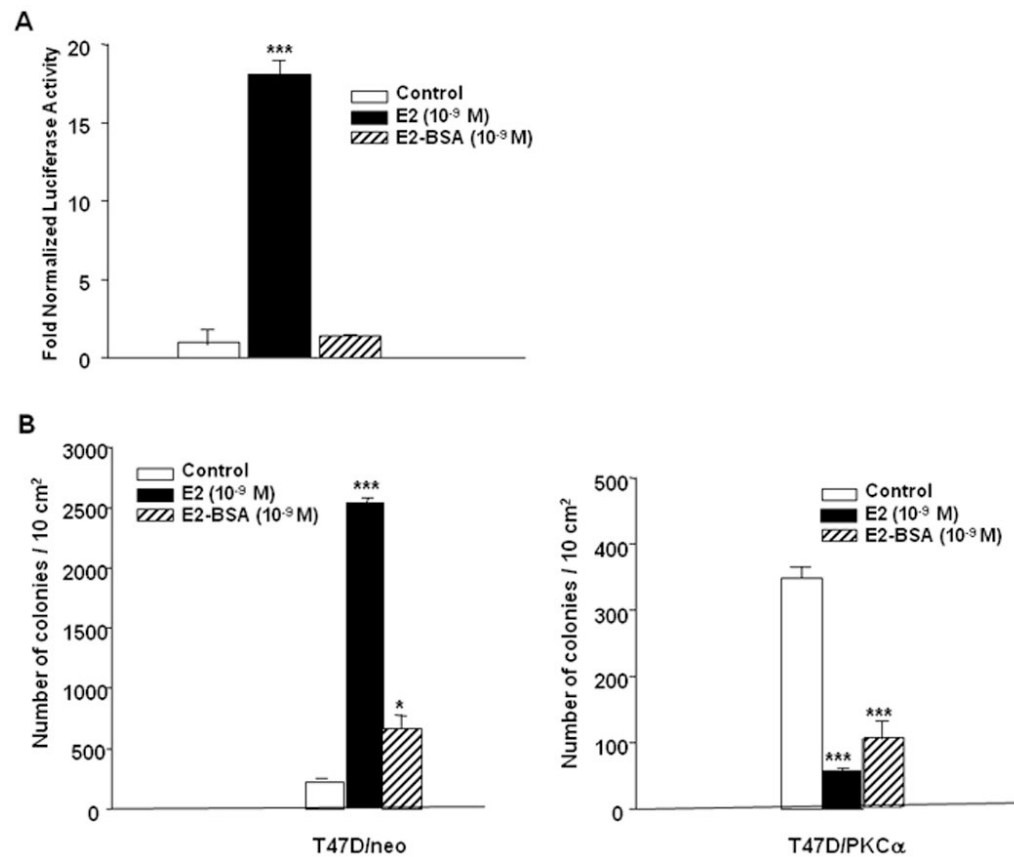


Figure 9. Both E2 and E2-BSA inhibit T47D:A18/PKCa colony formation in Matrigel

A.) E2-BSA does not induce ERE-luciferase activity in T47D:A18/neo cells. T47D:A18/neo cells were transfected with ERE-tk-LUC and β -galactosidase expression plasmids. Luciferase activity was measured after 20 hours treatment with E2 (10⁻⁹ M), E2-BSA (10⁻⁹ M) or vehicle (control) and normalized to β -gal. **B.**) Colony formation assays. T47D:A18/neo and T47D:A18/PKCa cells were grown in phenol-red free Matrigel in the presence of vehicle (Control), E2 (10⁻⁹ M), or E2-BSA (10⁻⁹ M). *: significant difference between treatment and control groups at $P < 0.05$; ***: $P < 0.001$ (one-way ANOVA Tukey-Kramer Multiple Comparison post-hoc test).