Aromatase Expression in Women's Cancers

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Abstract

strogen has been positively linked to the pathogenesis and growth of three common women's cancers (breast, endometrium and ovary). A single gene encodes the key enzyme for estrogen ✔ biosynthesis named aromatase, inhibition of which effectively eliminates estrogen production in the entire body. Aromatase inhibitors successfully treat breast cancer, whereas their roles in endometrial and ovarian cancers are less clear. Ovary, testis, adipose tissue, skin, hypothalamus and placenta express aromatase normally, whereas breast, endometrial and ovarian cancers overexpress aromatase and produce local estrogen exerting paracrine and intracrine effects. Tissue specific promoters distributed over a 93 kilobase regulatory region upstream of a common coding region alternatively control aromatase expression. A distinct set of transcription factors regulates each promoter in a signaling pathway- and tissue-specific manner. In cancers of breast, endometrium and ovary, aromatase expression is primarly regulated by increased activity of the proximally located promoter I.3/II region. Promoters I.3 and II lie 215 bp from each other and are coordinately stimulated by PGE₂ via a cAMP-PKA-dependent pathway. In breast adipose fibroblasts exposed to PGE₂ secreted by malignant epithelial cells, activation of PKC potentiates cAMP-PKA-dependent induction of aromatase. Thus, inflammatory substances such as PGE₂ may play important roles in inducing local production of estrogen that promotes tumor growth.

Introduction

In this chapter, we will review the pathologic significance and regulation of aromatase in ovarian hormone-related women's cancers (breast, endometrium and ovary). We will first provide a relevant brief overview of physiologic and genetic concepts regarding aromatase expression in human tissues.

The Aromatase Enzyme

The aromatase enzyme is localized in the endoplasmic reticulum of estrogen-producing cells.^{1,2} Aromatase enzyme complex is comprised of two polypeptides. The first of these is a specific cytochrome P450, namely aromatase cytochrome P450 (the product of the *CYP19* gene).¹ The second is a flavoprotein, NADPH-cytochrome P450 reductase and is ubiquitously distributed in most cells. Thus, cell-specific expression of aromatase P450 (P450arom) determines the presence or absence of aromatase activity. For practical purposes, we will refer to "P450arom" as "aromatase" throughout this text. Since only a single gene (*CYP19*) encodes aromatase in mice and humans, targeted disruption of this gene or inhibition of its product effectively eliminates estrogen biosynthesis in these species.¹

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Innovative Endocrinology of Cancer, edited by Lev M. Berstein and Richard J. Santen. ©2008 Landes Bioscience and Springer Science+Business Media.

In the human, aromatase is expressed in a number of cells including the ovarian granulosa cell, the placental syncytiotrophoblast, the testicular Leydig cell, as well as various extraglandular sites including the brain and skin fibroblasts.³ The principal product of the ovarian granulosa cells during the follicular phase is estradiol. Additionally, aromatase is expressed in human adipose tissue. Whereas the highest levels of aromatase are in the ovarian granulosa cells in premenopausal women, the adipose tissue becomes the major aromatase expressing body site after menopause (Fig. 1).⁴⁵ Although aromatase level per adipose tissue fibroblast may be small, the sum of estrogen arising from billions of adipose tissue fibroblasts in the entire body makes a physiologic impact. In adipose tissue, estrogenically weak estrone is produced from androstenedione of adrenal origin in relatively large quantities. However, at least half of this peripherally produced estrone is eventually converted to estradiol in extraovarian tissues (Fig. 1).⁶

The CYP19 (Aromatase) Gene

The aromatase gene is transcribed from the telomere to the centromere and the region encoding the aromatase protein spans 30 kb of the 3'-end and contains 9 exons (II-X).⁷ The ATG translation start site is located in coding exon II. The upstream (telomeric) 93 kb of the gene contains a number of promoters.^{2,3} The most proximal gonadspecific promoter II and two other proximal

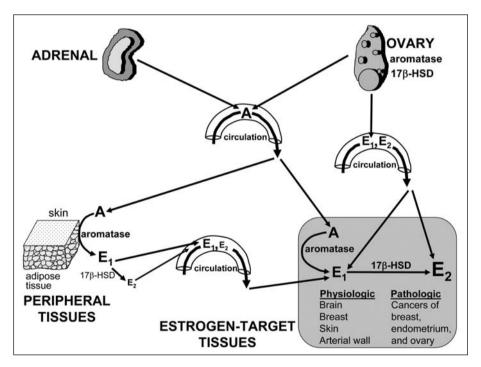


Figure 1. Tissue sites of estrogen production in women. The biologically active estrogen, estradiol (E_2) is produced at least in three major sites: 1-direct secretion from the ovary in reproductive age women; 2-by conversion of circulating androstenedione (A) of adrenal and/or ovarian origins to estrone (E_1) in peripheral tissues; and 3-by conversion of A to E_1 in estrogen target tissues. In the latter two instances, estrogenically weak E_1 is further converted to E_2 within the same tissue. The presence of the enzyme aromatase and 17 β hydroxysteroid dehydrogenase (17 β -HSD) is critical for E_2 formation at these sites. E_2 formation by peripheral and local conversion is particularly important for postmenopausal women and for estrogen-dependent diseases such as breast cancer, endometriosis and endometrial cancer. Figure used with permission from Bulun SE, Lin Z, Imir G. et al. Pharmacol Rev 2005; 57:359-383.¹⁵

promoters, I.3 (expressed in adipose tissue and breast cancer) and I.6 (expressed in bone) are found to be located within the 1-kb region upstream of the ATG translation start site in exon II, as expected (Fig. 2). Promoter I.2, the minor placenta-specific promoter, is located approximately 13 kb upstream of the ATG site in exon II. The promoters specific for the brain (I.f.), endothelial cells (I.7), fetal tissues (I.5), adipose tissue (I.4) and placenta (2a and I.1) are localized in tandem order at ~3, 36, 43, 73, 78 and 93-kb, respectively, upstream of the first coding exon, the exon II (Fig. 2).^{2,8} In addition to promoter II specific sequences, transcripts containing two other unique sequences, untranslated exons I.3 and I.4, are present in adipose tissue and in adipose tissue fibroblasts maintained in culture.⁸ Transcription initiated by use of each promoter gives rise to a transcript with a unique 5'-untranslated end that contains the sequence encoded in the first exon immediately downstream of this particular promoter (Fig. 2). Therefore, the 5'-untranslated region of aromatase mRNA is promoter specific and may be viewed as a signature of the particular promoter used. It should be emphasized again that all of these 5'-ends are spliced onto a common junction 38 bp upstream of the ATG translation start site.⁸ Consequently, the sequence encoding the open reading frame is identical in each case. Thus, the expressed protein is the same regardless of the splicing pattern (Fig. 2).

Normal Hormonal Pathways that Regulate Aromatase Expression

The primary site of aromatase expression in premenopausal women is the ovarian follicle, where FSH induces aromatase and thus estradiol production in a cyclic fashion.³ Ovarian aromatase expression is mediated primarily by FSH receptors, cAMP production and activation of the proximal promoter II (Fig. 3).³ Men and postmenopausal women also produce estrogen by aromatase that resides in extragonadal tissues such as adipose tissue and skin (Fig. 3).³ Estrogen produced in these extragonadal tissues are of paramount importance for the closure of bone plates

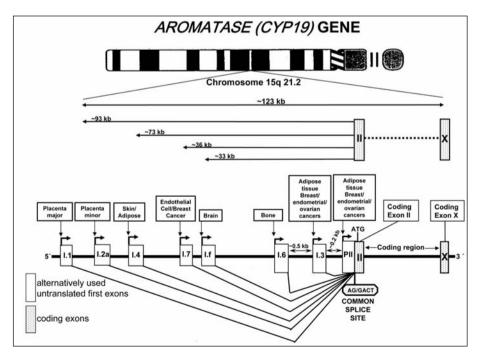


Figure 2. CYP19 (aromatase) gene. Expression of the aromatase gene is regulated by the tissue-specific activation of a number of promoters via alternative splicing. Figure used with permission from: Bulun SE, Lin Z, Imir G et al. Pharmacol Rev 2005; 57:359-383.¹⁵

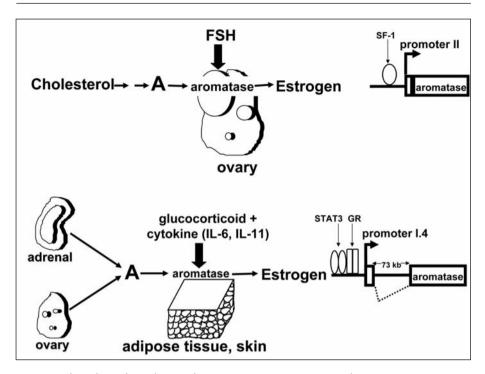


Figure 3. Physiological regulation of aromatase expression. FSH induces aromatase expression via through a cAMP-dependent pathway in ovarian granulosa cells via promoter II. Steroidogenic factor-I (SF-1) mediates this action of FSH. On the other hand, a combination of a glucocorticoid and a member of the class I cytokine family induces aromatase expression in skin and adipose tissue fibroblasts via promoter I.4 located 70 kb upstream of the coding region. Binding of signal transducers and activators of transcription (STAT)-3 and glucocorticoid receptor (GR) upstream of promoter I.4 mediate regulation of aromatase expression in these fibroblasts. Figure used with permission from: Bulun SE, Lin Z, Imir G et al. Pharmacol Rev 2005; 57:359-383.¹⁵

and bone mineralization in both men and postmenopausal women, since the phenotype of men with defective genes of aromatase or estrogen receptor- α include severe osteoporosis and extremely tall stature with growth into adulthood.⁹ A distal promoter (I.4) located 73 kilobases upstream of the coding region directs as was mentioned aromatase expression in adipose tissue and skin fibroblasts. Promoter I.4 in these tissues is regulated by combined action of a glucocorticoid and a member of the class I cytokine family [e.g., interleukin (IL)-6, IL-11, leukemia inhibitory factor (LIF), oncostatin-M] (Fig. 3).¹⁰

The alternative use of promoters comprises the basis for differential regulation of aromatase expression by various hormones, growth factors and cytokines in a tissue specific manner. For example, extremely high baseline levels of the placental promoter I.1 activity are maintained constitutively in the syncytiotrophoblast and a consequence of decreasing levels of inhibitory transcription factors as cytotrophoblasts differentiate to a syncytiotrophoblast.^{11,12} On the other hand, extremely low baseline levels of promoter II in the ovary are stimulated strikingly by FSH via a cAMP dependent pathway in the developing follicle (Fig. 3).³ Serum, cytokines and growth factors are inhibitory to promoter II. In case of adipose and skin fibroblasts, promoter I.4 is used in vivo and activated as was said above coordinately by a glucocorticoid in the presence of a cytokine (IL-6, IL-11, LIF, oncostatin M). Glucocorticoid receptors and the Jak-1/STAT-3 pathway mediate this induction.¹⁰

Promoter use in cultured adipose tissue fibroblasts is a function of hormonal treatments. For example, in vitro studies showed that PGE_2 or cAMP analogs stimulate aromatase expression strikingly via proximally located promoters II and I.3, whereas treatment with a glucocorticoid plus a member of the class I cytokine family switches promoter use to I.4.^{10,13}

Pathological Expression of Aromatase in Women's Cancers

Breast and endometrial cancers are highly responsive to estrogen for growth evident by high concentrations of estrogen receptors in these tissues.¹⁴ Malignant breast and endometrial tumors also produce large amounts of estrogen locally via overexpressing aromatase compared to their normal counterparts.¹⁵ In particular, aromatase overexpression in breast cancer tissue has been shown to be critical, since the use of aromatase inhibitors is clearly therapeutic in breast cancer. Aromatase is also overexpressed in endometrial cancer.¹⁶ Although preliminary trials showed promising results, the therapeutic role of aromatase inhibitors in endometrial cancer is not as clear yet.^{17,18}

Experimental and epidemiological evidence suggest that estrogen and progesterone are implicated in ovarian carcinogenesis. New data have indicated that estrogen favors neoplastic transformation of the ovarian surface epithelium while progesterone offers protection against ovarian cancer development.¹⁹⁻²³ Since a subset of ovarian cancers was linked to endometriosis and, aromatase is a key molecular target in endometriosis, aromatase expression in ovarian cancer may also be targeted for treatment in selected patients.¹⁵ In fact, recent pilot studies employing aromatase inhibitors have shown various degrees of clinical benefit for patients with advanced stages of ovarian cancer.²⁴⁻²⁷

Aromatase and Breast Cancer

Paracrine interactions between malignant breast epithelial cells, proximal adipose fibroblasts and vascular endothelial cells are responsible for estrogen biosynthesis and lack of adipogenic differentiation in breast cancer tissue. It appears that malignant epithelial cells secrete factors that inhibit the differentiation of surrounding adipose fibroblasts to mature adipocytes and also stimulate aromatase expression in these undifferentiated adipose fibroblasts.²⁸ The in vivo presence of malignant epithelial cells also enhances aromatase expression in endothelial cells in breast tissue.²⁹ We developed a model in breast cancer, which reconciles the inhibition of adipogenic differentiation and estrogen biosynthesis in a positive feedback cycle.

The desmoplastic reaction (formation of the dense fibroblast layer surrounding malignant epithelial cells) is essential for structural and biochemical support for tumor growth. In fact, the some pathologists refer to 70% of breast carcinomas as "scirrhous" type indicating the rock-like consistency of these tumors.³⁰ This consistency comes from the tightly packed undifferentiated adipose fibroblasts around malignant epithelial cells. Malignant epithelial cells achieve this by secreting large quantities of TNF and IL-11 that inhibit the differentiation of fibroblasts to mature adipocytes. Thus, large numbers of these estrogen producing cells are maintained proximal to malignant cells.^{28,29} At the same time, a separate set of factors secreted by malignant epithelial cells activates aromatase expression in surrounding adipose fibroblasts.^{28,29}

Malignant epithelial cells induce aromatase via activation of aberrant promoters in breast cancer tissue and adipose fibroblasts proximal to tumor (Fig. 4). The breast adipose tissue in disease-free women maintains low levels of aromatase expression primarily via promoter I.4 that lies 73 kb upstream of the common coding region (Fig. 4).⁸ The proximal promoters I.3 and II are used only minimally in normal breast adipose tissue.⁸ Transcription via activity of promoters II and I.3 in the breast tumor fibroblasts and malignant epithelial cells, however, is strikingly increased (Fig. 4).³¹⁻³⁴ Additionally, the endothelial-type promoter I.7 is also upregulated in breast tumor tissue (Fig. 4).³⁵ Thus, it appears that the prototype estrogen-dependent malignancy (breast cancer) takes advantage of four promoters utilized in various cell types for aromatase expression (Fig. 4). The sum of aromatase mRNA species arising from these four promoters markedly increase total aromatase mRNA levels in breast cancer compared with the normal breast that uses almost exclusively promoter I.4 (Fig. 4).

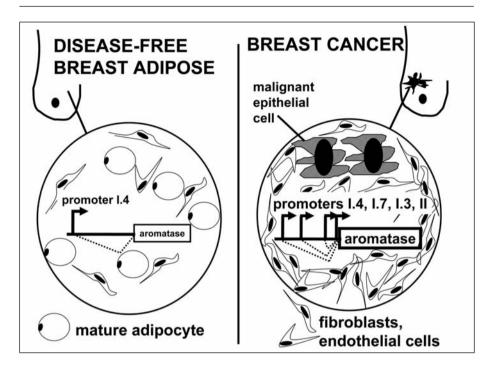


Figure 4. Promoter use for aromatase expression in normal and malignant breast tissues. The levels of total aromatase mRNA levels in breast cancer tissue are strikingly higher than normal breast tissue. The normal breast adipose tissue maintains low levels of aromatase expression primarily via promoter 1.4. Promoters 1.3 and 11 are used only minimally in normal breast adipose tissue. Promoter II and 1.3 activities in the breast cancer, however, are strikingly increased. Additionally, the endothelialtype promoter 1.7 is also upregulated in breast cancer. Thus, it appears that the prototype estrogendependent malignancy (breast cancer) takes advantage of four promoters (II, 1.3, 1.7 and 1.4) for aromatase expression. The sum of aromatase mRNA levels in breast cancer compared with the normal breast. Figure used with permission from: Bulun SE, Lin Z, Imir G et al Pharmacol Rev 2005; 57:359-383.¹⁵

Tissue Origins of Estrogen in Postmenopausal Women with Breast Cancer

Estrogen arises from two sources in a postmenopausal woman with breast cancer. First, estrogen that arises from extraovarian body sites such as subcutaneous adipose tissue and skin reaches breast cancer by way of circulation in an endocrine manner. Second, estrogen locally produced in breast cancer tissue makes an impact via paracrine or intracrine mechanisms.

Endocrine Effect

Aromatase in adipose tissue and skin. The potential roles of extraovarian aromatase in human physiology and pathology were also recognized initially in the 1960s.³⁶ These studies demonstrated that the conversion rate of plasma androstenedione to estrone in humans increased as a function of obesity and aging.^{5,37} These same studies also revealed the importance of extraovarian tissues (primarily adipose tissue and skin) as the origin of estrogen in postmenopausal women (see Fig. 1). Extraovarian estrogen formation was shown to be correlated positively with excess body weight in both pre and postmenopausal women and may be increased as much as ten-fold in morbidly obese postmenopausal women.^{5,37} This elevation in association with both obesity and aging bears a striking relationship to the incidence of endometrial hyperplasia and cancer, which are more commonly observed in elderly obese women.³⁸ It is now recognized that the

continuous production of estrogen by the adipose tissue in these women is one of the risk factors of endometrial hyperplasia and cancer.

Evidence is also suggestive of a role of estrogens produced by adipose tissue in the pathogenesis of the breast cancer. For example, breast cancer incidence correlates positively with the body fat content or serum estradiol levels in postmenopausal women suggesting that estrogens collectively produced in all extraovarian sites reach the breast tissue by circulation in an endocrine fashion and stimulate tumor growth.^{39,40} A role for adipose tissue estrogen biosynthesis in promoting the growth of breast cancer is implied because of the palliative effects of adrenalectomy in the past. As estrone production by adipose tissue is dependent on plasma androstenedione secreted by the adrenal cortex as substrate, the role of adrenalectomy is explicable in terms of the denial of substrate precursor for adipose tissue estrogen biosynthesis. Today, reduction of estrogen biosynthesis in the adipose tissue is accomplished by the use of aromatase inhibitors in the treatment of breast cancer (Fig. 5).^{41,42}

Paracrine/Intracrine Effect

Local aromatase in breast cancer tissue. Mechanisms giving rise to increased local concentrations of estrogen in breast cancer via aromatase overexpression within the tumor tissue has been demonstrated by a number investigators.^{43,45} These studies showed strikingly increased local levels of estrone, estrone sulfate and eastradiol in breast tumor tissue compared with circulating estrogen levels.^{43,45}

A series of paracrine interactions between malignant breast epithelial cells and surrounding adipose stroma were uncovered and explained increased local estrogen levels in the breast bearing a cancer. For example, independent studies from at least six different laboratories were indicative of striking increases in aromatase enzyme activity and mRNA levels in breast fat adjacent to cancer compared with those in distal fat or disease free breast adipose tissue (Fig. 5).^{32-34,46-49} We also found that an elevation in aromatase expression in adipose stroma surrounding malignant epithelial cells is regulated by complex cellular, molecular and genomic mechanisms.^{29,31,47} Interestingly, the overall aromatase expression in breast adipose tissue in mastectomy specimens bearing a breast tumor was significantly higher than that in benign breast tissue removed for reduction mammoplasty.^{31,50,51}

Estrogens can act both directly or indirectly on human breast cancer cells to promote proliferation. Breast cancer cells in culture elaborate a number of growth stimulants in response to estrogen, which can act in an autocrine and paracrine manner to stimulate their growth. However, there is also evidence that estrogens can directly induce proliferation of breast cancer cells.

The pathologic significance of local aromatase activity in breast cancer was recognized based on the following in vitro data. MCF-7 breast cancer cells, which were stably transfected to express an MMTV-promoter-driven human aromatase cDNA and inoculated into oophorectomized nude mice, remained dependent on circulating androstenedione for their rapid growth.⁵² Another evidence for the importance of local aromatase expression in the breast tissue came from an in vivo mouse model demonstrating that aromatase overexpression in breast tissue is sufficient for maintaining hyperplasia in the absence of circulating estrogen and that aromatase inhibitors abrogated hyperplasia.⁵³ These transgenic mice with MMTV-promoter-driven local aromatase in breast tissue are more prone for breast cancer development.⁵⁴

Role of Adipose Tissue in Aromatase Expression in Malignant Breast Tumor Tissue

Cell Types That Express Aromatase in Breast Cancer

Breast adipose tissue is primarily comprised of mature lipid containing cells and other stromal elements. This latter group of cells in the breast adipose tissue was characterized using immunohistochemical methods.⁵⁵ Ninety percent of these resident cells of adipose tissue are fibroblasts, i.e., the potential precursors of mature adipocytes and another 7% represented by endothelial cells. The majority (80-90%) of aromatase transcripts in adipose tissue was demonstrated to reside in fibroblasts compared with mature adipocytes.⁵⁵ Moreover, aromatase enzyme activity was found to reside primarily in the fibroblast component of the adipose tissue in a previous study from the same laboratory.⁵⁶

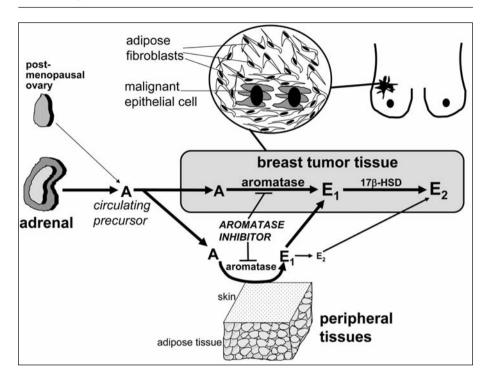


Figure 5. Origins of estrogen in postmenopausal breast cancer. This figure exemplifies the important pathologic roles of extraovarian (peripheral) and local estrogen biosynthesis in an estrogen-dependent disease in postmenopausal women. The estrogen precursor androstenedione (A) originates primarily from the adrenal in the postmenopausal woman. Aromatase expression and enzyme activity in extraovarian tissues such as fat increases with advancing age. The aromatase activity in skin and subcutaneous adipose fibroblasts gives rise to formation of systemically available estrone (E₁) and to a smaller extent estradiol (E₂). The conversion of circulating A to E₁ in undifferentiated breast adipose fibroblasts compacted around malignant epithelial cells and subsequent conversion of E₁ to E₂ in malignant epithelial cells provide high tissue concentrations of E₂ for tumor growth. The clinical relevance of these findings is exemplified by the successful use of aromatase inhibitors to treat breast cancer. Figure used with permission from: Bulun SE, Lin Z, Imir G et al. Pharmacol Rev 2005; 57:359-383.¹⁵

Immunoreactive aromatase was localized to both the malignant epithelial cells and surrounding fibroblasts in breast tumor tissues.⁵⁷⁻⁵⁹ Different antibodies, however, showed variable affinity to malignant epithelial cells vs fibroblasts.⁶⁰ Immunoreactive aromatase was also observed in endothelial cells in normal breast tissue and breast tumors. Recently published data using RNA from lasercaptured breast tumor cells showed aromatase mRNA in both stromal and malignant epithelial cells in tumor tissues from 3 patients.⁶⁰ Markedly high levels of aromatase enzyme activity have been consistently detected in breast adipose fibroblasts freshly isolated from breast tissue with or without cancer.^{55,56} Aromatase enzyme activity in malignant breast epithelial cells, on the other hand, in experimental conditions is either undetectable or extremely low.⁶¹

Adjacent adipose tissue including the dense fibroblast layer seems to account for the majority of aromatase expression in breast tumors for the following reasons. First, the quantity of adjacent adipose tissue surrounding a clinically detectable breast tumor is comparatively very large, e.g., the volume of adipose tissue within 1-inch radius of a 1 ml breast tumor is 129 ml. Second, the most intense aromatase immunostaining was observed in the adipose tissue fibroblasts located in and around the fibrous capsule (i.e., desmoplastic reaction) surrounding malignant cells.⁴⁸

Third, levels of aromatase expression and activity in fibroblasts isolated from breast adipose tissue or tumor are 10-15 times higher than those found in malignant epithelial cells or cell lines.⁶¹

Impaired Adipogenic Differentiation in Breast Cancer

Link to aromatase overexpression. Extraordinarily large quantities of TNF and IL-11 are produced and secreted by malignant breast epithelial cells²⁸ (Fig. 6).²⁸ These two cytokines mediate one of the most commonly observed biological phenomena in breast tumors, namely the desmoplastic reaction. Desmoplastic reaction or accumulation of fibroblasts around malignant epithelial cells serves to maintain the strikingly hard consistency in many of these tumors (i.e., the traditional macroscopic description of malignant breast tumors as "scirrhous cancer") and increased local concentrations of estrogen via aromatase overexpression localized to these undifferentiated fibroblasts. The inhibition of differentiation of fibroblasts to mature adipocytes mediated by TNF and IL-11 is the key event responsible for desmoplastic reaction, because neither malignant cell-conditioned media nor these cytokines caused the proliferation of adipose tissue fibroblasts.²⁸ Moreover, blocking both TNF and IL-11 in cancer cell-CM conditioned medium using neutralizing antibodies is sufficient to reverse this antidifferentiative effect of cancer cells completely (Fig. 6).²⁸ In summary, desmoplastic reaction primarily occurs via the action of cytokines (TNF and IL-11) secreted by the malignant epithelial cells to inhibit the differentiation of adipose tissue fibroblasts to mature adipocytes. This tumor-induced block in adipocyte differentiation is mediated by the selective inhibition of expression of the essential adipogenic transcription factors, namely, C/EBPa and PPARy (Fig. 6).28

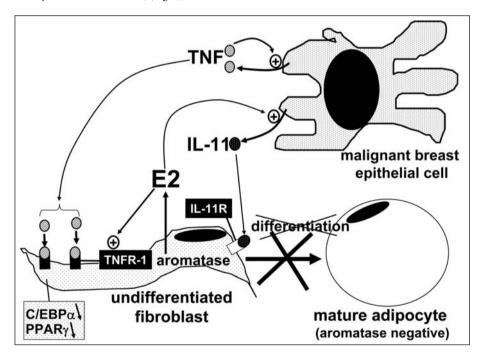


Figure 6. Detail of epithelial-stromal interaction via estrogen and cytokines in breast cancer. Estradiol (E_2) increases secretion of antiadipogenic cytokines (IL-11) from malignant epithelial cells and upregulates their antiadipogenic-type receptors (TNF-receptor type 1, TNFR1) in undifferentiated fibroblasts. These redundant mechanisms give rise to accumulation of undifferentiated fibroblasts around malignant epithelial cells (desmoplastic reaction), which express aromatase and form E_2 . Figure used with permission from: Bulun SE, Lin Z, Imir G et al. Pharmacol Rev 2005; 57:359-383.¹⁵

Estrogen per se seems to potentiate this antiadipogenic action via indirect mechanisms. For example, treatment of T47D breast cancer cells with estradiol increased the mRNA levels of IL-11 by 3-fold.⁶² Moreover, the cellular actions of TNF are mediated by two distinct receptors TNFR1 (also known as p60 in humans) and TNFR2 (p80). TNFR1 but not TNFR2 was found to be responsible for the inhibition of adipocyte differentiation using mutants of TNF specific for the stimulation of either receptor type.⁶³ We recently demonstrated that TNFR1 is responsible for inhibition of adipocyte differentiation in breast cancer.⁶⁴ Interestingly, estradiol enhances this anti-adipogenic effect by inducing TNFR1 levels in adipose fibroblasts (Fig. 6).⁶⁴

Thus, large amounts of antiadipogenic cytokines (e.g., TNF and IL-11) secreted by malignant epithelial cells serve to maintain increased numbers of the aromatase-expressing cell type, i.e., undifferentiated adipose fibroblast, in breast tumor tissue. This is further enhanced by stimulatory effects of estrogen on IL-11 production in cancer cells and on the TNF receptor type that mediates adipogenic inhibition (Fig. 6).

Transcriptional Mechanisms Responsible for Elevated Aromatase Expression in Breast Cancer

Alternative promoter use is a major mechanism that mediates increased aromatase expression in breast cancer. The normal breast adipose tissue maintains low levels of aromatase expression primarily via promoter I.4 that lies 73 kb upstream of the common coding region (see Fig. 4). The proximally located promoters I.3 and II are used only minimally in normal breast adipose tissue. Promoter II and I.3 activities in the breast cancer, however, are strikingly increased.³¹⁻³⁴ Additionally, the endothelial-type promoter I.7 is also upregulated in breast cancer.³⁵ Thus, as was mentioned above it appears that the prototype estrogen-dependent malignancy breast cancer takes advantage of four promoters (II, I.3, I.7 and I.4) for aromatase expression (see Fig. 4). The sum of aromatase mRNA species arising from these four promoters markedly increase total aromatase mRNA levels in breast cancer compared with the normal breast that uses almost exclusively promoter I.4 (see Fig. 4)

Upregulation of Promoters I.3 and II

Using an in vivo approach, we and two other groups demonstrated by quantitative exon specific RT-PCR that the use of the proximal promoters II/I.3 are strikingly upregulated in adipose tissue adjacent to breast cancer and in breast cancer tissue per se.^{31,33,34} As noted earlier, promoters II and I.3 are located within 215 bp from each other and are coordinately induced by cAMP-dependent or independent mechanisms in adipose fibroblasts in breast tumors. These promoters possibly share common regulatory DNA motifs.

Increased promoter I.3/II activity is in part, the basis for increased aromatase expression in peri-and intratumoral adipose tissue fibroblasts (see Fig. 4).³¹ Over the past several years, others and we sought to elucidate the mechanisms underlying this cancer-induced increase in promoters I.3/II activity in adipose tissue fibroblasts.

 PGE_2 induces aromatase via promoters I.3 and II employing both cAMP/PKA- and PKC-dependent pathways.¹³ In this model, PGE_2 stimulates binding activity of an orphan nuclear receptor LRH-1 to a nuclear receptor half-site (-136/-124 bp) upstream of promoter II.^{65,66} Treatment with PGE_2 strikingly increased both LRH-1 expression and its binding activity to the aromatase promoter II in cultured adipose fibroblasts. LRH-1 overexpression significantly increased aromatase promoter II activity and aromatase enzyme activity in cultured adipose fibroblasts (Fig. 7). From an in vivo perspective, LRH-1 was upregulated in undifferentiated fibroblasts in breast tumor tissue compared with those in disease-free breast tissue. The recently reported increases in COX-2 expression and beneficial effects of nonsteroidal anti-inflammatory in breast cancer support this model.^{67,68} In vivo evidence for increased PGE₂ formation in breast cancer, however, is still lacking at this time.

In addition to the LRH-1-binding site, we isolated two *cis*-acting elements that conferred stimulation of promoter I.3/II use in response to cAMP plus or minus PKC activation, the signaling pathways induced by $PGE_2^{29,69}$ Two critical elements were determined as a C/EBP site

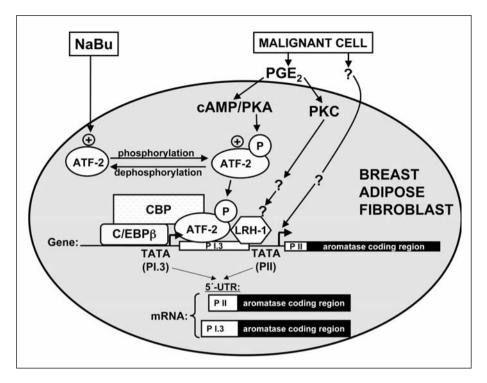


Figure 7. Effects of PGE₂, PKA, PKC and sodium butyrate (NaBu) on aromatase expression in breast adipose fibroblasts. We recently found that NaBu profoundly decreased promoter 1.3/ II-specific aromatase mRNA expression induced by PGE₂ or a surrogate hormomonal cocktail made of dibutyryl cAMP (Bt₂cAMP) plus the PKC activator phorbol diacetate (PDA). MCM, Bt₂cAMP+PDA or NaBu regulated aromatase mRNA levels or enzyme activity only specifically via promoters 1.3/II but not other promoters. Recruitment of phosphorylated ATF-2 by a CRE (-211/-199) in the promoter 1.3/II region conferred the response to malignant epithelial cells conditioned medium, PGE₂ or Bt₂cAMP+PDA. Malignant cell-conditioned medium, PGE₂ or Bt₂cAMP+PDA stabilized a complex comprised of phosphorylated ATF-2, C/EBPß and CBP in the common regulatory region of promoters I.3/II. The inhibitory effect of NaBu on transcription was not accompanied by comparable changes in overall histone acetylation patterns of promoters I.3/II. NaBu treatment, however, consistently decreased ATF-2 phosphorylation and disrupted the activating complex. Taken together, these findings represent a novel mechanism of NaBu action and provide evidence that aromatase activity can be attenuated in a signaling pathway and tissue specific fashion. Our data also suggested that malignant cells secreted substances other than PGE2. These unknown substances were associated with signaling pathways other than cAMP-PKA in the activation of aromatase promoters I.3 and II.

(-317/-304) and a CRE (-211/-197), since mutation of either element abolished cAMP-induced promoter activity (Fig. 7).⁶⁹

PGE2 or a cAMP analog (plus or minus PKC activator) induced phosphorylation of ATF-2 and its binding to the CRE (Fig. 7).⁶⁹ This CRE is occupied by nonphosphorylated ATF-2 in fibroblasts treated with benign epithelial cell-conditioned medium associated with inactivation of promoters I.3 and II. Moreover, chromatin immunoprecipitation PCR showed that the activator transcriptional complex in a malignant environment contains C/EBP β , phosphorylated ATF-2 and p300/CBP, whereas the inactivator complex (benign environment) contained nonphosphorylated ATF-2 (Fig. 7).⁶⁹ We recently found that sodium-butyrate (NaBu) profoundly decreased cAMP analog (plus or minus PKC activator)-induced promoter I.3/II-specific aromatase mRNA expression. A cAMP analog (plus or minus a PKC activator) or NaBu regulated aromatase mRNA levels or enzyme activity only via promoters I.3/II but not promoters I.1 or I.4 in breast, ovarian, placental and hepatic cells. Mechanistically, recruitment of phosphorylated ATF2 by a CRE (-211/-199) in the promoter I.3/II region conferred the response to a cAMP analog (plus or minus PKC activator). Treatment with a cAMP analog (plus or minus PKC activator) stabilized a complex comprised of phosphorylated ATF-2, C/EBP β and CBP in the common regulatory region of promoters I.3/II (Fig. 7). The inhibitory effect of NaBu on transcription was not accompanied by comparable changes in overall histone acetylation patterns of promoters I.3/II. NaBu treatment, however, consistently decreased ATF-2 phosphorylation and disrupted the activating complex (Fig. 7). Together, these findings represent a novel mechanism of NaBu action and provide evidence that aromatase activity can be attenuated in a signaling pathway- and tissue-specific fashion.⁶⁹

In an alternative experimental model, we found that conditioned medium from malignant breast epithelial cells (MCF-7 or T47D) markedly induced aromatase expression in adipose tissue fibroblasts via promoters I.3 and II (Fig. 7).^{29,69} Malignant epithelial cell-conditioned medium also induced phosphorylation of ATF-2 and its binding activity to the promoter I.3/II region. As in the case of treatment with a cAMP analog (plus or minus a PKC activator), incubation of breast adipose fibroblasts with malignant epithelial cell-conditioned medium stabilized a complex made of phosphorylated ATF-2, C/EBP β and CBP at the *aromatase* promoter I.3/II region.⁶⁹ NaBu also disrupts this transcriptional complex assembled in response to malignant epithelial cell conditioned medium.⁶⁹ We hypothesize that a hormonal cocktail secreted from malignant epithelial cells induces aromatase in undifferentiated adipose fibroblasts via redundant pathways. Although PGE₂ seems to be an important component of this cocktail, other substances can also induce aromatase expression via promoters I.3/II.²⁹ For example, the addition of a COX2 inhibitor or an adenylyl cyclase inhibitor does not reverse induction of aromatase expression by malignant epithelial cellconditioned medium.²⁹

A unified model for promoter II/I.3 activation in breast cancer therefore predicts that malignant epithelial cells secrete a number of factors including PGE_2 (Fig. 7). These factors induce a number of signaling pathways in a redundant fashion to activate the transcription of the aromatase gene via promoter I.3/II in adipose fibroblasts. PGE_2 possibly arising from malignant epithelial cells is a candidate factor for activation of promoters I.3 and II in breast cancer. This, however, has not been demonstrated in vivo. Neither PGE_2 nor its downstream regulators cAMP or LRH-1-binding site in promoter II were found to be essential for activation of promoters II in adipose fibroblasts treated with malignant cell-conditioned medium.²⁹ In disease-free breast tissue, incorporation of a number of transcriptional repressors into the multimeric complex that occupies promoter I.3/II region is associated with inhibition of transcription. Malignant epithelial cell conditioned medium, on the other hand, gives rise to replacement of this inhibitory complex by an activator transcriptional complex comprised of distinct factors such as phosphorylated ATF-2, C/EBP β , p300/CBP and possibly LRH-1 (Fig. 7).⁶⁹

In summary, the proximal promoters II and I.3 clustered within a 215 bp region are coordinately regulated. They remain quiescent in fibroblasts of normal breast tissue via redundant binding of multiple transcriptional inhibitors (Fig. 7). In a malignant breast environment, however, these promoter regions are occupied by multiple transcriptional enhancers as a result of activation of multiple signaling pathways in a fail-safe fashion to increase aromatase expression in breast fibroblasts (Fig. 7).

Regulation of Promoters II and I.3 in MCF-7 Cells

A group of investigators studied the regulation of a number of gene reporter constructs of the promoter II/I.3 region.⁷⁰ They found that ERR alpha-1 and CREB1 upregulate and EAR2, COUP-TFI, RAR gamma, Snail and Slug proteins downregulate this promoter region.⁷⁰ The in vivo relevance of these findings will become clearer in future, once the relative significance of aromatase enzyme activity and estrogen biosynthesis is demonstrated in malignant epithelial cells.

Upregulation of Promoter I.7 in Breast Cancer

Studies summarized above employed exon specific RT-PCR analysis of 5'-untranslated ends of aromatase mRNA in breast cancer tissues. This limited strategy permitted only the detection of promoters previously identified from healthy tissues.^{31,33,34} Discovery-driven approaches designed to identify novel promoter regions in breast cancer or adjacent adjpose tissues, however, have not been published until recently. To identify novel promoter regions in cancer tissues and proximal fat, we employed the 5'-rapid amplification of cDNA ends (RACE) procedure using total RNA isolated from breast cancer and proximal adipose tissue samples. We cloned a novel 101bp untranslated first exon (I.7) that comprises the 5'-end of 29-54% of aromatase mRNA isolated from breast cancer tissues.³⁵ The levels of aromatase mRNA with exon I.7 were significantly increased in breast cancer tissues and adipose tissue adjacent to tumors. We identified a promoter immediately upstream of exon I.7 and mapped this to about 36 kb upstream of ATG translation start site of the aromatase gene.³⁵ Promoter I.7 is a TATA-less promoter containing *cis*-regulatory elements found in megakaryocytic and endothelial type promoters. Maximal promoter activity could be demonstrated in human microvascular endothelial cells. Binding of the transcription factor GATA-2-to a specific GATA cisregulatory element in this promoter was critical for its regulation in endothelial cells.³⁵ In conclusion, promoter I.7 is a GATA-2-regulated endothelial type promoter of the human aromatase gene and may increase estrogen biosynthesis in vascular endothelial cells of breast cancer. The activity of this promoter may also be important for intracrine and paracrine effects of estrogen on blood vessel physiology.

Summary of Regulation of Aromatase Expression in Breast Cancer

Several alternative cellular and molecular mechanisms serve to maintain excessive levels of aromatase activity in breast stroma proximal to malignant epithelial cells. First, malignant epithelial cell derived factors induce aromatase over expression via the transcription factors LRH-1, C/EBPβ and phosphorylated ATF-2 (see Fig. 5). These factors are incorporated into a multimeric transcriptional complex that occupies the aromatase promoter I.3/II region in adipose tissue fibroblasts adjacent to epithelial cells. Second, aromatase is overexpressed in vascular endothelial cells of tumor tissue via binding of GATA2 and other endothelial type transcription factors to promoter I.7. These factors may also mediate angiogenesis in tumor tissue. Moreover, estrogen is known to induce the angiogenic factor VEGF in cancer cells (see Fig. 5).⁷¹⁻⁷⁴ Third, we demonstrated recently that antiadipogenic cytokines IL-11 and TNF secreted by malignant epithelial cells block the differentiation of the aromatase-expressing cells (fibroblasts) to mature adipocytes that do not express aromatase (see Fig. 5). Thereby, these cytokines secreted abundantly by malignant epithelial cells serve to maintain a dense layer of aromatase-expressing fibroblasts proximal to malignant epithelial cells to provide structural and hormonal support. Fourth, the expression of IL-11 in malignant epithelial cells and antiadipogenic type TNF receptors in adjacent adipose tissue fibroblasts are upregulated by estrogen produced as a consequence of elevated aromatase activity in breast tumors. This positive feedback involving complex epithelial-stromal interactions favor higher numbers of undifferentiated fibroblasts, angiogenesis and increased local estrogen concentrations in breast tumors (see Fig. 5). These four mechanisms interact to maintain high levels of estrogen production in a breast tumor.74

Aromatase Inhibitors in the Treatment of Breast Cancer

Today, aromatase inhibitors are the most effective endocrine-treatment of estrogen-responsive breast cancer (Fig. 7).⁷⁵ Six recent head to head randomized clinical trials published since 2000 demonstrated the superiority of aromatase inhibitors to tamoxifen in the treatment of breast cancer.⁷⁶⁻⁸³ Longterm side effect profiles of these agents will determine whether aromatase inhibitors will replace tamoxifen or other selective estrogen receptor modulators in the long run.

There are two intriguing implications of these results. First, it is pharmacologically more efficacious to block estrogen formation rather than its action at least by currently approved estrogen antagonists or SERMs. Second, the local effect of aromatase inhibitors at the target tissue level to block local estrogen formation possibly represents the most critical mechanism for the superior therapeutic potential of aromatase inhibitors (Fig. 7).

Targeting aromatase in breast cancer as a therapeutic strategy was first conceptualized in the 1960s.⁷⁵ Aminoglutethimide was the first aromatase inhibitor tested for this purpose. Although the first generation aromatase inhibitor aminoglutethimide was as efficacious as tamoxifen in the treatment of breast cancer, its adverse side effects precluded its widespread use.⁷⁵ Tamoxifen was introduced in the 1970s and became the gold standard for hormonal treatment of breast cancer.⁷⁵ Second generation aromatase inhibitors were tested in Europe in the 1980s and were found to be as efficacious as tamoxifen.⁷⁵ Finally, the third generation aromatase inhibitors were approved in the U.S. to treat postmenopausal breast cancer in the 1990s and proven to be superior to tamoxifen.^{76,85} These new inhibitors have a benign side effect profile and suppress estrogen production in extraovarian tissues and within the breast cancer tissue itself. This effectively blocks estrogenic action, reduces recurrences and prolongs disease free survival in postmenopausal women with breast cancer.^{76,77} Aromatase inhibitors are also effective in the treatment of breast cancer that became resistant to treatment with tamoxifen.⁸⁰

In these studies, tumors that express estrogen receptor (ER) were more responsive to aromatase inhibitors compared with the tumors with an unknown receptor status.^{76,77} Future studies are required to determine whether aromatase inhibitors might be beneficial in ER-negative tumors via ER-independent mechanisms.

Aromatase and Endometrial Cancer

The role of aromatase expression and the therapeutic use of aromatase inhibitors have been well defined for breast cancer and endometriosis. Aromatase is also expressed in endometrial cancer tissue and aromatase inhibitors have been used to treat endometrial cancer.^{16-18,86} The pathologic significance of local estrogen biosynthesis via aromatase expression in endometrial cancer tissue or the therapeutic value of aromatase inhibitors in its management, however, is not clear at the moment.

Endometrial cancer is the most common gynecological malignancy found in women.⁸⁷ These tumors rather often have high concentrations of estrogen receptors, their growth is clearly enhanced by estrogen and unopposed estrogen exposure (in the absence of progesterone) predisposes women to development of endometrial cancer.⁸⁷ Although there is no consistent evidence of increased concentrations of circulating estrogen in women with endometrial cancer, the local concentration of estradiol in endometrial cancer tissues was reported to be higher than that in blood and in the endometrium of cancer-free women.⁸⁸⁻⁹² It is therefore conceivable that endometrial cancer itself synthesizes estradiol in situ, which then contributes to growth and possibly carcinogenesis.

A conversion study demonstrated significant conversion of androstenedione to estrone in endometrial cancer tissue.⁹³ Aromatase protein and mRNA were detected in endometrial cancer using immunohistochemistry and RTPCR, whereas aromatase expression was low or undetectable in endometrial hyperplasia (a precursor lesion of endometrial cancer).^{16,94} These observations are suggestive that intratumoral aromatase may play a role in the pathology of endometrial cancer. Immunoreactive aromatase was found in malignant epithelial, endometrial stromal and myometrial cells. Aromatase in stromal but not epithelial cells correlated positively with advanced surgical stage and poor survival (Dr. Makio Shozu, personal communication).

Currently available 3rd generation aromatase inhibitors may be used for endocrine treatment of endometrial cancer.⁹⁵ Treatment of endometrial cancer tissues in vitro with aromatase inhibitors demonstrated that in situ depletion of estrogen results in decreased cell proliferation of tumor cells.⁹⁶ Treatment of women with endometrial cancer with aromatase inhibitors blocked estrogen production in vivo in tumor tissue.⁹⁷ Safety data from the clinical trials of postmenopausal women with breast cancer indicated a preventive role of aromatase inhibitors in that an aromatase inhibitor reduced the risk of endometrial cancer.⁹⁸

On the other hand, the therapeutic efficacy of aromatase inhibitors in advanced endometrial cancer is not clear. The Gynecologic Oncology Group (GOG) performed a phase II trial of anastrozole in advanced, recurrent, or persistent endometrial cancer.¹⁸ Twenty-three patients were entered, all with grade 2 or 3 cancers. Two partial responses were noted.¹⁸ In a recent report, two cases of reproductive-aged women with grade 1 endometrial cancer who were treated with medroxyprogesterone acetate and anastrozole daily for 3 and 6 months subsequently reverted to normal endometrium. A progestin combined with the elimination of production of estrogen may be an effective therapy in well differentiated endometrial cancer in the obese premenopausal woman.¹⁷ Thus, there are some early preliminary encouraging results. It is, however, too early to predict whether aromatase inhibitors will be used widely in the treatment of endometrial cancer.

Aromatase and Ovarian Cancer

Estrogen, Progesterone and Ovarian Cancer

Since premenopausal ovary contains large amounts of estrogen and progesterone and ovarian cancer largely develops in postmenopausal women, it was hypothesized that ovarian cancer was not an ovarian steroid-responsive disease. In view of the protective effect of oral contraceptives and pregnancy, the disruptive and inflammatory effects of incessant ovulation on the surface epithelium were hypothesized to represent the primary mechanism of ovarian carcinogenesis.^{19,20,23}

Accumulating evidence, however, are suggestive of significant roles of estrogen and progesterone in ovarian cancer. A retrospective 1979-1998 cohort study of 44,241 postmenopausal women revealed that estrogen only hormone replacement therapy (HRT), particularly for 10 or more years, significantly increased risk of ovarian cancer. Relative risks for 10 to 19 years and 20 or more years were 1.8 (95% CI, 1.1-3.0) and 3.2 (95% CI, 1.7-5.7), respectively (p-value for trend <0.001), This study did not show an increased risk in women who used short-term estrogen plus progestin HRT, but authors suggested that risk associated with longer-term estrogen plus progestin HRT warrants further investigation.⁹⁹ The estrogen plus progestin WHI HRT study found a trend for increased ovarian cancer risk, which though was not significant.¹⁰⁰ These are some of the first pieces of important epidemiological evidence supporting that estrogen might plays a critical role in the pathophysiology of ovarian cancer. This has also been supported by other epidemiological and experimental data.^{19.20,23}

Estrogens appear to favor neoplastic transformation of the ovarian surface epithelium while progesterone offers protection against ovarian cancer development.^{21,22} Specifically, estrogens, particularly those present in ovulatory follicles, are both genotoxic and mitogenic to ovarian surface epithelial cells. In contrast, pregnancy-equivalent levels of progesterone are highly effective as apoptosis inducers for ovarian epithelial and ovarian cancer cells. In this regard, high-dose progestin may exert an exfoliation effect and rid an aged ovarian surface epithelium of premalignant cells. A limited number of clinical studies have demonstrated efficacies of antiestrogens, aromatase inhibitors and progestins alone or in combination with chemotherapeutic drugs in the treatment of ovarian cancer. As a result of increased life expectancy in most countries, the number of women taking HRT in general continues to grow. Since in the U.S. after WHI a serious drop in the number of HRT prescriptions was registered. Therefore, knowledge of the mechanism of action of steroid hormones on the ovarian surface epithelium and ovarian cancer is of paramount significance to HRT risk assessment and to the development of novel therapies for the prevention and treatment of ovarian cancer.^{19,20,23}

A Possible Link between Endometriosis, Ovarian Cancer, Aromatase and Prostaglandin Formation

Another piece of evidence regarding the role of estrogen in ovarian carcinogenesis comes from the biological and epidemiological links of endometrioid ovarian cancer to endometriosis.^{19,20,23} It was proposed that endometrioid and/or undifferentiated ovarian cancer may develop from the endometriotic implants seeded on the ovarian surface epithelium.^{19,20,23} Endometriosis is an estrogen-dependent disease that affects 6-10% of U.S. women of reproductive age (approximately 4-7 million).¹⁰¹ Endometriosis is a systemic disorder that is characterized by the presence of endometrium like tissue in ectopic sites outside the uterus, primarily on pelvic peritoneum and ovaries.¹⁰¹ Local estrogen formation via aromatase expression in endometriosis is extraordinarily important for its pathogenesis and treatment.^{86,102-106} In fact, work from our laboratory and other investigators over the past 10 years uncovered a molecular link between inflammation and estrogen production in endometriosis.⁸⁶ This is mediated by a positive feedback cycle that favors expression of aromatase and COX-2 and continuous local production of estradiol and PGE₂ in endometriotic tissue.¹⁰⁷⁻¹⁰⁹ In this vicious cycle, PGE₂ induces aromatase, whereas the product of aromatase enzyme, i.e., estradiol, induces COX-2 and thus PGE₂ formation. It is possible that PGE₂-induced aromatase expression may also play a biological role and represent a therapeutic target in a subset of ovarian carcinomas related to endometriosis.

Aromatase Inhibitors for the Treatment of Ovarian Cancer

Ovarian cancer is the fourth most common cause of cancer death in women. Most patients present at an advanced stage and the disease commonly relapses after primary surgery and chemotherapy. Relapsed ovarian cancer is not curable in the majority of women and responses to salvage chemotherapy are often sustained for less than a year at the expense of toxicities that may diminish quality of life.

In experimental models of ovarian cancer, it was demonstrated that moderate-high expression of ER α is associated with a growth response to estrogen and these models are growth-inhibited by antiestrogen strategies both in vitro and in vivo.¹¹⁰⁻¹¹² In addition, a number of proteins are estrogen regulated and these include the PR, the EGF receptor and HSP27.¹¹³⁻¹¹⁶

Clinical studies of tamoxifen in chemoresistant ovarian cancer have suggested that a subset of unselected patients respond to tamoxifen treatment, but the characteristics of responding tumors have not been defined. In one study, 105 patients in first relapse received tamoxifen and an overall response rate of 17% was reported.¹¹⁷ This was later reanalyzed to give a 13% response rate in cisplatin-resistant disease and 15% in cisplatin-sensitive disease.¹¹⁸ Another trial of tamoxifen reported a 17% response rate in chemoresistant disease.¹¹⁹ ER α has been suggested to be linked with clinical response, but to date no significant association has been demonstrated.¹¹⁷

The therapeutic role of aromatase inhibitors have recently been explored in advanced ovarian cancer. Phase II studies suggested that aromatase inhibitors are at least as effective as other hormonal treatments in advanced ovarian cancer.²⁴⁻²⁷ Because of the significant epidemiological association of estrogen with ovarian cancer, the minimal side effects of aromatase inhibitors and demonstrated activity of hormonal therapies in other endocrine associated malignancies, further study is needed.

Conclusion

Estrogen is positively linked to the pathogenesis and growth of three common women's cancers (breast, endometrium and ovary). A single gene encodes the key enzyme for estrogen biosynthesis named aromatase, inhibition of which effectively eliminates estrogen production in the entire body. Aromatase inhibitors successfully treat breast cancer, whereas their roles in endometrial and ovarian cancers are less clear and need further exploration.

Acknowledgement

This research work was supported, in part, by the NIH grants CA67167, HD38691 and HD46260 and grants from the AVON Foundation, Northwestern Memorial Foundation, Lynn Sage Cancer Research Foundation and the Friends of Prentice (to SEB).

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