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To cite this article: Amy E. Siebert, Amelita L. Sanchez, Sumi Dinda & Virinder K. Moudgil (2011) Effects of Estrogen Metabolite 2-Methoxyestradiol on Tumor Suppressor Protein p53 and Proliferation of Breast Cancer Cells, *Systems Biology in Reproductive Medicine*, 57:6, 279-287, DOI: [10.3109/19396368.2011.633152](https://doi.org/10.3109/19396368.2011.633152)

To link to this article: <https://doi.org/10.3109/19396368.2011.633152>



Published online: 14 Nov 2011.



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INVITED RESEARCH ARTICLE FROM THE 2011 MICHIGAN ALLIANCE FOR REPRODUCTIVE TECHNOLOGIES AND SCIENCES SYMPOSIUM, CURRENT TRENDS IN DEVELOPMENTAL AND REPRODUCTIVE TOXICOLOGY

Effects of Estrogen Metabolite 2-Methoxyestradiol on Tumor Suppressor Protein p53 and Proliferation of Breast Cancer Cells

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An endogenous 17 β -estradiol (E₂) metabolite, 2-methoxyestradiol (2-ME₂), has been reported to exhibit estrogen receptor (ER)-independent anti-angiogenic and anti-tumor effects. Several mechanisms have been proposed for 2-ME₂ actions, but there is a lack of evidence for a common pathway for all of the cell-types sensitive to this metabolite. We have examined potential alterations in p53 in response to 2-ME₂, E₂ and the microtubule disruptor taxol in T47D breast cancer cells. Cells were cultured for six days in medium depleted of endogenous steroids or effectors. Semi-confluent cells were treated with 2-ME₂ (1 nM – 10 μ M), 10 nM E₂ and/or 1 μ M taxol and subjected to SDS-PAGE and Western blot analysis, quantitative analysis, or laser-scanning confocal microscopy. Western blot analysis revealed a concentration-dependent biphasic trend in p53 levels. Addition of 10 nM – 1 μ M 2-ME₂ induced significant up-regulation in p53, and this response gradually diminished to levels comparable to the control upon treatment with higher concentrations (2.5 – 10 μ M). The observed upregulation of p53 induced by 2-ME₂ is inhibited by concurrent treatment with 1 μ M taxol. Cell quantitation revealed a significant decrease (50 – 90%) in cell number upon treatment with 1 – 10 μ M 2-ME₂ with minimal effect at lower concentrations. No additional effect on cell proliferation was observed when taxol was combined with 10 nM or 1 μ M 2-ME₂. In a concentration dependent manner, treatment with 2-ME₂ for 24 h differentially influenced cellular localization of p53. These results may aid in further understanding the relationship between steroid receptors, tumor suppressor proteins, and effects of hormone metabolites on breast cancer cells.

Keywords 2-Methoxyestradiol, breast cancer, estrogen, p53, tumor suppressor protein

Abbreviations 2-ME₂: 2-methoxyestradiol; ATCC: American type culture collection; COMT: catechol-o-methyltransferase; Cs: control cells cultured in 5% fetal bovine serum (FBS) treated with Dextran-coated charcoal (DCC); Cw: control cells cultured

in 10% FBS; DAPI: 4',6-diamidino-2-phenylindole; E₂: 17 β -estradiol; ECL: enhanced chemiluminescence; ER: estrogen receptor; HBSS: Hanks' balanced salt solution; p53: 53-kDa tumor suppressor protein; PBS: phosphate buffered saline; PMSF: phenylmethylsulfonyl fluoride; PR: progesterone receptor; PVDF: polyvinylidene fluoride; RIPA: radio-immunoprecipitation assay; SDS-PAGE: sodium dodecyl sulfate polyacrylamide gel electrophoresis; TBS: tris-buffered saline.

Introduction

Estrogen exerts multiple effects on physiological functions in reproductive and other tissues [Agrawal and Alvin Jose 2010; Depalo et al. 2009]. It is widely known that 17 β -estradiol (E₂) is a mitogenic steroid that enhances proliferation of target cells. Several E₂ metabolites have been shown to be more potent estrogenic compounds than their precursor [Nandi et al. 1995; Seegers et al. 1989]. However, unlike these growth-enhancing estrogen metabolites, 2-methoxyestradiol (2-ME₂) is a highly potent anti-proliferative metabolite that induces apoptosis in tumor cells *in vitro* as well as *in vivo* [Pribluda et al. 2000]. The formation of 2-ME₂ occurs from the catechol-o-methyltransferase (COMT)-mediated o-methylation of the catecholesterogen 2-hydroxyestradiol, a major metabolite formed in humans by the hepatic hydroxylation of E₂ [Mannisto and Kaakkola 1999]. In addition to hydroxylation in the liver, 2-hydroxyestradiol is also formed in the uterus and breast, where it can then be further metabolized by COMT to form 2-ME₂ or to other products by conjugation enzymes [Mannisto and Kaakkola 1999].

Studies have shown that 2-ME₂ is a potent inhibitor of proliferation of transformed and endothelial cells [Fotsis et al. 1994; Lottering et al. 1992], as well as angiogenesis *in vivo* [Fotsis et al. 1994]. There are indications that the anti-proliferative effects of 2-ME₂ are due to the targeted disruption of cellular events associated with proliferation

Received 10 July 2011; accepted 29 September 2011.

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[Fotsis et al. 1994; Mueck and Seeger 2010]. The metabolite has been shown to affect cell cycle events in S-phase, arrest mitotic cells during metaphase by disrupting correct microtubule assembly, and induce cellular alterations that are indicative of apoptosis [Attalla et al. 1996; Cushman et al. 1995; D'Amato et al. 1994; Hamel et al. 1996; Lottering et al. 1992; Mueck and Seeger 2010; Seegers et al. 1989]. These observations suggest that 2-ME₂ is selective for only actively proliferating cells [Fotsis et al. 1994; Mueck and Seeger 2010].

Though the anti-proliferative and anti-angiogenic effects of 2-ME₂ are well established [Cushman et al. 1995; D'Amato et al. 1994; Fotsis et al. 1994; LaVallee et al. 2002; Lottering et al. 1992; Pribluda et al. 2000; Seegers et al. 1989], the mechanism(s) by which this molecule elicits its effects is poorly understood. Several mechanisms have been proposed for 2-ME₂ actions, but there is a lack of evidence for a common pathway for all of the cell-types sensitive to this metabolite. One of the proposed mechanisms is the up-regulation of the tumor suppressor protein p53 [Mukhopadhyay and Roth 1997; Seegers et al. 1997; Shimada et al. 2003]. The p53 phosphoprotein is a nuclear transcription factor that functions, in part, by responding to DNA damage and inducing apoptosis [Riley et al. 2008]. The level of p53 protein is indicative of the cellular response to stressors [Vogelstein et al. 2000]. The transcriptional and translational activation of the p53 protein after exposure to genotoxic and nongenotoxic stressors induces the expression of proteins involved in redox regulation, cell-cycle arrest, DNA repair, protein degradation, and apoptosis [El-Deiry et al. 1993; Erster et al. 2004; Grover et al. 2009; Halaby and Yang 2007]. The p53 protein's apoptotic function is primarily, but not exclusively, mediated via activation of mitochondrial apoptotic machinery through the intrinsic pathway [Erster et al. 2004; Grover et al. 2009; Hamel et al. 1996; Miyashita and Reed 1995].

Mutations of the p53 gene contribute to approximately fifty percent of all human cancers and are seen in a wide variety of tumors such as breast, lung, esophageal, liver, bladder, and ovarian carcinomas, as well as brain tumors, sarcomas, lymphomas, and leukemias [Hollstein et al. 1991]. Although the exact mechanism by which normal tissue transforms to a malignant state is unclear, the primary events associated with carcinogenesis may involve mutations in the p53 gene [Moudgil et al. 2001]. Earlier studies have shown abnormally high levels of the p53 protein in human tumor cell lines and tissues [Donehower and Bradley 1993]. These results suggest an alteration in regulation or structure of the p53 gene, due to the known increase in cellular stability of the mutant protein product compared to the normal wild type (wt) protein [Donehower and Bradley 1993].

Previously, Lottering et al. [1992] showed that estrogen receptor (ER)-positive MCF-7 breast cancer cells, expressing wt p53, were susceptible to the anti-mitotic effects of 2-ME₂. Despite being a natural metabolite of E₂, the antiproliferative and cytotoxic effects induced by 2-ME₂ are independent of ER status and are not considered to be ER mediated [Cushman et al. 1995; LaVallee et al. 2002]. There is a

paucity of information regarding the influence of 2-ME₂ on tumor suppressor function in hormone-responsive breast cancer cells and thus further studies are warranted. Our laboratory initially reported the hormonal regulation of the p53 tumor suppressor protein in T47D breast cancer cells [Hurd et al. 1995; Moudgil et al. 2001]. The T47D cell line is estrogen receptor (ER) and progesterone receptor (PR)-positive and expresses p53 protein with a missense mutation at codon 194 converting leucine to phenylalanine [Nigro et al. 1989]. In the present study, the hormone dependent T47D breast cancer cells were utilized to examine the effects of 2-ME₂ on cell proliferation and on the expression and localization of p53 protein. Studies were also conducted to compare the observed effects of 2-ME₂ with those of exogenous E₂ as well as the extensively studied microtubule-stabilizing agent taxol. The physiological effects of E₂ are mediated via ER, and microtubules are the only known biochemical target of taxol. Comparison of the observed effects of these compounds may aid in elucidating mechanisms or potential pathways involved in the actions of this metabolite. Understanding the mechanism of p53 accumulation may lead to further examination of the potential of 2-ME₂ in new therapeutic modalities.

Results

Effects of 2-ME₂ on p53 Protein: Concentration Dependency

Our laboratory has previously shown that factors present in growth serum appear to be required for maintaining high p53 levels and that depletion of factor(s) by charcoal treatment decreases p53 to basal levels in T47D cells [Moudgil et al. 2001]. Results in Figure 1 show that when T47D cells were cultured in 5% fetal bovine serum (FBS) stripped with Dextran-coated charcoal (DCC-FBS) for six days (labeled as Cs), the level of p53 was approximately one-half of that observed when cells were grown in 10% FBS (labeled as Cw) for the total duration of the study. We have monitored the alteration in p53 protein level in cells grown for six days in 5% DCC-FBS to allow for a more direct assessment of the effects of 2-ME₂ in these studies. When T47D cells were treated with 1 nM – 10 μM 2-ME₂ for 24 h, the E₂ metabolite caused a consistent biphasic trend in the levels of p53. As compared to Cs, a significant increase in the p53 level was observed upon addition of 10 nM – 1 μM 2-ME₂. The observed increase in p53 level was not selective to 2-ME₂, as E₂ has been shown to reverse the effect of charcoal treatment on p53 levels [Hurd et al. 1995; Moudgil et al. 2001]. This response is reflected in results seen in Figure 1 showing a 70% increase in the level of p53 upon addition of 10 nM E₂. Interestingly, higher concentrations of 2-ME₂ (2.5–10 μM) appear to be less effective in altering observed p53 levels.

Effects of 2-ME₂ on T47D Cell Proliferation: Concentration Dependency

In order to determine a physiological function of 2-ME₂, quantitative analysis of the effects of this metabolite on cell proliferation was performed. Results presented in Figure 2

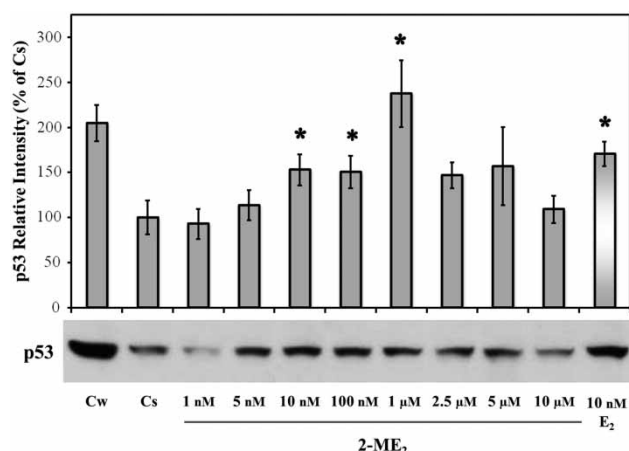


Figure 1. Concentration-dependent effects of 2-ME₂ on p53. T47D cells were plated for two d in medium (- phenol red) containing 10% fetal bovine serum (FBS) followed by culturing for six d in medium containing 5% DCC-FBS. Semi-confluent cells were treated with 1 nM – 10 μM 2-ME₂ or 10 nM E₂ for 24 h. Cellular extracts were prepared and subjected to protein quantitation, SDS-PAGE, and Western analyses as described in Materials and Methods. Lanes labeled Cw and Cs represent control lanes with cells grown in the absence of ligands in medium containing either 10% FBS or DCC-FBS, respectively. The relative intensities of p53 as compared to Cs are displayed as the mean ± SEM. The sample sizes ranged from 3 - 15 integrated density measurements per group. *indicates significant difference with the control at $p < 0.05$ (Kruskal-Wallis Test followed by post-hoc analysis using Mann-Whitney U-Test). Representative Western blot is shown.

show the growth response of T47D cells cultured in the presence of 1 nM – 10 μM 2-ME₂ or 10 nM E₂. Although the use of physiological steroid hormones in experimental studies employs steroid concentrations in the nanomolar range, the metabolite 2-ME₂ is effective in cell free systems at higher

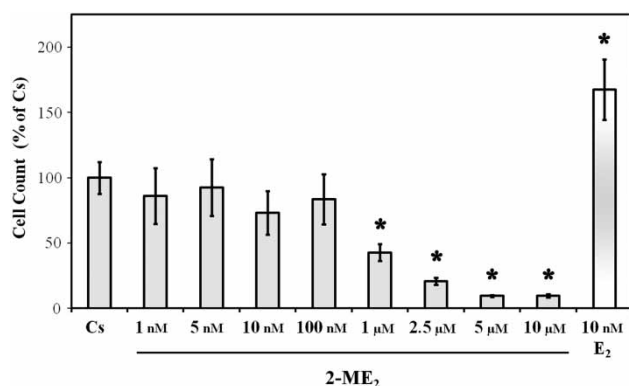


Figure 2. Effects of 2-ME₂ on T47D cell proliferation. T47D cells were seeded at 15,000 cells/well in 12 well growth plates for two d in medium (- phenol red) containing 10% FBS followed by culturing for six d in medium containing 5% DCC-FBS supplemented with 1 nM – 10 μM 2-ME₂ or 10 nM E₂. Control and treatment medium was changed every 48 h. Cells were trypsinized and counted by Coulter Counter (model # Z2) after six d of treatment. Mean cell populations ($\bar{x} \pm$ SEM) were plotted as a percentage of Cs with 100% on the ordinate scale representing cells cultured in the absence of ligands ($\bar{x} = 2.1 \times 10^5$ cells/well). The sample sizes ranged from 8-20 cell counts per group. *indicates significant difference with the control at $p < 0.05$ (Kruskal-Wallis Test followed by post-hoc analysis using Mann-Whitney U-Test).

concentrations. Previous studies have shown inhibition of cellular proliferation by use of 1 – 10 μM 2-ME₂ [Azab et al. 2009; LaVallee et al. 2002; Mueck and Seeger 2010; Pribluda et al. 2000; Van Zijl et al. 2008]. Thus the use of 2-ME₂ at the concentrations utilized in our studies is consistent with similar concentrations employed by other investigators in the literature. T47D cells were cultured for two days in 10% FBS followed by culturing for six days in 5% DCC-FBS growth medium containing the exogenous ligands as described in Materials and Methods. Culturing T47D cells in 5% DCC-FBS greatly reduces their growth rate suggesting that charcoal treatment removes factors that are required for maximal proliferation of this cell line [Hurd et al. 1995; Kodali et al. 1994]. The addition of 10 nM E₂, induced a significant proliferative response in cells cultured in the absence of serum growth factors and reflects findings reported previously by this laboratory [Hurd et al. 1995]. When T47D cells were cultured in the presence of 1 – 100 nM 2-ME₂, no significant effect on the cell population was observed. However, treatment with 1 – 10 μM 2-ME₂ caused significant growth inhibition (50 – 90%) in this cell line.

Effects of 2-ME₂, E₂ and Taxol on p53

The use of microtubule-disrupting agents such as taxol in the treatment of neoplastic disease has long been established. Previous studies from our laboratory have demonstrated that 1 μM taxol was equally effective in blocking T47D cell proliferation when cultured in 5% FBS or DCC-FBS, indicating that taxol effects are not dependent on charcoal-sensitive heterocyclic compounds [Kodali et al. 1994]. In order to gain insight into possible similarities between 2-ME₂ and other microtubule-disrupting agents, T47D cells cultured in 5% DCC-FBS were treated with 10 nM and 1 μM 2-ME₂ and 1 μM taxol alone or in combination for 24 h (Fig. 3). For

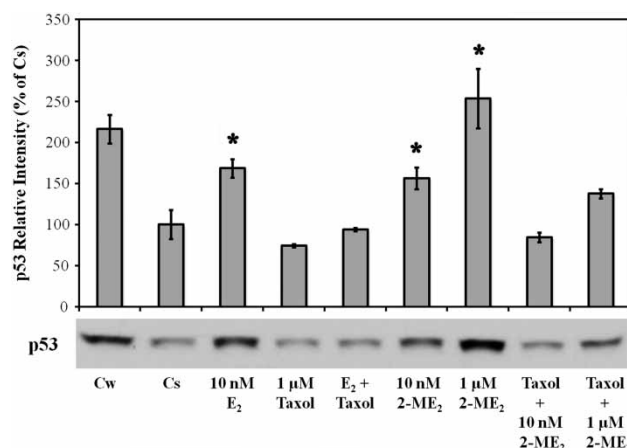


Figure 3. The effects of 2-ME₂, E₂, and Taxol on p53 protein. Culture and treatment of T47D cells with 2-ME₂, E₂, and taxol was performed as described in Figure 1. Cellular extracts were prepared and subjected to protein quantitation, SDS-PAGE and Western analyses. The relative intensities of p53 as compared to control (Cs) are displayed as the mean ± SEM. The sample sizes ranged from 3 - 15 density measurements per group. *indicates significant difference with the control at $p < 0.05$ (Kruskal-Wallis Test followed by post-hoc analysis using Mann-Whitney U-Test). Representative Western blot is shown.

comparative purposes, 10 nM E₂ alone and in combination with 1 μ M taxol was also included in this study. Treatment with 1 μ M taxol alone did not significantly affect p53 protein level. However, taxol was able to inhibit up-regulation of p53 induced by 10 nM E₂. Interestingly, 1 μ M taxol also had an inhibitory effect on the ability of 2-ME₂ (10 nM and 1 μ M) to increase p53 level.

Effects of 2-ME₂, E₂, and Taxol on T47D Cell Proliferation

To determine whether an additional benefit exists when 2-ME₂ is combined with taxol, a quantitative analysis of cell growth was performed. T47D cells were cultured for six days in 5% DCC-FBS and in the presence of 10 nM and 1 μ M 2-ME₂, 10 nM E₂, or 1 μ M taxol, either alone or in combination. Addition of 1 μ M taxol caused a decrease in T47D cell number (Fig. 4), which was consistent with previously published results from our laboratory [Kodali et al. 1994]. When used in combination with E₂, taxol significantly reversed the proliferative effects of E₂, causing an 85% inhibition in cell growth as compared to the control. No additional effects were observed upon treatment of cells with 2-ME₂ and taxol in combination.

Immunolocalization of p53 in T47D Cells: the Effects of 2-ME₂, E₂, and Taxol

To determine if the effects of 2-ME₂ on the level of p53 correlate with alterations in the cellular localization of the tumor suppressor, immunolabeling of p53 protein in T47D cells was performed followed by laser-scanning confocal microscopy. Consistent with the transcriptional function of this nuclear phosphoprotein, results in Figure 5 reveal p53 localized in the nuclei of T47D cells cultured in 5%

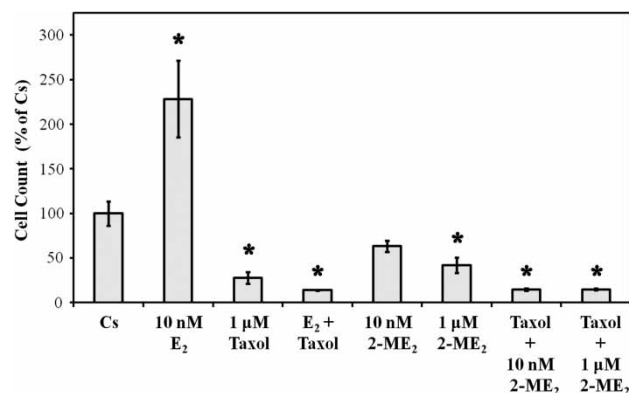


Figure 4. The effects of 2-ME₂, E₂, and Taxol on T47D cell proliferation. Cells were seeded into 12 well growth plates in medium containing 10% FBS at a density of 20,000 cells/well followed by culturing for 6 d in media that contained 5% DCC-stripped FBS and 2-ME₂, E₂, or taxol, either alone or in combination. Control and treatment media was changed every 48 h. Cells were trypsinized and counted by Coulter Counter (model # Z2) after six d of treatment. Mean cell populations ($\bar{x} \pm$ SEM, $n = 6$) were plotted as a percentage of control cells with 100% on the ordinate scale representing cells cultured in the absence of ligands ($\bar{x} = 3.2 \times 10^5$ cells/well). *indicates significant difference between treatment and control at $p < 0.05$ (Kruskal-Wallis Test followed by post-hoc analysis using Mann-Whitney U-Test).

DCC-FBS. This nuclear localization appears predominantly dispersed throughout the nuclear compartment which can be seen in the DAPI (nuclear counterstain) and p53 merged images. The addition of 2-ME₂, E₂, and/or taxol for 24 h, did not alter the cellular localization of this protein as it remained in the nucleus. However, treatment with 10 nM 2-ME₂ for 24 h, induced an alteration in the subnuclear localization of p53. Distinct p53 foci are visible that appear to overlap negative region/s in DAPI stained nuclei in approximately 10-20% of T47D cells. This observation potentially suggests p53 accumulation within nucleolar regions of cells [Maggi and Weber 2005]. This effect is consistently observed upon exposure of T47D cells to 10 nM 2-ME₂ for 24 h, and is also exhibited in cells treated with this concentration of the metabolite in combination with 1 μ M taxol.

Discussion

The p53 tumor suppressor protein is a nuclear phosphoprotein, which functions as a transcription factor that indirectly regulates the cell cycle. The transcriptional activity of p53 is determined by its ability to bind to sequence-specific DNA response elements [Cho et al. 1994; Hupp et al. 1992], and is modulated by the intranuclear concentration of the protein. Thus, measured levels of p53 are an important determinant of its activity [Hurd et al. 1995]. An alteration in p53 protein expression may indicate a potential role for this protein in the cellular response to new therapeutic modalities.

Previous studies in this laboratory have demonstrated that supplementation with physiological concentrations of E₂ (0.1 - 1 nM), induces an accumulation of the p53 protein in T47D breast cancer cells as well as an increase in cellular proliferation [Dinda et al. 2010; Hurd et al. 1995; Moudgil et al. 2001]. Among the steroid hormones known to significantly influence breast function, E₂ plays a crucial role in the progression of hormone-dependent breast cancer. Recent reports indicate that metabolites of E₂ may also have their own roles in the continuous growth or regression of cancer. It is possible that tumor progression may be regulated to some extent by estrogen metabolism, with some metabolites having mitogenic properties and others, such as 2-ME₂, inducing inhibitory responses. It is important to assess the role of E₂ metabolites in both normal and cancerous entities to distinguish the metabolic balance that may be required.

In the presence of 10 nM - 1 μ M 2-ME₂, there is an accumulation of the p53 protein in T47D cells (Fig. 1). One possible explanation for the observed increase in p53 protein expression may involve the c-Jun N-terminal kinase (JNK) signaling pathway [Fukui and Zhu 2009; Shimada et al. 2003]. It has been shown that activated JNK increases p53 stability by mediating p53 phosphorylation [Fuchs et al. 1998b; Fuchs et al. 1998c]. This induces conformational alterations in the protein inhibiting complex formation with inactive JNK and Mdm2 [Fuchs et al. 1998b;

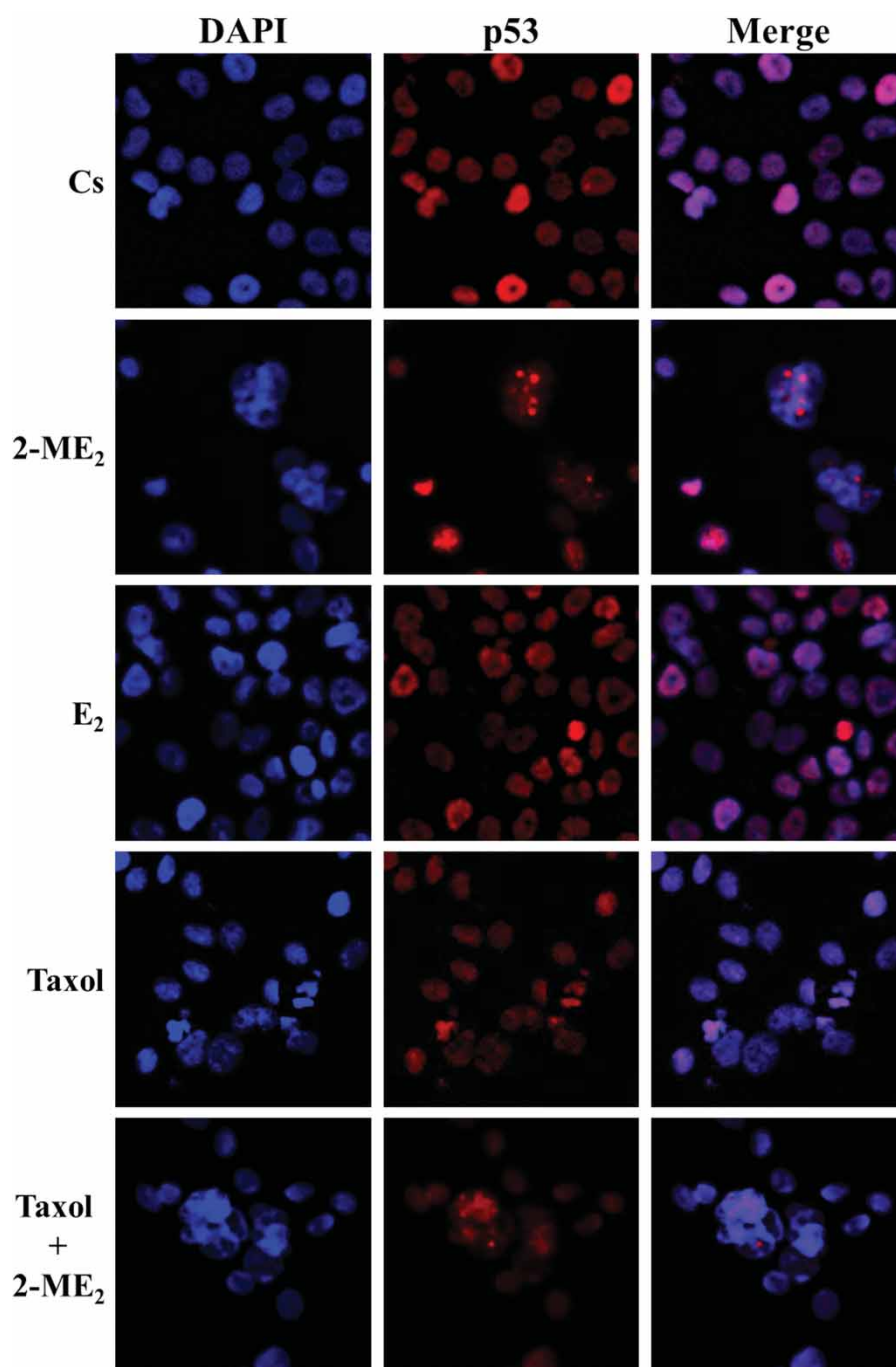


Figure 5. Immunolocalization of p53 in T47D cells: the effects of 2-ME₂, E₂, and Taxol. Cells were seeded at 15,000 cells/well on coverslips in 12 well growth plates for two d in medium containing 10% FBS, followed by culturing for six d in medium containing 5% DCC-stripped FBS. Cells were treated with 10 nM 2-ME₂, 10 nM E₂, and/or 1 μ M taxol for 24 h. Localization of p53 was determined by immunolabeling and the distribution was analyzed using a Nikon Digital Eclipse C1 plus modular confocal microscope. DAPI (blue) and Cy3 were employed to detect the immunolocalization of p53 (red). The acquired images were processed using Nikon Elements AR software. The merged images illustrate localization of p53 relative to DAPI nuclear counterstain.

Fuchs et al. 1998c]. Thus, the level of p53 is found to be inversely correlated with its direct association with inactive JNK and Mdm2 due to targeted ubiquitination of the tumor suppressor protein [Fuchs et al. 1998a; Haupt et al. 1997; Milne et al. 1995]. It was recently reported that 2-ME₂ induced phosphorylation/activation of JNK, ERK (extracellular signal-regulated kinase), and p38 in the

ER-negative MDA-MB435s breast cancer cell line [Fukui and Zhu 2009]. It has also been shown in the LNCaP prostate cancer cell line, 2-ME₂ up-regulated p53 through p38/JNK-dependent nuclear factor kappaB (NFkappaB)/activation protein-1 (AP-1) activation [Shimada et al. 2003]. Inhibition of JNK in LNCaP prostate cancer cells diminished induction of p53 protein by 2-ME₂ [Shimada

et al. 2003], and expression of a phosphorylation-deficient JNK mutant in ER-positive MCF-7 breast cancer cells inhibited the taxol-induced up-regulation of p53 protein levels [Fuchs et al. 1998b].

It is known that taxol-induced apoptosis requires JNK activation [Lee et al. 1998; Noguchi et al. 1999]. When 2-ME₂ was used in concurrent combination with 1 μ M taxol, the observed upregulation of p53 with 10 nM and 1 μ M 2-ME₂ alone was inhibited in T47D cells (Fig. 3). Differential effects on p53 have been reported upon treatment with taxol [Giannakakou et al. 2001] and p53 may or may not be inducible in various cell lines regardless of mutant or wt status. The presence of insulin has also been shown to modulate the taxol-induced response in MCF-7 cells inhibiting the upregulation of the tumor suppressor protein which is commonly found in this cell line [Miglietta et al. 2004]. This indicates that variations in p53 induction could be a result of the conditions by which these cell lines are cultured and reflects the high degree of sensitivity and complexity of these processes. Further studies are warranted to determine whether insulin or JNK activation play a role in the observed increase in p53 levels upon addition of 2-ME₂ in T47D cells.

The decrease in p53 levels upon 24 h treatment with 2.5 – 10 μ M 2-ME₂ in T47D cells, may be due to a process reported to be shared with taxol. 2-ME₂ is a known inhibitor of hypoxia-inducible factor-1 (HIF-1 α) [Mabjeesh et al. 2003; Mueck and Seeger 2010]. This inhibition is reported to be a post-transcriptional process that occurs following microtubule disruption and is shared with other microtubule disrupting agents including taxol [Mabjeesh et al. 2003]. Inhibition of specific protein biosynthesis may be a common process among microtubule disrupting agents, as this effect has also been shown by taxol on ER α [Martin et al. 2004]. The underlying mechanism that mediates this post-transcriptional process is currently unknown.

The data presented above shows the anti-proliferative effects of 2-ME₂. Other laboratories have previously shown that treatment with nanomolar concentrations of 2-ME₂ induced a mitogenic response in T47D cells that they believe is mediated by ER when cultured in the absence of exogenous E₂ [Han et al. 2005; Liu et al. 2005; Liu and Zhu 2004; Vijayanathan et al. 2006]. This study has consistently found an anti-proliferative effect of 2-ME₂ at concentrations of 1-100 nM in T47D cells, however, these results are not significant when compared to the control. Lottering et al. [1992] also reported antiproliferative effects in MCF-7 cells when treated with 10 nM 2-ME₂. Differences in these observations may be based on the culturing conditions or confluency of cultures during treatment. Fotsis et al. [1994] found that treatment with 2-ME₂ did not induce an inhibitory effect on cell proliferation in confluent monolayer endothelial and fibroblast cultures, but inhibition was apparent in cultures with lower cell densities. However, when confluent cultures were treated with 2-ME₂ in the presence of exogenous basic fibroblast growth factor, an inhibitory effect on proliferation that reflected non-confluent cultures was observed [Fotsis et al. 1994]. This indicates that 2-

ME₂-induced cytostatic effects occur only in actively cycling cells. It is possible that when lacking specific growth factor stimulation, physiological concentrations of 2-ME₂ act as a weak ER agonist inducing a mild mitogenic effect in quiescent cultures of ER-positive breast cancer cells. When cells are treated in the presence of exogenous E₂, the stimulatory effect induced by E₂ in these confluent cultures may then promote the anti-mitotic effects of 2-ME₂ leading to inhibition of proliferation and cell death.

The level of p53 protein is indicative of the cellular response to stressors [Vogelstein et al. 2000], and the localization of p53 is also a crucial factor in this response. In these studies, p53 localization remained nuclear upon 24 h treatment with 2-ME₂, E₂ and taxol. It is possible, that the protocol or antibody utilized may dictate this observation. Regardless, an interesting observation of the subnuclear localization of p53 was made in T47D cells that were treated with 10 nM 2-ME₂ for 24 h. It appears that this protein may be accumulating, possibly through sequestration, in regions that overlap those of the nucleoli. Further studies are needed in order to confirm if p53 is indeed localizing to nucleolar compartments or intranucleolar cavities. It has been reported that p53 localizes and/or accumulates in the nucleolus or within intranucleolar cavities in several cell lines under specific experimental conditions [Benninghoff et al. 1999; Horky et al. 2001; Karni-Schmidt et al. 2007; Karni-Schmidt et al. 2008; Klibanov et al. 2001; Latonen et al. 2003; Pokrovskaja et al. 2001; Rubbi and Milner 2000; Wesierska-Gadek et al. 2002]. The physiological relevance to this process and how p53 is targeted to nucleolar regions remains to be elucidated. Efforts are now underway to examine 2-ME₂ effects in MCF-7 cell line with wt expression of p53. We are also attempting to investigate mechanism of induction of p53 by 2-ME₂ in T47D cells. Results of these initial studies will form the basis of comparing effects of 2-ME₂ on p53 levels in its wt and mutated form.

In summary, these results presented above suggest that 2-ME₂ regulates p53 at the molecular level. Whether 2-ME₂ regulates p53 transcription, or whether the mechanism is indirect or post-transcriptional calls for future investigation. The observation that 2-ME₂ increased p53 levels in breast cancer cells, as well as blocking cellular proliferation, has implications in understanding the role of estrogen metabolite(s) in the regulation of tumor progression. Additional studies are warranted to determine the mechanism of 2-ME₂ action to provide more insight into the role of 2-ME₂ in p53 expression as well as its implication in hormone responsive breast cancer.

Materials and Methods

Cell Culture and Treatment with Ligands

T47D human breast cancer cells (HTB-133) were obtained from ATCC (Manassas, VA, USA). Cells were routinely cultured in RPMI-1640 with phenol red (Hyclone, Logan, UT, USA), containing 2 mM L-glutamine, 25 mM Hepes, antibiotic antimycotic solution (100 units/ml penicillin, 0.1

mg/ml streptomycin, and 0.25 mg/ml amphotericin B) (Hyclone), and 0.14 IU/ml insulin (Sigma, St. Louis, MO, USA) supplemented with 10% (v/v) FBS (Hyclone) and incubated at 37°C in the presence of 5% CO₂.

In all studies, T47D cells were subcultivated into phenol red-free RPMI-1640 media (Hyclone) containing the above components with alterations in the type and percentage of FBS utilized. Cells were plated in 10% FBS media and allowed to attach and grow for 48 h. To ensure steroid-free treatment conditions, the cell culture medium was changed to contain 5% DCC-FBS as described [Kodali et al. 1994], which depletes the serum of small heterocyclic molecules including serum growth factors. To ensure that the protein of interest was expressed only at basal levels prior to treatment with ligands [Hurd et al. 1995; Moudgil et al. 2001] the cells were cultured in 5% DCC-FBS for a total of six d, with fresh medium added every 48 h. On the 6th day, semi-confluent cells were treated with various ligands for 24 h. Ligands were dissolved in ethanol (2-ME₂, E₂) or methanol (taxol) and stock solutions were prepared to a 1,000-fold higher concentration. Aliquots (10 µl) of ligands were added directly to 10 ml of the culture medium (< 0.1% ethanol or methanol vehicle) and incubated for 24 h.

Protein Extraction of Cells

The culture medium was removed by aspiration, and the cells were washed with 10 ml ice cold Hank's balanced salt solution (HBSS). Cells were lysed in 500 µl of RIPA lysis buffer (50 mM Tris-HCl, 150 mM NaCl, 1.0% Triton X-100, 0.25% sodium deoxycholate, 5.0 mM EDTA) in the presence of 1 mM leupeptin, 30 µg/ml pepstatin-A, 30 µg/ml chymostatin, and 0.3 mM PMSF [Dinda et al. 1997; Dinda et al. 2010; Hurd et al. 1999; Moudgil et al. 2001], or 10 µl/ml Halt protease inhibitor cocktail (Thermo Fisher Scientific, Hanover Park, IL, USA) and cells were removed from the culture dish using disposable cell scrapers. A high speed supernatant of the extracts was prepared via centrifugation at 15,000 × g for 15 min at 4°C, and sample aliquots were frozen in liquid nitrogen and stored at -20°C until further use.

Quantification of Proteins

The protein concentration in individual samples was determined by using a standard protein assay (Bio-Rad, Hercules, CA, USA) based on the method of Bradford [1976]. The linear regression method was used to calculate the protein concentration of the individual samples, which was equalized prior to loading.

SDS-PAGE and Western Analyses

The supernatant was denatured (3 min, 100°C) and 25 µg aliquots of total protein/lane were loaded on a 4-20% polyacrylamide gel under denaturing conditions and electrophoresed for the separation of proteins. Proteins were wet transferred to Immobilon-P polyvinylidene fluoride (PVDF) membrane (Millipore, Bedford, MA, USA) by using a Bio-Rad trans blot cell in a tris-glycine buffer system containing 0.025% SDS and 15% methanol. Membranes were blocked for 1 h in

Tris-buffered saline (TBS)-tween (0.1%) plus 5% Carnation instant non-fat dry milk, then probed for 1 h in the same solution containing a 1:1000 dilution of anti-p53 monoclonal antibody (80/p53, BD Transduction Laboratories, San Jose, CA, USA) [Dinda et al. 2002; 2010; Hurd et al. 1995; Moudgil et al. 2001]. The membrane was washed for 30 min with three changes of TBS-tween, re-blocked for 15 min with TBS-tween plus 5% Carnation instant non-fat dry milk and then incubated in the same solution plus a 1:1000 dilution of horse radish peroxidase (HRP)-conjugated goat anti-mouse IgG_{2a} secondary antibody (Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA). After a final wash, p53-specific bands were visualized using autoradiography by the enhanced chemiluminescence (ECL) method according to instructions from Amersham. After immunoblotting, the PVDF membranes were then stained with coomassie blue to ensure the correct normalization against total protein levels and full transfer of the protein. The Western blots were subjected to quantitation of the protein band density using the ImageJ program (NIH, Bethesda, MD, USA).

Cell Proliferation Assays

All growth studies were conducted in 12-well culture plates. To ensure active, non-confluent cell populations during treatment duration, wells were initially seeded with 1.5×10^4 or 2.0×10^4 cells per well in 1 ml culture medium containing 10% FBS. The ligands were suspended in 5% DCC-FBS media and added to the wells 48 h later. Control wells received only 5% DCC-FBS culture medium. The culture medium in the absence or presence of ligands, was changed every 48 h. The proliferation studies were limited to eight d total duration to correlate with results of Western analyses. On the 8th day, the cells were washed with HBSS, trypsinized, and removed from individual wells of the culture plate. The resulting cell suspension was quantified by the Coulter counter model Z₂ (Coulter Electronics Inc., Hialeah, FL, USA).

Immunofluorescence and Confocal Microscopy

T47D cells were plated on cover slips in 12-well plates (15,000 cells/well) and cultured for 48 h in 10% FBS. The cell culture medium was then changed to 5% DCC-FBS and fresh medium was added at 2-d intervals. The cells were cultured in 5% DCC-FBS for a total of six d. On day six, the ligands were suspended in 5% DCC-FBS media, and semi-confluent cells were treated for 24 h.

Immunolabeling was performed as previously described for p53 [Dinda et al. 2002; 2010]. Briefly, the cells were fixed on cover slips for 10 min with 1% formalin in phosphate buffered saline (PBS), permeabilized with ice cold acetone and methanol (50:50) and washed three times with PBS. Staining procedures were performed in a humidified chamber at 25°C. Cells were incubated in 10% goat serum (Sigma) to suppress nonspecific binding of IgG, followed by 2 h incubation with 1:200 dilution anti-p53 (80/p53) monoclonal antibody. After washing with PBS, cells were incubated for 3 h with 1:200 dilution anti-mouse IgG conjugated with Cy3 (Jackson ImmunoResearch Laboratories,

West Grove, PA, USA). Cover slips were washed in PBS, and incubated for 2 min in 1 µg/ml 4',6-diamidino-2-phenylindole (DAPI) dissolved in PBS. Cells were washed three times in PBS, mounted with Fluoromount-G (Electron Microscopy Sciences, Hatfield, PA, USA) and stored in the dark at 4°C.

The distribution of three-dimensional fluorescent structures was analyzed using a Nikon Digital Eclipse C1 plus confocal microscope. NIS elements software (Nikon Instruments, Melville, NY, USA) was used for noise reduction and three-dimensional reconstruction of the images.

Statistical Analyses

The results are expressed as mean ± SEM. Statistical significance was determined by Kruskal-Wallis test followed by post-hoc analysis using Mann-Whitney U-test. Differences were considered significant at $p < 0.05$. Statistical analyses were carried out using SPSS for Windows version 11.5 (SPSS Inc., Chicago, IL, USA).

Acknowledgments

This research was made possible by support from Oakland University Cancer Research Program and Provost's Graduate Student Research Awards to AES and ALS. SD acknowledges support from Center for Biomedical Research, Oakland University.

Declaration of interest: The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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