



Ligand structure influences autologous downregulation of estrogen receptor- α messenger RNA

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Abstract

A series of A- and D-ring substituted estrogen analogues have been examined for their effect on estrogen receptor- α (ER α) mRNA downregulation. Recently it has been proposed that ER α autologous downregulation occurs via transcriptional repression exerted by the binding of the ER α -ligand complex to the 5' region of the coding region of the ER α gene. Placement of the phenolic hydroxyl group on the various carbons of the aromatic A-ring of estratrien-17 β ol (carbons 1–3) produced ligands which diminished the steady state level of ER α mRNA in relation to their affinity for receptor. 4-Hydroxyestratrien-17 β ol, on the other hand, was inactive in the downregulation of ER α mRNA. Although this A-ring isomer brought about apparent processing of the nuclear receptor, the ER α reappeared in the cytosol within 24 h. Unlike the stimulation of genes regulated via estrogen response elements, maximum autologous negative regulation of the ER α gene required the presence of an hydroxyl group on carbon 17 of the D-ring. These results suggest that the conformational alterations elicited in the ER α molecule by various ligands create surfaces capable of interacting with other transcription factors in a manner which is different when the receptor functions via a response element mechanism relative to interactions during autologous negative regulation of the ER α gene. © 1999 Published by Elsevier Science Ltd. All rights reserved.

1. Introduction

Cells of hormone-dependent breast tumors contain a specific protein member of the nuclear-receptor superfamily which binds estradiol-17 β (E₂) with high affinity. Activation of this receptor by ligand binding results in conformational changes which enable the complex to interact with, and perform as, transcriptional regulation factor(s). Positive regulation of estrogen responsive genes has been documented by numerous laboratories [1–6]. Information pertaining to negative regulation of transcription by estrogens is limited. Estrogens are known to reduce the transcription

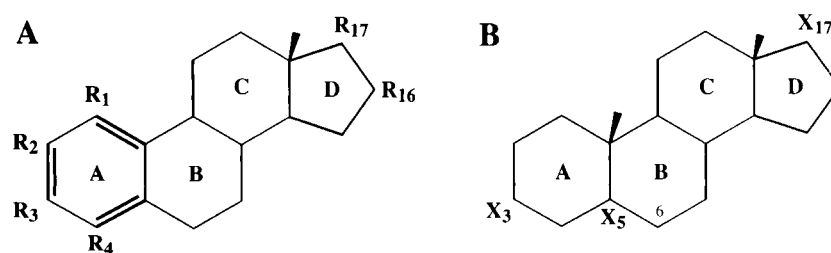
of c-jun [7] and the β -subunits of glycoprotein hormones [8] as well as various other uncharacterized proteins in MCF-7 cells [9]. However, the mechanism for this negative regulation remains obscure. E₂ has been shown to decrease the steady state levels of both estrogen receptor- α (ER α) mRNA and protein in many estrogen responsive tissues and cell lines (chick oviduct, endometrial cancer cells and MCF-7 and other cancer cells) by many investigators [10–13].

ER 'processing' was initially defined as the loss of approximately 50% of ligand binding, or monoclonal antibody recognizable, ER α within 6 h of E₂ exposure, which is not explained by the return of these 'processed' receptors to the cytosol [14]. However, the cloning of the ER α cDNA allowed a more detailed examination of this phenomenon and processing recently has been associated with downregulation of ER α mRNA [13,15,16]. The observed decrease in the rate of ER α gene transcription was cycloheximide resistant and dependent on the concentration of E₂

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A: ESTROGENS						
	R₁	R₂	R₃	R₄	R₁₆	R₁₇
Estratriene	H ₂	H ₂	H ₂	H ₂	H ₂	H ₂
3-Hydroxyestratriene	H ₂	H ₂	OH	H ₂	H ₂	H ₂
Estratriene-17β-ol	H ₂	H ₂	H ₂	H ₂	H ₂	β-OH
Estradiol-17β (E ₂)	H ₂	H ₂	OH	H ₂	H ₂	β-OH
Estradiol-17α	H ₂	H ₂	OH	H ₂	H ₂	α-OH
Estradiol-16α	H ₂	H ₂	OH	H ₂	α-OH	H ₂
1-Hydroxyestratrien-17β-ol	OH	H ₂	H ₂	H ₂	H ₂	β-OH
2-Hydroxyestratrien-17β-ol	H ₂	OH	H ₂	H ₂	H ₂	β-OH
4-Hydroxyestratrien-17β-ol	H ₂	H ₂	H ₂	OH	H ₂	β-OH
Estrone	H ₂	H ₂	OH	H ₂	H ₂	keto
Estriol	H ₂	H ₂	OH	H ₂	α-OH	β-OH

B: ANDROGENS			
	X₃	X₁₇	X₅
5-Androstene-3β,17β-diol	β-OH	β-OH	C5-C6 double bond
5α-Androstane-3β,17β-diol	β-OH	β-OH	α-H
5α-Androstane-3α,17β-diol	α-OH	β-OH	α-H

Fig. 1. Structures of estrogen analogues and androstanediols.

[15,16]. The absence of an estrogen response element (ERE) in the promoter of the gene suggests a unique ER α -DNA interaction in the regulation of the ER α gene. Recently, it has been proposed that ER α (or glucocorticoid receptor) autologous downregulation occurs via a transcriptional repression exerted by the estrogen (or glucocorticoid) receptor on their genes [17,18]. It is proposed that these receptors bind to specific DNA sequences within the gene. The rate at which functional ER α is synthesized and degraded (turnover) in estrogen responsive cells is a major factor in regulating the level of ER α and therefore cell responsiveness to E₂. Modulation of receptor level is

quite possibly one mechanism which governs target gene transcription in response to estrogens.

Recent reports from this laboratory have demonstrated an influence of estrogen structure on the positive regulation of certain estrogen responsive genes (progesterone receptor [6], cathepsin D and pS2 [5] and tissue plasminogen activator [3]) in MCF-7 cells. As with positive regulation, the ligand-dependent downregulation of ER α may involve an estrogen structure-dependent mechanism. The studies described herein examined the effect of ER α downregulation of 14 estrogen ligands with modifications to the A- and D-rings.

2. Materials and methods

2.1. Steroids, antiestrogens and inhibitors (Fig. 1)

Estratriene, estrone, E₂, estradiol-16 α and estradiol-17 α were purchased from Research Plus (Bayonne, NJ). Other estrogens used in these investigations were synthesized in this laboratory. 1-, 2- and 4-hydroxyestratrien-17 β -ol, estratrien-17 β -ol and 3-hydroxyestratriene were synthesized according to published procedures [19–21]. The purity of each estrogen analogue was guaranteed by thin layer chromatography and crystallization. Chromatography was carried out on thin layer chromatography plates (silica gel), developed three times with hexane:ethylacetate (5:5), which resolved approximately one mg of the compounds (3–4 μ mol). After exposure to iodine, spots were visualized under ultraviolet light (254 nm). This procedure enabled detection of 0.1 nmol of E₂. Thus, each analogue was determined to contain less than one part in 10,000 of contaminating estrogens [6].

The androstane diols (5-androsten-3 β ,17- β -diol, 5 α -androstan-3 β ,17- β -diol and 5 α -androstan-3 α ,17- β -diol) were purchased from Aldrich Chemical Co. (Milwaukee, WI) and purified by thin layer chromatography as described above [6]. The antiestrogen, 4-hydroxytamoxifen, was a gift from Stuart Pharmaceuticals (Division of ICI Americas, Wilmington, DE) and ICI-164,384 was kindly supplied by Dr. A.E. Wakeling, Imperial Chemical Industries (Alderly Park, UK). Cycloheximide and actinomycin-D were purchased from Sigma, St. Louis, MO.

2.2. Cell culture

A subclone of MCF-7 designated E3 [22] was maintained in a custom formulation of Eagles modified MEM (Gibco Formula 86-5126EF, Gaithersburg, MD) with Hank's balanced salt solution (HBSS), L-glutamine, 25 mM HEPES without phenol red and supplemented with nonessential amino acids (Sigma, St. Louis, MO), 5% donor calf serum (Flow Labs, Lot #9010763, McLean, VA), 0.5 μ g/ml gentamicin sulfate (Sigma), and 0.01 μ g/ml phenol red (Sigma). Cells were incubated at 37°C in closed flasks. All experiments were carried out on cells which had been passed between 170 to 200 times. For experiments, cells were passed at approximately 2×10^6 cells/75 cm² tissue culture flask (Corning, Corning, NY) and allowed to grow for 5–6 days (approximately 70% confluent) in MEM supplemented with 5% donor calf serum, 0.5 μ g/ml gentamicin and phenol red. Cells were then transferred to a phenol red-free media that contained dextran-coated charcoal (DCC) treated [23] calf serum (Gibco Lot #38N1091). Cultures were allowed to grow for three days. After this estrogen withdrawal period,

the cells were washed with phosphate buffered saline (PBS; 137 mM sodium chloride, 2.7 mM potassium chloride, 6.5 mM dibasic sodium phosphate, 1.5 mM monobasic potassium phosphate, pH 7.4). The cells were then exposed to the estrogens (10^{-11} to 10^{-6} M), added in 20 μ l ethanol, for 18 h in the media containing DCC treated serum. Experiments were carried out in duplicate.

2.3. ER-EIA

The treated cells were washed with cold saline and removed from flasks in 5 ml MEM utilizing a wide bore syringe followed by centrifugation of the cell suspension at $150 \times g$ for 10 min at 4°C. For the determination of cytosolic receptor cells in the pellet were ruptured with a Dounce homogenizer in ice-cold immunoassay buffer (10 mM tris, 1.5 mM EDTA, 5.0 mM sodium molybdate pH 7.4 with 1.15 μ M monothioglycerol). The homogenate was centrifuged for 1 h at $100,000 \times g$ at 47°C and the supernatant assayed for ER α using the enzyme immunoassay (ER-EIA) kit supplied by Abbot Laboratories (Chicago, IL). This kit utilized the sandwich technique employing the D547 and H222 ER α specific antibodies. The pellets were utilized for DNA determination by the method of Burton [24].

Determination of the nuclear ER α was carried out according to the protocol described by Kral et al. [25]. Briefly, the cellular pellet was resuspended in 1 ml of 0.5% NP₄₀ cell lysis buffer (10 mM HEPES, pH 7.4 at 47°C, 10 mM NaCl, 5 mM monothioglycerol, 0.5% NP₄₀ and 5 mM magnesium chloride) and kept cold for 10 min with occasional mixing. Following a second lysis, the nuclear pellet (150 g, 10 min) was extracted at 4°C with lysis buffer containing 0.4 M KCl (2 ml) by mixing every 2 min for 10 min. This mixture was centrifuged and the nuclear extract assayed (EIA) for the amount of nuclear estrogen receptor complex and the pellets were utilized for DNA determinations. The ER-EIA Kit contains lyophilized ER α standard from MCF-7 cells from which a standard curve was constructed. This method has been shown to recover 95% of the receptor in MCF-7 cells [25].

2.4. Messenger RNA isolation and Northern analysis

Messenger RNA was isolated from cell lysates according to a modified one-step procedure of Chomczynski and Sacchi [26] and the mRNA resolved by electrophoresis (3.0–6.0 μ g per lane) on a 1.2% agarose gel as reported previously [3]. Northern analysis was performed [3] with radioactive probes prepared with the cDNAs for the human ER α and the 36B4 housekeeping gene. The washed and dried membranes were exposed to film (Kodak Diagnostic Film X-

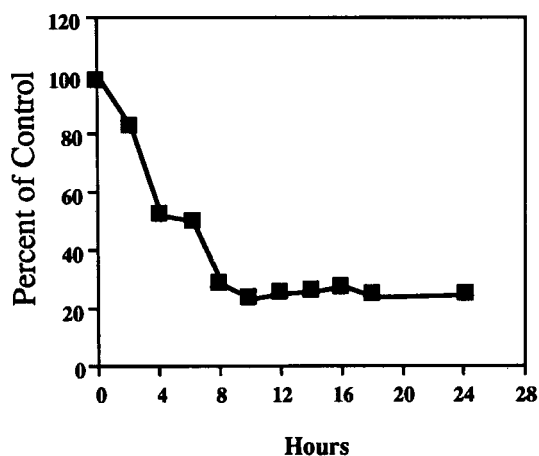


Fig. 2. Plot of the time-dependent downregulation of ER α mRNA in MCF-7 cells (clone E-3) by E $_2$. Confluent cultures maintained in estrogen free media 72 h were exposed to 10 $^{-8}$ M E $_2$ for the noted times and harvested. mRNA was collected as described in the Materials and methods. Desitometric analysis of the autoradiographs from Northern blots of the mRNA samples were normalized to the internal standard (36B4) and plotted as percentage of the normalized control (time=0). Points are the average of duplicate flasks.

OMAT, Rochester, NY) for 18 h before developing and scanned with a Molecular Dynamics densitometer employing the Imagequant[™] software (Sunnyvale, CA). Recorded density units were related to the amount of RNA applied to each lane and to the radioactivity in each band from the hybridized cDNA.

2.5. Preparation of DNA probes for Northern analysis

The 1.9 kb cDNA for ER α and the estrogen insensitive gene, 36B4 (both cDNAs graciously supplied by P. Chambon), were inserted into pBR322. These plasmids were transformed into competent cells of *E. coli* strain DH5alpha (Gibco/BRL), amplified, extracted and purified using procedures described previously [3]. Restriction digests were performed [3] utilizing PstI for 36B4; and *EcoRI* for ER α cDNA and fragments purified by agarose gel electrophoresis. The inserts were cut out and recovered via electroelution for 10–12 h using an Elutrap (Schleicher and Schuell).

Radiolabeled probes of each cDNA were made [3] using a commercially available random primed labeling kit (Boehringer Mannheim, Indianapolis, IN). A 1 μ l sample from each probe was counted in a Packard Tri-Carb 4530 scintillation counter (Packard Instruments, Downers Grove, IL) to determine the radioactivity. Approximately 2.0 \times 10 7 cpm of each probe was added to the hybridization buffer after denaturation at 100°C for 4 min.

3. Results

3.1. The effect of E $_2$ and antiestrogen

Exposure of cultures of MCF-7 cells to 10 $^{-8}$ M E $_2$ for varying periods of time brought about a reduction in the expression of ER α mRNA (Fig. 2). After 2 h the level of ER α mRNA had decreased to 82% of that originally in the MCF-7 cells. This negative regulation continued, reaching a minimum value after 10 h (Fig. 2). The low level of ER α mRNA expression was maintained for the remaining 14 h of the experiment. Others have shown ER α mRNA expression to be repressed for at least 48 h after exposure of MCF-7 cells to E $_2$ [13,15]. E $_2$ downregulation of ER α mRNA was maximal at a concentration of 10 $^{-11}$ M, with higher concentrations being equally effective (Fig. 3(A), depicted as percent maximum downregulation).

Two vastly different antiestrogens effectively stimulated the downregulation of ER α mRNA after 18 h exposure. ICI164,384 a ‘pure’ antiestrogen reduced the cellular levels of ER α comparable to the effect of E $_2$ at a concentration of 10 $^{-10}$ M (Fig. 4). Likewise the partial agonist, 4-hydroxytamoxifen, decreased ER α to 50% of its initial level in untreated cells when present in culture media at 10 $^{-9}$ M. The apparent downregulation (following a 12 h exposure) of ER α mRNA by 4-hydroxytamoxifen has been reported [16]. The effect of ICI164,384 on ER α protein has been attributed to an increased rate of turnover of the receptor [27,28]. However, other antiestrogens such as LY117018 are ineffective [15]. Progesterone, which is often shown to possess antiestrogen activity blocks the E $_2$ autologous downregulation of ER α in these cells [15]. Yet in T47D cells, progestins are effective in the suppression of this receptor by blocking ER α synthesis [15].

3.2. Effect of structurally altered estrogens and androstane diols

The estratriene nucleus, devoid of hydroxyl groups, was capable of effecting a minimal downregulation of ER α mRNA over a concentration range of 10 $^{-9}$ –10 $^{-7}$ M (Fig. 3(B)). This repression of ER α mRNA accumulation occurred regardless of the immeasurably weak affinity of estratriene for ER α (relative binding affinity, RBA < 0.0005, relative to E $_2$, K_a = 3.7 \times 10 9 M $^{-1}$, RBA = 1.0). Repositioning the phenolic hydroxyl of E $_2$ to other A-ring carbons yielded estrogens with varying effects on ER α mRNA repression. 1-Hydroxyestratrien-17 β -ol was 10,000-fold less active than E $_2$, decreasing ER α mRNA maximally at 10 $^{-7}$ M (Fig. 3(C)). This compound binds weakly to ER α , with a RBA of 0.005. 2-Hydroxyestratrien-17 β -ol, which binds 150 times tighter (RBA = 0.71) to the receptor than 1-hydroxyestratrien-17 β -ol was 100-fold more

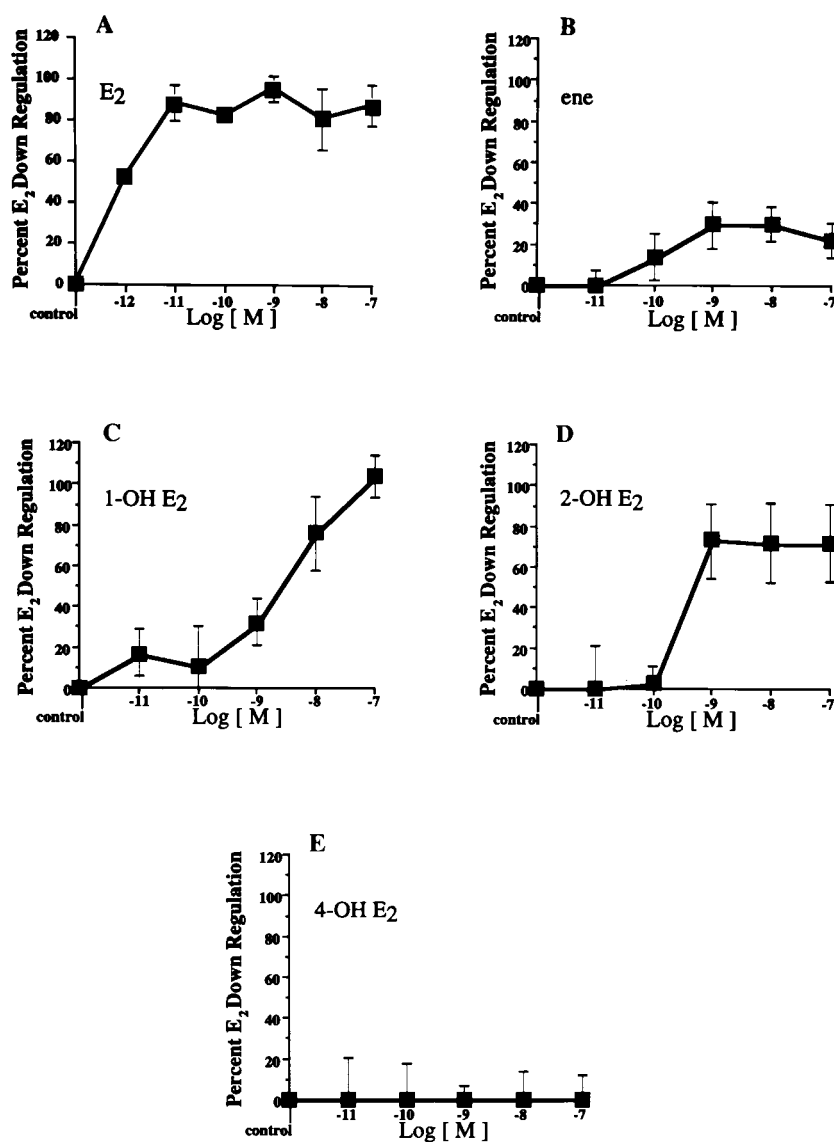


Fig. 3. Percent maximum downregulation of ER α mRNA in MCF-7 cells (clone E3) pulsed with different concentrations of estrogen analogs. Cultures were maintained and exposed to estrogens for 18 h as described in Materials and methods. mRNA was collected as described in Materials and methods. Densitometric analysis of the autoradiographs from Northern blots of the mRNA samples were normalized to the internal standard (36B4) and plotted as a percentage of maximum E₂ (10⁻⁸M) downregulation. In these experiments, the normalized E₂ density ranged from 35–65% of control values. Cultures were treated with (A) E₂, (B) estratriene (ene), (C) 1-hydroxyestratrien-17 β -ol (1-OHE₂), (D) 2-hydroxyestratrien-17 β -ol (2-OHE₂) and (E) 4-hydroxyestratrien-17 β -ol (4-OHE₂). Standard deviations of several determinations are shown as brackets.

active, downregulating ER mRNA maximally at 10⁻⁹ M (Fig. 3(D)). Interestingly, 4-hydroxyestratrien-17 β -ol, which bound ER α an order of magnitude better than 1-hydroxyestratrien-17 β -ol (RBA=0.07) was completely inactive in the downregulation of ER α mRNA, over the entire concentration range tested (Fig. 3(E)). This A-ring isomer, which has been shown to function as an antiestrogen [3], effectively blocked E₂ downregulation of ER α mRNA when its concentration in the culture medium reached 10⁻⁷ M, a level at which 4-hydroxyestratrien-17 β -ol competed with the

added 10⁻⁸ M E₂ for receptor (the K_a of 4-hydroxyestratrien-17 β -ol is 1/10th that of E₂, Fig. 5).

Placement of a phenolic hydroxyl group on position-3 of estratriene yielded a monohydroxyestrogen capable of downregulating ER mRNA half maximally, at concentrations of 10⁻¹⁰M and above (Fig. 6(A)). 3-Hydroxyestratriene was incapable of decreasing ER mRNA to the level brought about by 10⁻¹¹M E₂ when added to cultures of MCF-7 cells at concentrations up to 10⁻⁷M. This compound, 3-hydroxyestratriene, has a high affinity for ER α , binding approximately 80% as

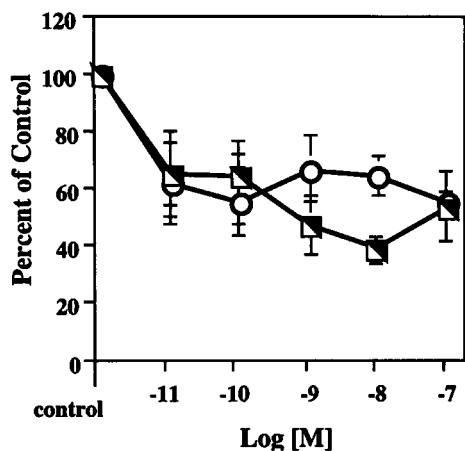


Fig. 4. Downregulation of ER α mRNA by antiestrogens. Confluent cultures were maintained in estrogen free media for 72 h then exposed to antiestrogens for 18 h and harvested. mRNA was collected as described in the Materials and methods. Downregulation of ER α mRNA by ICI 164,384 (○) and 4-hydroxytamoxifen (■) is shown. Standard deviations are depicted as brackets for four to six experiments.

well as E₂ (RBA=0.79). On the other hand, placing the D-ring hydroxyl group of E₂ on estratriene yielded a monohydroxyestrogen, estratrien-17 β -ol (RBA=0.11) with maximal negative regulation of ER α mRNA at 10⁻⁸M (Fig. 6(B)). In fact, all ligands tested which possessed a 17-hydroxyl group (e.g. E₂, Fig. 3(A), estradiol-17 β , RBA=0.22 and estriol, RBA=0.17, Fig. 6(C and E)) brought about maximal negative regu-

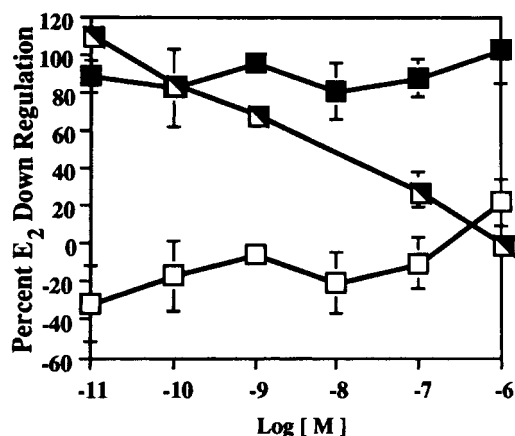


Fig. 5. Downregulation of ER α mRNA in MCF-7 cells (clone E-3) by E₂, 4-hydroxyestratrien-17 β -ol and combinations. Confluent cultures were maintained in estrogen free media for 72 h then were supplemented with increasing concentrations of 4-hydroxyestratrien-17 β -ol (□); increasing concentrations of 4-hydroxyestratrien-17 β -ol + 10⁻⁸M E₂ (■) or increasing concentrations of E₂ (■) and harvested after 18 h. mRNA was collected as described in the Materials and methods. Densitometric analysis of the autoradiographs from Northern blots of the mRNA samples were normalized to the internal standard (36B4) and plotted as percentages of the normalized control sample (35–65%). Standard deviations of several experiments are shown as brackets.

Table 1
EC₅₀ values of A- and D-ring substituted estrogens

Analogue	RBA ^a	EC ₅₀ ^b
(1) Estradiol-17 β	1.0	1.1 × 10 ⁻¹¹
(2) Estradiol-16 α	0.80	1.0 × 10 ⁻¹⁰
(3) 3-Hydroxyestratriene	0.79	1.0 × 10 ⁻¹⁰
(4) 2-Hydroxyestratrien-17 β -ol	0.67	3.1 × 10 ⁻⁹
(5) Estrone	0.27	7.9 × 10 ⁻¹⁰
(6) Estradiol-17 α	0.27	7.7 × 10 ⁻¹⁰
(7) Estriol	0.17	1.4 × 10 ⁻⁹
(8) Estratrien-17 β -ol	0.11	4.5 × 10 ⁻¹⁰
(9) 4-Hydroxyestratrien-17 β -ol	0.071	1.0 × 10 ⁻⁶
(10) 5-Androstene-3 β , 17 β -diol	0.0073	3.7 × 10 ⁻⁹
(11) 5 α -Androstane-3 β , 17 β -diol	0.0049	1.0 × 10 ⁻⁸
(12) 1-Hydroxyestratrien-17 β -ol	0.0046	8.4 × 10 ⁻⁹
(13) 5 α -Androstane-3 α , 17 β -diol	< 0.0005	6.1 × 10 ⁻⁷
(14) Estratriene	< 0.0005	1.0 × 10 ⁻⁶

^a Relative binding affinities determined by the competitive binding assay as described in Vanderkuur et al. [6]. E₂, K_a=3.7=1.5 × 10⁹M⁻¹ is set at 1.0.

^b EC₅₀ is defined as the concentration of analogue which produces half maximal response. EC₅₀ was calculated for each analogue as log [(% of maximal E₂ response)/(100-(% of maximal E₂ response))]. Compounds that did not reach 50% maximal response were given an arbitrary property value of 1 × 10⁻⁶ M (Sybyl Molecular Modeling Program — Theory Manual, 1992).

lation of ER α mRNA. As with 3-hydroxyestratriene, a 16 α -hydroxyl (estradiol-16 α , RBA=0.80, Fig. 6(D)) or a 17-keto substituent (estrone, RBA=0.22, Fig. 6(F)) produced half-maximal downregulation of the ER α mRNA.

Without the aromatic character of the A-ring of E₂, steroids such as 5 α -androstane-3 β , 17 β -diol (RBA=0.005) or 5-androstene-3 β , 17 β -diol (RBA=0.007) were capable of downregulating ER mRNA maximally at a 10⁻⁷ M concentration, yielding an EC₅₀ between 10⁻⁸ and 10⁻⁹ M (Table 1). Repositioning the orien-

Table 2
Down-regulation of ER α mRNA in MCF-7 cells exposed to inhibitors of RNA and protein synthesis

Treatment ^a	Relative percentage of control ^b		
	-E ₂	+E ₂	Effect of added E ₂ % Change ^c
None	100	34.2 ± 17.7	-65.8
Actinomycin-D	17.0 ± 15.5	18.4 ± 14.2	+8.2
Cycloheximide	41.0 ± 12.0	19.6 ± 12.0	-52.2

^a MCF-7 cells were exposed to no inhibitor (none), actinomycin-D (4.0 μ M), or cycloheximide (50 μ M) in estrogen free media for 20 min prior to exposure ethanol vehicle (-E₂) or 10⁻⁸M E₂ (+E₂). The cells were harvested after 18 h exposure.

^b Value represents the relative percentage of the corrected band density from Northern analysis of mRNA isolated from control cells which were not treated with inhibitor or exposed to estradiol.

^c Difference in the level of ER α mRNA in control (-E₂) and E₂ treated (+E₂) cells in control (none) or inhibitor treated cells.

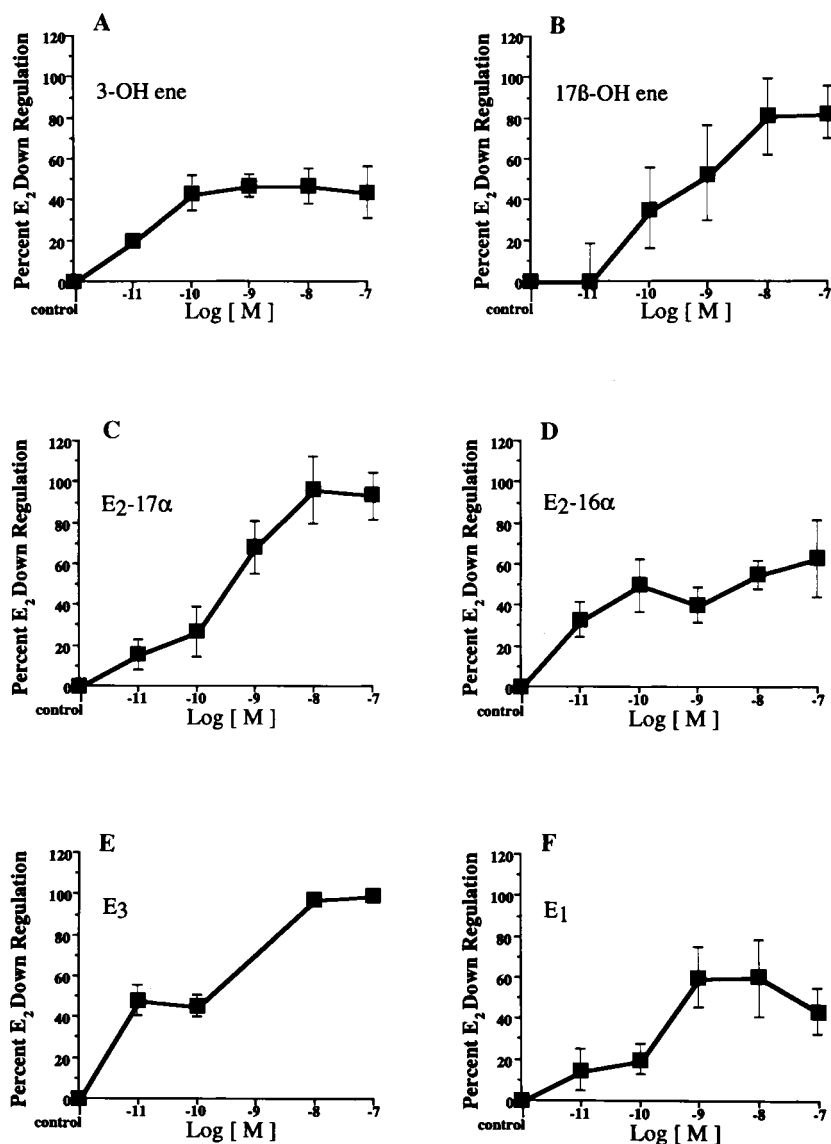


Fig. 6. Percent maximum downregulation of ER α mRNA in MCF-7 cells (clone E3) pulsed with different concentrations of estrogen analogues. Cultures were maintained and exposed to estrogens for 18 h and the mRNA was collected as described in Materials and methods. Densitometric analysis of the autoradiographs from Northern blots of the mRNA samples were normalized to the internal standard (36B4) and plotted as a percentage of maximum E₂ downregulation. In these experiments the normalized E₂ density ranged from 35–65% of control values. Cultures were treated with (A) 3-hydroxyestratriene (3-OHene), (B) estratrien-17 β -ol (17 β -OHene), (C) estradiol-17 α (E₂-17 α), (D) estradiol-16 α (E₂-16 α), estriol (E₃), (E) estrone (E₁). Standard deviations of several experiments are shown as brackets.

tation of the 3 β -hydroxyl group of 5 α -androstane-3 β , 17 β -diol to the 3 α configuration, resulted in a compound, 5 α -androstane-3 α ,17 β -diol (RBA = < 0.0005), which brought about less than half maximal downregulation of ER α mRNA accumulation at the highest concentration tested 10⁻⁷ M (Table 1).

3.3. Effect of inhibitors

Actinomycin-D is known to reduce the E₂ induced downregulation of ER α mRNA in MCF-7 cells

[14,29]. Confirming these observations, we found the estrogen-dependent reduction in the steady state level of ER α mRNA was totally blocked by 4.0 μ M actinomycin-D (Table 2). Incubation of MCF-7 cells with cycloheximide has no apparent effect on the estrogen induced downregulation of ER α mRNA under the conditions of our experiments (Table 2 and Ref. [13]). However, both actinomycin-D and cycloheximide caused an apparent reduction in the overall yield of ER α mRNA in these experiments (Table 2).

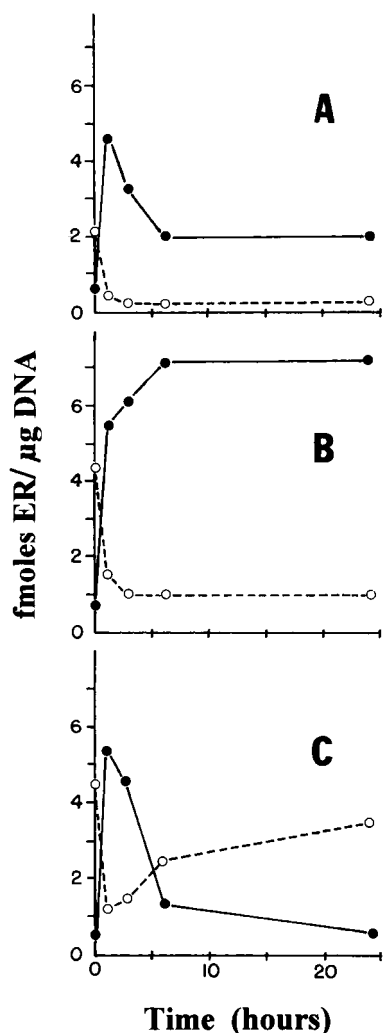


Fig. 7. Levels of cytosolic (○—○) and nuclear (●—●) ER α during a 24-h exposure of MCF-7 cells to: (A) E₂ (10⁻⁸M), (B) 4-hydroxytamoxifen (10⁻⁸M) and (C) 4-hydroxyestratrien-17 β -ol (10⁻⁷ M). EIA were carried out as described in Materials and methods. Points are the mean of several determinations with standard deviations of less than 10%. Patterns shown were independent of ligand concentration.

3.4. Processing of ER in MCF-7 cells

The often reported 'classical' loss of ER in MCF-7 cells treated with E₂ is depicted in Fig. 7(A). Within 1 h of exposure to E₂, a maximum level of tightly bound receptor is found in the nuclear fraction, followed within 6 h by a diminished level (50%) of detectable ER which persists for 24 h. Concurrently, cytosolic ER remained low from 1 to 24 h. In these experiments 4-hydroxytamoxifen was capable of bringing about tight nuclear binding. However, processing of this nuclear receptor did not occur (Fig. 7(B)). On the other hand, 4-hydroxyestratrien-17 β ol, promptly brought about the binding of ER to DNA, which disappeared from the nuclear fraction by 6 h. However, in the case of this A-ring isomer the level of nuclear

ER continued to diminish for 24 h (Fig. 7(C)). Unlike any of the other ligands examined in these experiments, immediately after the binding of ER to nuclei (1 h), receptor began to reappear in the cytosol following administration of the 4-hydroxy isomer of E₂. By 24 h, 3.5 fmol/ μ g DNA of cytosolic ER was detected, or 80% of the initial ER was present (cytosol and nuclear). All of the other estrogen analogues yielded a pattern of processing similar to that of E₂ (data not shown).

4. Discussion

ER processing has been documented in mammary carcinoma cells and many other experimental systems [12,14,30]. This phenomenon undoubtedly influences a target cell's response to estrogen stimulation. The purpose of this study was to examine the effects of structurally altered estrogens on the homologous downregulation ('processing') of ER α mRNA. The MCF-7 cell line chosen for these studies expresses ample levels of ER α mRNA and protein. ER β protein has not been detected in these cells and only trace quantities of the mRNA for this receptor subtype has been detected by PCR [31].

Binding affinities for ER α of the various estrogens and androstane diols employed in these experiments have been calibrated [6], and shown, with two exceptions, to decrease as much as 200-fold less than the K_a of E₂. The exceptions being estratriene and 5 α -androstane-3 α ,17 β -diol which possessed affinities too low to measure. Nevertheless, the range of concentrations of these compounds added to cultures of MCF-7 cells was 10⁴-fold above the level (10⁻¹¹ M) at which E₂ displayed its maximum repression of ER α mRNA levels. Therefore each ligand (with the exception of estratriene and 5 α -androstane-3 α ,17 β -diol) examined would be expected to influence the 'processing' of ER α mRNA in this system, assuming in each case, the activity was related directly to the K_a for ER α . Overall, a plot which depicts the capacity of each compound to stimulate negative regulation of ER α mRNA shows excellent correlation ($R^2=0.82$) with the binding affinity for receptor of the estrogens and androstane diols (Fig. 8). The only anomaly was 4-hydroxyestratrien-17 β -ol which bound receptor with an affinity comparable to, or greater than, a number of estrogens that were active in the downregulation of ER α mRNA. Yet, this dihydroxyestrogen was inactive. Previously, it had been demonstrated that 4-hydroxyestratrien-17 β -ol was incapable of stimulating the induction of several mRNA's (cathepsin D and pS2, [5] and tissue plasminogen activator [3], also Fig. 8) whereas all other estrogen analogues tested induced these genes. In fact,

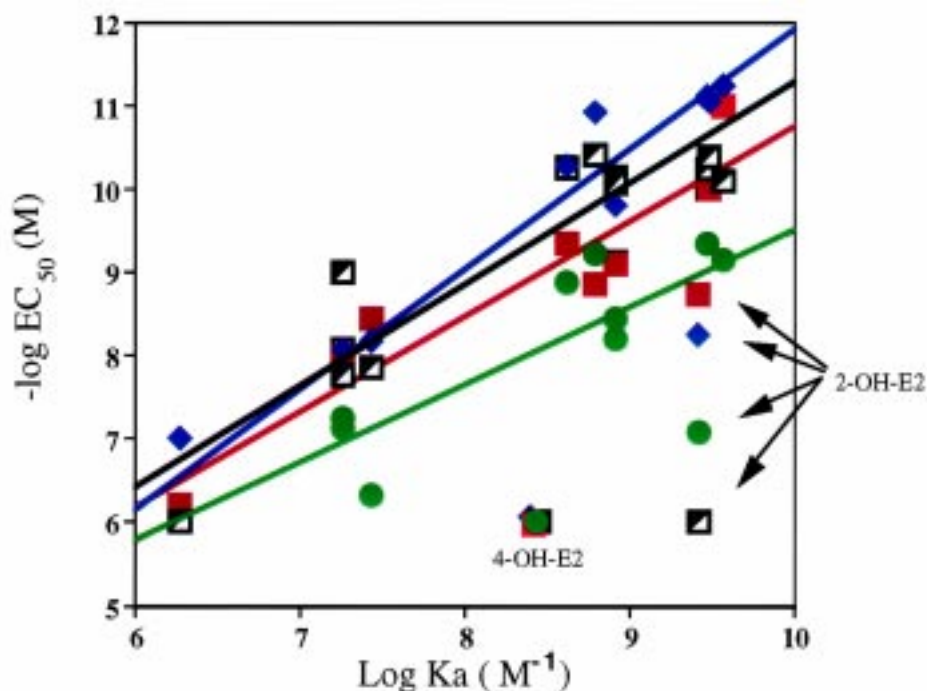


Fig. 8. Relation of K_a of selected estrogens and androstane diols to the induction of Cath D mRNA (■); pS2 mRNA (◇); t-PA mRNA (○) and the downregulation of ER mRNA (□). EC_{50} is defined as the effective concentration which produced a half-maximal response. The EC_{50} of each analogue was calculated as $\log [\% \text{ of maximal } E_2 \text{ response}] / [100 - (\% \text{ of maximal } E_2 \text{ response})]$. Analogs which had an inductive effect less than 50% maximal response were given a low and arbitrary property value of 10^{-6} [3,5]. K_a 's were determined from the RBAs using the K_a of E_2 ($3.7 \times 10^9 M^{-1}$). R^2 values for the lines determined from the Cath D mRNA, pS2 mRNA, t-PA mRNA and ER mRNA results are 0.85, 0.97, 0.89 and 0.92, respectively, when the points for 2-hydroxyestratrien-17 β -ol and 4-hydroxyestratrien-17 β -ol are omitted. The ligands listed in Table 1 were used for these experiments.

there was a strong correlation between gene induction and receptor affinity in these previous experiments.

Although the estrogen analogues employed in these experiments have not, as yet, been examined by others, there exist abundant studies of ligand structure–estrogen receptor affinity relationships. These published data have recently been utilized to create an estrogen pharmacophore [32]. Regarding the K_a of estrogen analogues for receptor, these data demonstrated, as had Anstead et al. [32], that the C3- and 17 β -hydroxyl groups are major contributors to ligand affinity, with the phenolic group contributing the greatest energy as an H-donor. The proximity of the 2-hydroxyl group apparently allows it to participate as an H-donor as well. A 3D QSAR study of 42 estrogen-like analogues (including those in this report) has been carried out in our laboratory [33] in which contour plots created for receptor affinity, as well as for a biologic response (growth) were compared. Most notable in these plots are the expanded steric and electropositive tolerance zones around the A-ring. Also, steric and electrostatic contours around the D-ring are depicted in this model which help explain the preference for 17 β -hydroxysteroids over the less biologically active 17 α and 16 α -hydroxy compounds.

As we have suggested, endogenous gene regulation by estrogens is not related directly to the affinity of ligand for ER α , nor to the presence of either hydroxyl group, but this activity may be determined by the geographic location of the electronegative isopotential above the A-ring and the influence a variation in this structural characteristic may exert on the conformation of the receptor complex and its capacity to interact with the transcription apparatus [3,5,6,33].

The atypical effect of 4-hydroxyestratrien-17 β -ol on the conformation of full length ER α has been demonstrated in gel-shift assays [34]. In experiments wherein the concentrations of ligands were adjusted to assure receptor occupancy, 4-hydroxyestratrien-17 β -ol did not induce a shift of the ERE–ER complex relative to that brought about by E_2 . Nevertheless, certain other estrogens (3-hydroxyestratriene, estratrien-17 β -ol and estradiol-17 α) active in gene regulation experiments also did not induce a shift in the DNA–ER complex. It can be concluded that, indeed, the conformational changes generated in the ERE–ER complex by 4-hydroxyestratrien-17 β -ol are different from those induced by E_2 . However, the changes in conformation highlighted by gel-shift assays are not necessarily related to the gene regulatory activity.

The downregulation of ER α has been attributed to a diminished steady state level of the receptor's mRNA brought about by estrogenic ligands when bound to ER α [4,13–16]. In the absence of an estrogen response element in the promoter of this gene, the initial autologous negative regulation of ER α is the result of the binding of receptor complex to the 5'-end of the coding sequence of the ER α gene, thereby suppressing the transcription rate [17,18]. Recovery of transcription from this negative regulation within 3 h [13], suggests that the prolonged depression of ER α mRNA results from a second, posttranscriptional event [13,16,17]. Therefore, the typical picture of ER α processing involves the rapid tight binding of cellular ER α to the nuclear compartment following the interaction of receptor with an estrogen ligand. Within 6 h, 50–60% of this nuclear-bound protein is seen to become undetectable by antibody or tritiated ligand assays (Fig. 7, Refs. [4,6,13–15]). This diminished receptor status remains for at least 48 h [13,15].

The described pattern is typical of most estrogenic ligands including 9 of the 10 estrogen analogues studied in these experiments (Table 1). An unusual pattern was elicited by 4-hydroxyestratrien-17 β -ol which did bring about the rapid tight nuclear binding of ER α . However, as the nuclear receptor decreased to levels as low as 15% of total ER α , cells treated with this A-ring isomer displayed a reappearance of ER α in the cytosol (Fig. 7(C)). Not only was this ligand unable to downregulate the early transcription of ER α mRNA (Fig. 3(E)), but the typical posttranscriptional depression of receptor was not apparent. Nevertheless, at a concentration commensurate with its binding affinity, this A-ring isomer was able to block the downregulation of ER α mRNA by E $_2$ (Fig. 4). Indicating that the ER α -4-hydroxyestratrien-17 β -ol complex formed an initial tight association within the nuclear compartment. This analogue, however, did not facilitate the posttranscriptional suppression of ER α noted by others [13,17].

Antiestrogens, such as 4-hydroxytamoxifen and ICI164,384 display yet another effect on ER α synthesis and processing. These ligands downregulate ER α mRNA in a concentration-dependent fashion after an 18 h incubation (Fig. 4). The downregulation of ER α mRNA by 4-hydroxytamoxifen after a period of 12 h has been seen previously [16]. In our experiments this late decrease in ER α mRNA (to 50% the level in untreated cells) is not reflected in receptor protein as detected by monoclonal antibody over a period of 24 h (Fig. 7(B)). This lack of processing of ER α following a pulse of 4-hydroxytamoxifen has been reported by Kaneko et al. [17] in Rat1 + ER cells. Others have seen partial processing of ER α by tamoxifen [14]. Therefore, although this antiestrogen brought about a downregulation of ER α mRNA, it had no effect on

the posttranscriptional process, resulting in its inability to decrease the levels of ER α in MCF-7 cells during 24 h of treatment (Figs. 4 and 7(B)).

The nature of the posttranscriptional process involved in the downregulation of ER α is presently unknown. A recent publication [35] suggests that phosphorylation events directed to the transactivation function-2 domain may play a role in receptor downregulation. In addition, a number of phosphorylation sites in the A/B region of ER α are known to be hormone sensitive. Phosphorylation of ER α potentiates its affinity for DNA [36] and treatment of the receptor with phosphatase diminished its DNA-binding activity [37]. Interestingly, 4-hydroxyestratrien-17 β -ol was the only estrogen analogue which has been shown in unpublished investigations in this laboratory to block the phosphorylation of ER α .

Results from these studies of ER α mRNA downregulation by estrogen ligands reveal specific ligand structural requirements which have not been characteristic of gene regulation via the binding of receptor complex to the ERE. For example, the estratriene nucleus which possesses a low affinity for ER α and was inactive in target gene regulation [3,5,6], displayed autologous negative regulation of ER mRNA at concentrations as low as 10⁻¹⁰ M (Fig. 2(B)). In addition, unlike the regulation of the progesterone receptor [6], pS2 [5], cathepsin D [5] and tissue plasminogen activator [3] genes, the placement of the D-ring hydroxyl group was critical to maximal downregulation of ER α mRNA (Fig. 6). In the absence of a hydroxyl group on carbon-17, the ligand was only capable of generating half-maximal depletion of ER α mRNA (Fig. 5). Whereas these same D-ring variations in the estrogen molecule generated ligands which stimulated the above target genes to a degree proportional to their affinity [3,5,6]. This information suggests that the conformational alterations elicited in the ER α molecule by the various ligands, impinge upon other transcription factors differently when the receptor functions via the response element directed mechanisms than they may during autologous negative regulation of the ER α gene.

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