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Hormone response in ovarian cancer: time to reconsider as a clinical target?

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Abstract

Ovarian cancer is the sixth most common cancer worldwide among women in developed countries and the most lethal of all gynecologic malignancies. There is a critical need for the introduction of targeted therapies to improve outcome. Epidemiological evidence suggests a critical role for steroid hormones in ovarian tumorigenesis. There is also increasing evidence from *in vitro* studies that estrogen, progestin, and androgen regulate proliferation and invasion of epithelial ovarian cancer cells. Limited clinical trials have shown modest response rates; however, they have consistently identified a small subset of patients that respond very well to endocrine therapy with few side effects. We propose that it is timely to perform additional well-designed trials that should include biomarkers of response.

Introduction

Ovarian cancer is the sixth most common cancer worldwide among women in developed countries and the most lethal of all gynecologic malignancies (Jemal *et al.* 2011). About 90% of primary malignant ovarian tumors are epithelial carcinomas and are further classified as serous, endometrioid, clear cell, mucinous, transitional, mixed cell, or undifferentiated based on cell morphology (Cho & Shih 2009). Recent technological

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advances have shed light on both the cellular and the molecular biology of ovarian cancer such that it is now widely believed that 'ovarian cancer' is a general term for a group of molecularly and etiologically distinct diseases that share an anatomical location (Vaughan *et al.* 2011). In particular, the diverse histological types of epithelial tumors are believed to be derived from different tissues. For example, high-grade serous carcinomas are believed to arise from the ovarian surface epithelium and/or the distal fallopian tube (Bell 2005, Crum *et al.* 2007), whereas endometrioid and clear cell carcinomas are believed to arise from endometriotic lesions (Nezhat *et al.* 2008). In contrast, most mucinous tumors are believed to be metastases to the ovary from the gastrointestinal tract, including the colon, appendix, and stomach (Lee & Young 2003, Kelemen & Kobel 2011, Zaino *et al.* 2011).

Despite the differences in the putative tissues of origin of epithelial ovarian cancers (EOC), the presence of sex steroid hormone receptors in many of these tissues of origin for ovarian cancer (Catalano *et al.* 2000, Akahira *et al.* 2002, Wada-Hiraike *et al.* 2006, Horne *et al.* 2009, Shao *et al.* 2011), as well as in many malignant epithelial ovarian tumors (Rao & Slotman 1991), suggests a potential role for hormones in the origin and promotion of these diseases. However, at this point in time, detailed mechanistic studies are lacking, and models to study hormone response *in vitro* and *in vivo* are very limited. Few and often small clinical trials have not resulted in any advances in the use of endocrine treatment for ovarian cancer. In this review, we will summarize and discuss epidemiological evidence and laboratory data that collectively strongly support a critical role for hormones in ovarian cancer development and progression, focusing on the steroids estrogen, progesterone, and androgen. We will also provide an overview of clinical trials targeting the receptors of these steroids. Due to space limitations, we will not discuss the role of gonadotropins in this review and ask the readers to refer to previous reviews on this topic (Zheng *et al.* 2007, So *et al.* 2008, King & Wong 2011). We will end the review with our suggestions for future directions for the field, such as the need for further mechanistic studies and well-designed clinical trials, which should include the use of biomarkers.

The epidemiology of EOC

Reproductive and hormonally related risk factors

The most consistently reported reproductive and hormonally related factors found to protect against EOC are use of oral contraceptives (OCs), increasing parity, and having a tubal ligation. In contrast, increasing age and nulliparity have been consistently shown to increase EOC risk. Other hormonally linked factors, such as hormone replacement therapy (HRT) use, infertility, endometriosis, breast-feeding, hysterectomy, and central adiposity, have shown some association with EOC, but the data are neither as strong nor as consistent as OC use, parity, and tubal ligation. Table 1 summarizes the relationship between reproductive and hormonally related risk factors and EOC.

OCs, HRT, and exogenous hormones—Both prospective and case-control studies report about a 30% decrease in ovarian cancer risk with ever use of OCs (Collaborative Group on Epidemiological Studies of Ovarian Cancer 2008), and even short-term use (6 months or less) appears protective (Greer *et al.* 2005). Longer duration of use imparts increased protection, with a 20% decrease in risk for each 5 years of use. More recent OC use is associated with greater protection, but even after stopping its use for 30 or more years, risk is still reduced by about 15% (Collaborative Group on Epidemiological Studies of Ovarian Cancer 2008). Risk reduction appears consistent for all histological subtypes of EOC, except for possibly mucinous tumors. Estrogen dose does not appear to affect the OC-EOC association (Collaborative Group on Epidemiological Studies of Ovarian Cancer 2008) whereas some higher progestin dose formulations may confer greater protection, although

that association has not been consistent (Ness *et al.* 2000, Schildkraut *et al.* 2002, Pike *et al.* 2004, Lurie *et al.* 2007).

While the data on OC use and EOC have been consistent, the data for HRT have not. However, more recent studies, including the prospective Women's Health Initiative (WHI) (Anderson *et al.* 2003) and the Million Women Study (Beral *et al.* 2007), report an increase in risk for both estrogen-only (ET) and estrogen–progestin (EPT) formulations, although the risk associated with EPT was lower than that of ET. A recent meta-analysis of 14 published studies found risk increases 22% per 5 years of ET use compared with only 10% per 5 years of EPT use, suggesting that risk differs by regimen (Pearce *et al.* 2009).

Exogenous androgens may be associated with EOC. One case–control study found that use of Danazol, a synthetic androgen commonly used in the treatment of endometriosis, significantly increased EOC risk (Cottreau *et al.* 2003), although this finding has not been replicated (Olsen *et al.* 2008). Ever use of testosterone (tablets, patches, troches, or cream) has been associated with a threefold increase in EOC (Olsen *et al.* 2008).

Childbearing and breast-feeding—Like OC use, childbearing has been consistently shown to be associated with a reduced risk of EOC in both prospective and case–control studies, with an observed protective effect similar to or greater than that observed for OC use (about 30% for the first full-term pregnancy). Moreover, increasing parity is associated with increasing protection, with each additional full-term pregnancy conferring about a 10% decrease in risk (Braem *et al.* 2010, Tsilidis *et al.* 2011). Even an incomplete pregnancy appears to provide some protection against EOC, although the magnitude of the protective effect is less than that of a full-term pregnancy (Riman *et al.* 1998). Finally, multiple births in a single pregnancy (giving birth to twins, triplets, etc.) may be associated with a greater risk reduction than singleton births (Whiteman *et al.* 2000). In contrast to childbearing, the evidence for a relationship between breast-feeding and EOC is inconsistent, although most studies show a small negative association (Riman *et al.* 2004, Danforth *et al.* 2007).

Reproductive disorders and other reproductive factors—Factors affecting childbearing have also been shown to be associated with EOC. In most studies, infertility has been associated with an increased risk, which may be greatest among women who fail to conceive (Vlahos *et al.* 2010). In general, infertility treatment does not appear to increase EOC risk, although the subset of treated women who remain nulliparous may be at an increased risk (Vlahos *et al.* 2010).

Endometriosis, defined as the presence and growth of endometrial tissue outside the uterine cavity, has also been associated with EOC. A recent pooled analysis of 13 case–control studies showed a threefold increase in the incidence of clear cell EOC and a twofold increase in endometrioid EOC among women with a self-reported history of endometriosis (Pearce *et al.* 2012).

An increased risk of EOC was reported by one case–control study (Schildkraut *et al.* 1996) among women with polycystic ovary syndrome (PCOS), a condition associated with menstrual dysfunction, infertility, obesity, the metabolic syndrome, hyperandrogenism, and insulin resistance. However, the finding was based on a small number of cases ($n=7$) and the association was limited to nonusers of OCs and thin women. Further case–control and prospective studies have failed to confirm this relationship (Pierpoint *et al.* 1998, Olsen *et al.* 2008, Brinton *et al.* 2010).

Tubal ligation has been consistently shown to be associated with reduction in EOC risk (Cibula *et al.* 2011). This protection appears similar in magnitude to OC use and child

bearing (about 30%) and is protective in high-risk women (i.e. BRCA1/2 carriers) as well. Hysterectomy has also been shown to reduce EOC risk, although the magnitude of the association is not as great nor as consistent as that reported for tubal ligation (Riman *et al.* 2004). Finally, reproductive factors associated with other hormonally linked cancers, such as age at first menarche, age at menopause, and length of reproductive years, have not been consistently associated with EOC (Riman *et al.* 2004).

Weighing the hormone: EOC epidemiological evidence

The epidemiological relationships between reproductive and hormonally related risk factors and EOC suggest that hormones play a role in the etiology of the disease. Indeed, an early, small (31 cases and 62 controls), prospective study reported that circulating levels of androstenedione and DHEA were significantly associated with increased risk of EOC in a dose-dependent fashion, whereas levels of DHEAS, estradiol, estrone, and progesterone were not (Helzlsouer *et al.* 1995). However, three subsequent prospective studies, with a total of over 580 ovarian cancer cases, have failed to replicate these early positive hormone–EOC associations (Lukanova *et al.* 2003, Rinaldi *et al.* 2007, Tworoger *et al.* 2008). Thus, the evidence to date suggests that circulating hormone levels are not associated with EOC. Based on the existing data, however, the exact nature of the hormone–EOC link remains unclear, with some factors supporting and others refuting a direct association. Below we will describe and discuss current understanding of roles of hormones in EOC.

Estrogens—The evidence linking estrogens to EOC are mixed. Although pregnancy raises circulating estrogen levels, intraovarian levels are reduced. OCs reduce endogenous estrogen levels (Killick *et al.* 1987). In contrast, risk-conferring HRT raises circulating estrogen levels. Breast-feeding reduces both circulating and intraovarian estrogen levels (Rosenblatt & Thomas 1993). Endometriosis is associated with an increased local production of estradiol as well as increased expression of estrogen receptor α (ER α). Moreover, compared with levels in the early reproductive years, circulating peri-menopausal estradiol levels are higher, reflecting the period when EOC risk begins to sharply rise (Prior 2005). Together, these data suggest that estrogens may be associated with an increase in EOC risk.

Further evidence for an estrogen–EOC link comes from genetic susceptibility studies. Specifically, a large, international, pooled analysis of ten case–control studies comprising 4946 women with primary invasive EOC and 6582 controls found that women with the rs1271572 TT genotype were at a significantly increased risk of EOC compared with women with the G allele (Lurie *et al.* 2011). The association was even stronger among women \geq 50 years. rs1271572 is located in the promoter of the ER β gene (*ESR2*), where it maps to a binding region for MyoD and AP-4, previously shown to be important for expression of ER β (Li *et al.* 2000). ER β is believed to inhibit proliferation and motility of ovarian cancer cells as well as facilitate apoptosis (Bardin *et al.* 2004a, Cheng *et al.* 2004b, Treeck *et al.* 2007). Hence, there is a potential causal association of rs1271572 with EOC, especially among pre- and perimenopausal women, who have higher circulating estrogen concentrations compared with their postmenopausal counterparts, further supporting an association between estrogens and EOC.

However, other factors do not support an estrogen–EOC link. No prospective study has found an association between circulating estrogen levels and EOC. OC use reduces endogenous estrogen levels; however, OC estrogen dose does not alter the magnitude of the observed EOC protective effect (Collaborative Group on Epidemiological Studies of Ovarian Cancer 2008). Finally, reproductive factors, such as early menarche and late menopause, which are associated with greater estrogen exposure and are linked to estrogen-associated breast and endometrial cancers, have not been consistently associated with EOC.

Progesterone and progestins—Epidemiological data suggest that progestins and progesterone may have a protective role against EOC. Progestin-containing OCs raise circulating progesterone levels about threefold. Progestin-only OCs may be as protective as EPT regimens and high-dose progestin OCs may be more protective than low-dose OCs (Rosenberg *et al.* 1994). Pregnancy, a consistently strong protective factor, raises progesterone levels, too. Third trimester circulating progesterone levels are more than ten times higher than luteal phase levels during the menstrual cycle. Moreover, progesterone levels in non-singleton births, which are more protective against EOC than singleton births, are higher than in singleton births (Batra *et al.* 1978, Haning *et al.* 1985). Although the risk of EOC appears increased with HRT use, regimens containing a progestin confer a lower relative risk compared with ET regimens, suggesting that the progestin component may mitigate the deleterious effect of estrogens. Ovulatory infertility, which is associated with reduced progesterone production, may increase EOC risk (Brinton *et al.* 1989, Rossing *et al.* 1994). Finally, endometriosis is associated with resistance to progesterone, which has been suggested to be due to the presence of inhibitory PR-A isoform and absence of transcription activating PR-B isoform (Attia *et al.* 2000).

Although the epidemiological data linking progesterone with reduced EOC risk are substantial, they are not definitive. For example, the data on OC progestin dose and EOC are not consistent. Additional compelling data come from a large consortium of 12 ovarian cancer case–control studies: a functional SNP, +331/CT, in the progesterone receptor (PR) alters the relative transcription of the two PR isoforms, PR-A and PR-B. Variants associated with increased PR-B, which acts as a classical steroid receptor, in theory should be associated with decreased EOC risk. Conversely, variants associated with increased PR-A, which inhibits in part PR-B, should be associated with increased risk. However, in this large study involving 4788 cases and 7614 controls, PR variants were not found to alter EOC risk, except for possibly the endometrioid subtype (Pearce *et al.* 2008).

Androgens—Several pieces of epidemiological data support a role for androgens in EOC development. OCs, one of the strongest protective factors, reduce circulating androgen levels (Gaspard *et al.* 1983, Murphy *et al.* 1990, Coenen *et al.* 1996). Tubal ligation and hysterectomy are also associated with decreased circulating androgen levels (Laughlin *et al.* 2000, Davison *et al.* 2005, Danforth *et al.* 2010). The potential association between PCOS, a hyperandrogenic condition, and EOC provides further support for the androgen–EOC link. Finally, the possible increased risk associated with use of exogenous androgenic agents further supports the relationship.

Despite these epidemiological associations, the data are not conclusive. Circulating androgen levels have not been associated with increased risk in three large prospective studies. Notably, androgen levels decline with age, with the decline being greater in the earlier reproductive years than in later decades (Davison *et al.* 2005). This is in contrast to the age–EOC relationship wherein incidence and risk rise slowly in the early reproductive years, then sharply increase beginning at around age 40, and continue to increase through late age (Howlader *et al.* 2011). In addition, factors associated with androgen levels as well as those altering androgen levels have not been consistently associated with EOC. For example, PCOS has not been consistently associated with the disease and the data linking exogenous androgen use with EOC have not been confirmed. Furthermore, OC formulations that contain androgenic agents do not differ in the magnitude of their protective effect compared with non-androgenic formulations (Greer *et al.* 2005). Finally, a trinucleotide repeat polymorphism in exon one of the androgen receptor (AR) gene, the length of which is inversely associated with the ability of the AR–ligand complex to transactivate AR-responsive genes, has been inconsistently associated with EOC in both magnitude and

direction (Spurdle *et al.* 2000, Menin *et al.* 2001, Santarosa *et al.* 2002, Lee *et al.* 2005a, Schildkraut *et al.* 2007, Ludwig *et al.* 2009).

Summary of epidemiological data and the hormone–EOC associations—The epidemiological evidence suggests a role for androgens and estrogens in the etiology of EOC. Data further support a protective role for progesterone and progestins against EOC. However, while supportive, the data are inconclusive and in some cases contradictory. One possible explanation may be that the relationship between hormones and EOC risk depends more on the intraovarian environment than on circulating hormone levels (Lukanova & Kaaks 2005). It is also possible that the timing of exposures, for example, in the pre- vs the postmenopause years, may be critical. Moreover, the interaction of lifestyle and hormonal factors may alter the hormone–EOC link. However, due to study design issues such as sample size and recall of specific details when assessing exposures that precede diagnosis by many years, such interactions are challenging to assess. Finally, the heterogeneity of EOC may contribute to the inconsistent findings linking hormonally associated reproductive, lifestyle, and host factors with the diseases. To date, most epidemiological studies have considered EOCs as a single disease, which may partially explain inconsistencies among studies as well as some negative finding. Indeed, a very recent study suggests that association between circulating levels of sex steroid hormones and EOC differ by tumor histology and invasiveness. These data suggest that the previously reported negative findings (Lukanova *et al.* 2003, Rinaldi *et al.* 2007, Tworoger *et al.* 2008, Modugno & Edwards 2012) may be due to treating EOCs as a single entity. In the future, it will be important to determine potential associations with histological and molecular subtypes of EOC.

***In vitro* and *in vivo* studies of hormone action in ovarian cancer**

Role of ER signaling action in ovarian cancer cells

Estrogen exerts its effect through two receptors, ER α and ER β . A number of studies have addressed the expression of both isoforms in clinical samples and their functions in cell line models. ER β is highly expressed throughout the normal ovary, including in granulosa cells, theca cells, corpora lutea, oocytes, as well as cultures of primary ovarian surface epithelial (OSE) cells (Kuiper *et al.* 1996, Byers *et al.* 1997, Brandenberger *et al.* 1998, Hillier *et al.* 1998); however, its expression is progressively lost during ovarian cancer development and progression (Lau *et al.* 1999, Bardin *et al.* 2004b, Lazennec 2006, Chan *et al.* 2008). While this loss has been associated with loss at the genetic level, there is increasing evidence that lower expression of ER β can also result from epigenetic changes, namely hypermethylation of its promoter (Geisler *et al.* 2008, Suzuki *et al.* 2008, Yap *et al.* 2009). Of interest, a recent study showed nuclear localization of ER β in normal ovarian tissue, but cytoplasmic localization in the tumor tissue, which was associated with worse outcome (De Stefano *et al.* 2011). In contrast, ER α expression is maintained, or even increased, in a subset of ovarian tumors (Rao & Slotman 1991, Chan *et al.* 2008). As a result, there is an increase in the ER α /ER β ratio with malignant progression of the ovary. There is limited knowledge about the expression of ER α /ER β in different histological subtypes of ovarian cancer, and even less about their expression in different molecular subtypes, such as the proliferative, immunoreactive, differentiated, and mesenchymal subtypes in high-grade serous cancers, as defined by the TCGA analysis (Bell *et al.* 2011).

Although details are little understood, a number of studies clearly show that estrogen treatment exerts pro-proliferative action, which can be blocked with the antiestrogens tamoxifen and ICI 182 780 (Galtier-Dereure *et al.* 1992, Langdon *et al.* 1994). There is also increasing evidence for estrogen mediating increased motility and invasion of ovarian cancer cells (Hua *et al.* 2008, Zhu *et al.* 2012). In a recent *in vivo* study, using a mouse model in

which overexpression of SV40Tag in OSE results in poorly differentiated ovarian tumors, estrogen treatment caused an earlier onset of tumors, decreased overall survival (OS) time, and a distinctive papillary histology (Laviolette *et al.* 2010).

To confirm that estrogen-mediated growth stimulatory effects in ovarian cancer cell lines were indeed mediated by ER α , O'Donnell *et al.* (2005) treated ER α /ER β -expressing ovarian cancer cells with ER α and ER β -specific ligands. Treatment with PPT (the ER α ligand) but not with DPN (the ER β ligand) resulted in growth stimulation, confirming a role for ER α in the estrogen-mediated growth stimulation. Given the tumor suppressive-like activity of ER β , it is not surprising that overexpression of ER β can result in inhibition of ovarian cancer cell motility and invasion (Zhu *et al.* 2011). One might expect some growth inhibition with ER β ligands, but such studies have yet to be performed in ovarian cancer cells.

A number of studies have identified ER α target genes in ovarian cancer cells, revealing some overlap with the estrogen response in breast cancer cells but also unique targets. Early studies have shown regulation of genes involved in proliferation, invasion and cell cycle regulation such as cathepsin (Rocheffort *et al.* 2001), c-fos, pS2, cyclins (Albanito *et al.* 2007), TGF α (Simpson *et al.* 1998), fibulin (Clinton *et al.* 1996, Roger *et al.* 1998, Moll *et al.* 2002), c-myc (Chien *et al.* 1994), and SDF-1 (Hall & Korach 2003), PR (Langdon *et al.* 1998), and more recently members of the semaphorin family (Joseph *et al.* 2010). In addition, a number of IGFbps have been described to be regulated by estrogen (O'Donnell *et al.* 2005), and levels of IGFBP3, IGFBP4, and IGFBP5 were predictive in a letrozole trial in ovarian cancer patients (Walker *et al.* 2007a).

A gene expression array study has been reported, in which ER-positive PEO1 cells were treated with estradiol for 24 h, and target genes were identified using a 1.2K array. Using a threefold change as cutoff, the authors identified five induced and 23 downregulated genes. The induced genes were *TNFD7*, *TRAP1*, *FOSL1*, *TFAP4*, and cathepsin D, and among the repressed genes were *cyr61*, vimentin, fibronectin, *IGFBP3*, and several keratins (Hall & Korach 2003). The finding of repression was more dominant compared with induction, is of interest; however, given the 24-hour treatment, additional experiments are necessary to show that the identified genes are truly direct ER targets. Of note, there is one study detailing mechanism of estrogen-mediated repression of target genes, focusing on the folate receptor, in ovarian cancer cells (Kelley *et al.* 2003). The authors show a role for ER α and the corepressor SMRT, and a lack of involvement of coactivators, including those of the p160 family in the repression.

In contrast to breast cancer for which ligand-independent activation of ER activity has been well described, there is limited literature on such activity in ovarian cancer. One study has described ERK2-mediated phosphorylation of ER α in response to DC44 interaction with hyaluron and subsequent IQGAP1 recruitment (Bourguignon *et al.* 2005). There is some preliminary evidence that the observed interaction between hormone response and obesity in ovarian cancer could result from cross talk between ER α and leptin signaling. Using BG-1 cells as a model system, Choi *et al.* (2011) showed that treatment with ICI 182 780, a pure ER antagonist, blocked leptin-induced cell proliferation. The effect was mediated by ER α interaction between phosphorylated Stat3, which was at least in part mediated by ERK and PI3K pathways. Given the recent finding of BG-1 cells being identical to MCF-7 breast cancer cells, these data should be interpreted with caution, unless DNA fingerprinting was performed to confirm authenticity of the BG-1 cells as ovarian cancer cells (Korch *et al.* 2012).

Lack of response to estrogen and antiestrogen can result from primary resistance (e.g. due to lack of ER expression or activity) or can develop as secondary (i.e. acquired) resistance, for

example, resulting from activation of alternative pathways. However, there has been a paucity of studies on endocrine resistance in ovarian cancer, with few exceptions. The SKOV3 cells, for example, have been described to be resistant to estrogen and antiestrogen treatment, associated with loss of PR, and overexpression of Her2 and cathepsin D (Hua *et al.* 1995). Subsequently, Lau *et al.* (1999) described a 32 bp deletion in exon 1 of ER α in SKOV3 cells, which is potentially very exciting; however, no follow-up studies have been reported. Further studies of ER target genes *in vitro* but also *in vivo*, such as in established estrogen-sensitive xenografts from PE04, OVA-5, and OVCAR-3 cells (Ritchie & Langdon 2001) and in resistance models, and mechanistic analysis of ER and coregulator recruitment, for example, using chromatin immunoprecipitation studies, are warranted to move this field forward.

Progesterone action in normal ovary and ovarian cancer cells

There is evidence that loss of PR expression is associated with increasing grade of ovarian cancer (Langdon *et al.* 1998, Lau *et al.* 1999, Akahira *et al.* 2002). Also, in contrast to ER, where little is known about subtype-specific expression, PR levels seem to be higher in endometrioid tumors compared with other ovarian cancers (Langdon *et al.* 1998). The same study reported an association between PR, stage, and outcome – PR expression was higher in low-grade tumors, and high PR expression was associated with improved survival.

A number of studies showed that progesterone treatment of OSE and ovarian cancer cells results in decreased proliferation and increased apoptosis. Rodriguez *et al.* (1998) performed *in vivo* studies, using primates, which showed that progesterone treatment resulted in a four- to six-fold increase in the number of apoptotic ovarian epithelial cells compared with control and estrogen treated monkey. The same group subsequently showed that this effect can be enhanced by treatment with NSAIDs, a concept of potential interest for the prevention of ovarian cancer (Rodriguez *et al.* 2012). The induction of apoptosis was associated with decreased expression of TGF β 1 and increased expression of TGF β 2 and TGF β 3, suggesting that differential regulation of TGF β family members could play a role in progestin-mediated induction of apoptosis (Rodriguez *et al.* 2002). While details of underlying mechanisms are sparse, some studies have addressed this question and have shown enhanced TRAIL-mediated cell death (Syed *et al.* 2007), induction of p53 (Bu *et al.* 1997), and increased expression FasL (Syed & Ho 2003). Interestingly, in addition to the induction of apoptosis, activation of PR was also shown to induce cell cycle arrest and senescence (Takahashi *et al.* 2009).

As for ER, there is limited knowledge of PR downstream target genes in ovarian cancer. Syed *et al.* (2005) treated human OSE cells (HOSE 642, HOSE 6–8, and HOSE 12–12) and ovarian cancer cell lines (OVCA429, OVCA420, and OVCA432), previously described to be PR positive (Lau *et al.* 1999), with high-dose progesterone. Using a small gene expression array ($n=2000$ genes), the authors identified 171 progesterone-regulated genes in HOSE cells and 135 in ovarian cancer cells. They focused on *ATF3*, caveolin-1, *DLC1*, and nm23-H2, showing that these genes were indeed direct PR target genes, and given their antitumor and anti-invasive properties, the authors speculate that induction of these candidates could be related to progesterone-induced antitumor effects.

Importantly, there is some evidence that progesterone might synergize with chemotherapeutic drugs to induce apoptosis. For example, high doses of progesterone were shown to enhance cisplatin-induced apoptosis (Murdoch *et al.* 2008). However, this is not straightforward, as other studies have reported opposite effects. Peluso *et al.* (2008, 2009) for example, showed that progesterone treatment can result in decreased cisplatin-induced apoptosis. This apparent controversy might be, at least in part, due to progesterone's action on two receptors, PRA and PRB, and their relative expression in the target cells. In addition,

progesterone can also bind to a protein complex containing the progesterone membrane receptor component 1 (PGRMC1/mPR). Indeed, the Pelusso laboratory has published a few studies showing an involvement of PGRMC1 in sensitivity of ovarian cancer cells to cisplatin (Peluso *et al.* 2008, 2009). Interestingly, PGRMC1 shares homology with cytochrome b5-related proteins rather than with hormone receptors and can bind heme (Peluso 2011). Different cellular responses to progesterone could also result from altered expression of the more recently identified membrane PRs mPR α , β , and γ . The mPRs belong to the larger progestin and adipoQ receptor gene family (mPR α – PAQR7; mPR β – PARQ8; and mPR γ – PARQ5), and while they are not members of the classical G-protein family, they can activate G-proteins and affect cAMP levels. They bind progesterone with high affinity, but do not interact with synthetic progestin R5020, or the PR antagonist RU486. They are abundantly expressed in ovarian cancer cells including those that lack expression of classical PR, and in clinical specimens representing all major histological subtypes (Romero-Sánchez *et al.* 2008, Charles *et al.* 2010). Charles *et al.* (2010) showed that progesterone alone did not affect cAMP levels in ES-2 and SKOV3 cells; however, it enhanced isoproterenol-induced and β 1,2-adrenergic receptor-mediated increases in cAMP levels. This resulted in activation of JNK1/2 and p38 MAPK activity and subsequently induction of pro-apoptotic genes like *Bax*.

Thus, the effects of progesterone in ovarian cancer cells are mediated by the classical PR, by PGRMCs, and finally by mPRs, through both genomic and non-genomic actions (for review, see Peluso (2007)). This signaling is undoubtedly very complex, and while potentially attractive as a clinical target, significantly more research needs to be done before we can begin to understand intricate details of progesterone action in ovarian cancer.

Limited understanding of AR action in ovarian cancer cell lines

The AR is expressed in both normal ovary and ovarian cancer. Edmondson *et al.* (2002) have shown that OSE cells express AR and respond to androgen with increased proliferation and attenuated apoptosis. Epithelial cells, especially those within inclusion cysts arising after ovulation, appear to be exposed to high levels of androgen (Risch 1998). Ovarian cancer cells also express 17 β -hydroxysteroid dehydrogenase (17 β -HSD) converting androstenedione (which is a weak androgen) to testosterone (Blomquist *et al.* 2002, Chura *et al.* 2009). Importantly, there are also a number of studies showing overexpression of AR in ovarian cancer (Kühnel *et al.* 1987, Chadha *et al.* 1993, Ilekis *et al.* 1997, Lau *et al.* 1999, Lee *et al.* 2005b).

Although sparse, there is evidence from *in vitro* studies suggesting that androgens affect gene expression, growth, invasion, and survival in ovarian cancer cell lines. Shi *et al.* (2011) have shown that androgen's effect on survival of ovarian cancer cells was associated with increased expression, activity, and phosphorylation of telomerase. They also showed an androgen-mediated degradation of the cell cycle inhibitor p27 (Shi *et al.* 2011). Androgen was reported to stimulate DNA synthesis/S-phase fraction in low-passage OSE culture (Syed *et al.* 2001, Edmondson *et al.* 2002), and in ovarian cancer cells (Sheach *et al.* 2009); however, other studies failed to observe an effect of androgens on growth (Karlan *et al.* 1995). Finally, activation of AR has been shown to stimulate ovarian cancer cell invasion (Gogoi *et al.* 2008, Ligr *et al.* 2011).

There is evidence for cross talk between androgen signaling and other signaling pathways. For example, Evangelou *et al.* (2000) showed that dihydrotestosterone (DHT) treatment of Hey and SKOV-3 ovarian cancer cells, and of ascites-derived OVCAS-16 cells, prevented growth inhibitory effect of TGF- β , while DHT alone had no effect on growth. This effect of DHT was associated with downregulation of TGF- β 1 and TGF- β 2 receptors. Interestingly, the same group went on to show similar effects of DHT on blocking TGF- β -mediated

growth inhibition in cells isolated from ovarian surface epithelium of women undergoing oophorectomy for non-ovarian indications or with a germline BRCA mutation (Evangelou *et al.* 2003).

To the best of our knowledge, there is only one study reporting a genome-wide analysis of androgen target genes. Sheach *et al.* (2009) identified more than 100 AR target genes in OVCAR3 cells with the majority being related to transcription, proliferation, and G-protein signaling. The G-proteins that were significantly induced by androgen treatment in OVCA3 cells included GNAI3, ELKS, GSTPI, RERG, Rab25, Rab45, and Rab35. The induction of Rab25 is of special interest as this small GTPase has previously been shown to proliferation, survival, and invasion of ovarian cancer cells (Fan *et al.* 2006) and to be overexpressed in aggressive ovarian cancers (Cheng *et al.* 2004a).

Coregulator proteins in ovarian cancer

Although not a major focus of this review, there is little doubt that coregulator proteins – activating or repressing steroid receptors – do play a role in ovarian tumorigenesis. SRC3/AIB1 was shown to be amplified in up to 25% of ovarian cancer (Bautista *et al.* 1997, Tanner *et al.* 2000). The coregulator p44/Mep50/WDR77 increases activities of ER and AR and stimulates proliferation and invasion of ovarian cancer cells in the presence of estrogen or androgen (Ligr *et al.* 2011). It shows strong cytoplasmic localization in normal ovarian surface, and fallopian tube epithelia, while it is mainly in the nucleus in invasive ovarian carcinoma. The coactivator ARA70 was highly expressed in invasive ovarian tumors but not in the normal ovary (Shaw *et al.* 2001). And finally, corepressor proteins have also been described to be highly expressed in ovarian cancer – Havrilesky showed that 35% of tumors expressed NCoR, and 71% expressed NCoR2/SMRT (Havrilesky *et al.* 2001). Given the known critical role of coregulator proteins in sensitivity and resistance to hormones, future studies on their expression and function in ovarian cancer development and progression are warranted.

Summary: basic science of hormone response in ovarian cancer

Steroid hormone signaling in ovarian cancer is complex yet little understood. It is imperative that we develop and use *in vitro* as well as *in vivo* models representing the different histological and molecular ovarian cancer subtypes to decipher signaling pathways and downstream target genes of the steroid receptors before we can improve efficacy of endocrine treatment in ovarian cancer.

Clinical trials involving hormonal therapy

Introduction

Targeting hormone receptors has been a successful therapeutic strategy in endocrine-sensitive tumors such as breast, prostate, and endometrial cancers (Moreau *et al.* 2006, Decruze & Green 2007, Poole & Paridaens 2007). There is clear evidence that EOC is a hormone-responsive cancer, at least in a subset of tumors. Nuclear receptors are widely expressed in EOC: ER is expressed in 61–79% of EOC, with highest expression in serous and endometrioid subtypes (Glavind & Grove 1990, Rao & Slotman 1991, Lindgren *et al.* 2001, Lee *et al.* 2005b). PR is expressed in ~25–50% of all EOC, but is as high as 91% in endometrioid subtypes (Fujimura *et al.* 2001, Lindgren *et al.* 2001, Lee *et al.* 2005b), and expression of AR is as high as 90% (Kühnel *et al.* 1987). Despite the high expression of endocrine-responsive receptors, hormonal therapy has only a minor role in the treatment of EOC. Therapy targeting the ER, for example, has been employed since the 1960s, but variable clinical responses have limited their usefulness in ovarian cancer (Langdon & Smyth 2008). Given that hormonal therapy is a relative nontoxic anticancer therapy, which

is easy to administer and that it is well tolerated, the question is why hormonal therapy is not used more widely in ovarian cancer? Can it be made more effective? Below we summarize and discuss endocrine therapies in ovarian cancer, again focusing on ER, PR, and AR (also depicted and summarized in Fig. 1).

Targeting the ER in ovarian cancer

Tamoxifen, a selective ER modulator (SERM), that produces antiestrogen effects through competitive inhibition of ER has been used with variable results for the treatment of ovarian cancer. The majority of clinical trials involving tamoxifen were small phase II trials of patients with heavily pretreated recurrent disease. A prospective Gynecologic Oncology Group (GOG) study by Hatch *et al.* (1991), evaluated the response to tamoxifen in patients with recurrent or persistent disease following primary treatment. One hundred and five patients were treated with tamoxifen 20 mg twice a day after frontline chemotherapy. Study participants had an 18% objective response rate (RR), with 10% having a complete response (CR). Reanalysis of this study, focusing on patients with platinum-refractory disease, showed an objective RR of 13% (Markman *et al.* 1996). Of note, the objective RRs to second-line platinum-based therapy range from <10 to >40% (Markman & Bookman 2000). A similar objective RR of 17% (5/29) was seen in patients with unknown ER/PR status treated with tamoxifen after failure of cytotoxic therapy in three phase II clinical trials published by the Mid-Atlantic Oncology Program in 1993 (Ahlgren *et al.* 1993). Therefore, hormonal therapy is well within the response range of chemotherapeutic agents but considerably easier to administer and with fewer side effects. Unfortunately, few of these trials report ER status of the ovarian tumors. One study reported that patients with ER+ tumors had higher RRs to tamoxifen treatment than ER- tumors, although this difference was not statistically significant (Hatch *et al.* 1991). In another small study, all patients with stable disease (SD) were ER positive (Schwartz *et al.* 1982). ER status did not correlate with response in two other trials; however, ER status was known in only one-fourth to one-third of patients enrolled (Shirey *et al.* 1985, Weiner *et al.* 1987).

In 2010, the GOG published the results of a phase II trial of tamoxifen vs thalidomide in women with biochemical recurrence of EOC, fallopian tube, or primary peritoneal carcinoma based on rising CA125 after CR to frontline platinum/taxane therapy (Hurteau *et al.* 2010). The interim analysis did not show any difference in the benefit of thalidomide relative to tamoxifen, and the study was stopped early. At a median follow-up of 31 months, the progression-free survival (PFS) was 3.2 and 4.5 months while median survival was 24.0 and 33.2 months for thalidomide and tamoxifen arms respectively. Thalidomide was associated with an increased risk (HR=1.76, 95% CI=1.16–2.68) compared with tamoxifen. Also, there was significantly less toxicity associated with tamoxifen treatment, and more patients in this treatment arm received ≥ 3 cycles. As this trial did not contain a third arm (no treatment or another active agent), the possible interpretations of the tamoxifen results are limited. However, they are very encouraging and indicate that additional study of tamoxifen (and other endocrine treatments) in this patient population, i.e. asymptomatic patients with biochemical evidence of recurrent ovarian cancer, is warranted. Markman *et al.* (2004a) examined the evidence behind this management strategy with a retrospective review of 56 women with asymptomatic recurrent ovarian or primary peritoneal cancer treated with 20–40 mg Tamoxifen daily before initiation of cytotoxic chemotherapy. The median duration of treatment was 3 months, but 42% of patients were on Tamoxifen for ≥ 6 months and 19% for ≥ 12 months. The most common reasons for stopping single-agent tamoxifen were a continued rise of serum CA125, progression of disease on CT scan or physical exam, or the development of cancer-related symptoms. No standard of care exists for the management of asymptomatic recurrent disease, and therefore, trials testing hormonal therapy in this setting seem warranted.

While investigations continue to examine the utility of hormonal therapy, contemporary studies publishing randomized and nonrandomized trials evaluating the use of tamoxifen in EOC are lacking. A Cochrane review published in 2010 attempted to identify randomized and nonrandomized studies of more than ten patients with recurrent ovarian cancer treated with tamoxifen (Williams *et al.* 2010). Surprisingly, the search of articles published between 2002 and 2009 yielded no trials that satisfied the inclusion criteria. Publications during that period consisted only of observational data from single-arm studies of women treated with tamoxifen. Additionally, no data on tamoxifen's effect on symptom control, quality of life, or prolongation of life were available from the uncontrolled, non-comparative trials that were screened as part of this Cochrane review. A Cochrane review using the same inclusion criteria as the more recent publication did identify 11 nonrandomized series, one nonrandomized phase II study, and two randomized trials from 1997 to 2002 (Williams 2001; see Table 2). In total, 60 of the 623 (9.6%) women treated with tamoxifen had an objective response and SD of 4 weeks or more was seen in 131/411 (31.9%) women from eight studies. Due to lack of data, duration of response, survival, symptoms palliation, and quality of life were not assessed.

The majority of women who develop ovarian cancer are postmenopausal at the time of diagnosis. In postmenopausal women, the major source of circulating estrogen is from the peripheral conversion (in skin and adipose tissue) of androstenedione by the enzyme aromatase. Additionally, aromatase expression has been shown in malignant ovarian epithelial cells resulting in intratumoral production of estrogen (Cunat *et al.* 2005). The use of aromatase inhibitors (AI), such as letrozole and anastrozole, for the treatment of ER+ breast cancer has been a great success, and several studies have investigated the use of these agents in the treatment of recurrent or persistent ovarian cancer (see Table 3). Similar to other hormonal therapies, the results of AI treatment are variable. In seven phase II trials, with known ER status, the complete and partial RRs were 0–4 and 0–11% respectively (Bowman *et al.* 2002, del Carmen *et al.* 2003, Papadimitriou *et al.* 2004, Gourley *et al.* 2006, Verma *et al.* 2006, Smyth *et al.* 2007, Ramirez *et al.* 2008). However, benefit from AI therapy can be measured by prevention of tumor progression in addition to tumor regression. In a phase II trial evaluating the utility of letrozole in a selected group of ER+ patients with recurrent ovarian cancer, a 17% RR was seen by CA125 criteria and 26% of patients had SD at 6 months of treatment (Smyth *et al.* 2007). Response to AI therapy correlated with ER α expression with those patients having the highest expression of ER α having the greatest RR, 33%, compared with 0% RR in those with low expression. In another study, response to letrozole was associated with increased tumor production of aromatase (Walker *et al.* 2007b). The ideal use of AIs may be following initial cytoreductive surgery and adjuvant chemotherapy for stabilization of any residual disease, potentially serving to prolong PFS; however, a trial comparing DFS in ER+ patients treated with AIs vs placebo is still warranted before such therapy can be justified.

More recently, the pure antiestrogen fulvestrant (ICI 182 870) was tested in a phase II study. Multi-recurrent ER+ EOC were treated with single-agent fulvestrant until intolerance or disease progression (Argenta *et al.* 2009). Of the 26 patients in the study, there was one CR (4%), one PR (4%), and nine patients had SD (35%). While the median time to progression was 2 months, two patients remained on treatment for >250 days. The drug was well tolerated and there were no grade 3 or 4 toxicities. Similar to the AI trials, the authors of this study suggest that the clinical utility of this agent may be optimized by its use early in the adjuvant treatment setting or as long-term consolidation therapy after primary disease remission. Sustained benefit in selected patients suggests that a subset of patients may exist that may prove to benefit substantially from this approach. It is most critical to be able to identify this group of patients that is most likely to respond. There are a few studies that have attempted to identify biomarkers for endocrine treatment response in ovarian cancer

(Bowman *et al.* 2002, Smyth *et al.* 2007, Walker *et al.* 2007a) but such studies are clearly in their infancy and will need to be expanded.

PR ligand in EOC

Several large clinicopathological studies have shown that PR expression is associated with an increased OS in EOC (Münstedt *et al.* 2000, Lee *et al.* 2005b, Tangjitgamol *et al.* 2009). One study showed that PR expression was also associated with improved response to chemotherapy (Tangjitgamol *et al.* 2009). These findings, together with *in vitro* studies suggesting that PR induces apoptosis (and potentially senescence) in ovarian cancer cells, strongly suggest modulation of PR levels and/or activity as a form of endocrine treatment of EOC. Niwa *et al.* (2008) reported their findings of the effect of combination medroxyprogesterone acetate (MPA) with primary adjuvant chemotherapy in advanced EOC in 2008. Both PFS and OS were significantly longer in the patients treated with combination MPA and platinum-based chemotherapy compared with the control group. These effects were more pronounced in the group with higher PR expression.

Zheng *et al.* (2007) recently reviewed the utility of PR ligands in ovarian cancer treatment by examining 13 clinical trials that included 432 patients with recurrent or refractory ovarian cancer treated with megestrol acetate or MPA. Ten patients (2.3%) had CR, 21 (4.9%) had a partial response, and 47 (10.9%) had SD. The authors concluded that the efficacy of progestational agents in recurrent EOC has not been established based on the currently available literature (trials are summarized in Table 3).

The antiprogestin, mifepristone, has also been used in the treatment of platinum-resistant ovarian cancer. In 2000, Rocereto *et al.* (2000) reported an overall RR of 26.5%, with 8.8% (3/34) having a CR, including one patient that remained without evidence of disease for over 3 years. These promising results could not be reproduced in a recent phase II trial in patients with recurrent or persistent ovarian, peritoneal, and fallopian tube cancers, in which only one response was seen among the 22 patients (Rocereto *et al.* 2010). Efforts are also ongoing testing mifepristone specifically in endometrioid adenocarcinomas (Ramondetta *et al.* 2009).

The *in vitro* data, and the promising evidence of treatment response in subset of patients, strongly suggest that further investigation of progestins, either alone or in combination with chemotherapy, is warranted. Given recent preclinical data, and basic science findings, one can expect to see drugs that target different PR isoforms (PR-A vs PR-B) and that might target other progesterone binding receptors (i.e. mPRs and PGRMCs).

Role for targeting AR in ovarian cancer?

In vitro studies have shown that androgen increases proliferation of normal human OSE cells and human ovarian cancer cells (Syed *et al.* 2001, Edmondson *et al.* 2002), making targeting AR a promising treatment strategy. There is increasing evidence that AR is not only a great target in prostate cancer but in other hormone-driven tumors, such as breast cancer (Nahleh 2008). However, the use of antiandrogens in the management of ovarian cancer has been limited to several small clinical trials in the recurrent setting. Two trials from the late 1990s evaluated the use of this flutamide, a nonsteroidal drug with antiandrogen properties, in recurrent ovarian cancer; Vassilomanolakis *et al.* (1997) reported a RR and disease stabilization of 4.3 and 8.7% respectively, while Tumolo *et al.* (1994) had slightly higher RRs and SD at 6.3 and 28%. These phase II trials were small, including only 24 and 32 patients respectively. A more recent trial, published in 2007, examined the use of bicalutamide, an antiandrogen, with goserelin, a GNRH agonist, in women with EOC who were in their second or higher disease remission (Levine *et al.* 2007). The use of these agents did not appear to prolong PFS in this group of patients, and there was no association

between AR repeat number, genotype, or haplotype and PFS. Like for ER and PR-targeting trials, the identification of biomarkers of response will be critical. Given recent study by Elattar *et al.* (2012), nuclear expression of AR might be a viable biomarker for androgen sensitivity. DHT treatment of primary ovarian cancer cultures established from ascitic fluid resulted in increased in S-phase, and this was strongly associated with nuclear AR levels. Thus, in summary, AR modulation as a viable treatment option has not been validated at this time; however, additional studies potentially including novel drugs such as abiraterone, and including the use of biomarkers, are needed.

Combination therapies

PR can be induced with estrogen (and with tamoxifen in situation where it functions as an agonist), and several trials have studied the utility of combination hormonal therapy targeting this cross talk. Although no clinical response was seen with combination tamoxifen and progesterone (Belinson *et al.* 1987, Jakobsen *et al.* 1987), a significant clinical response was seen in one trial of 65 women with refractory ovarian cancer treated with sequential ethinyl estradiol and medroxyprogesterone, with a RR of 14% and SD in 20% of patients (Freedman *et al.* 1986). These findings were later confirmed with a similar although smaller clinical trial (Fromm *et al.* 1991).

Several small phase II trials have looked at the addition of goserelin to tamoxifen, again in the recurrent EOC setting. One trial reported one CR (3.8%), two PR (7.7%), and ten patients with SD (38.5%) with a median progression-free interval (PFI) of 4 months and OS of 13.6 months for the entire cohort (Hasan *et al.* 2005). A similar trial published 6 years earlier had similar findings with a PFI of 5 months and OS of 8 months (Hofstra *et al.* 1999).

Few trials have examined the utility of combining hormonal therapy to chemotherapy vs chemotherapy alone in the adjuvant setting for advanced ovarian cancer. There was no difference in progression-free or OS in 100 women with stage III or IV EOC randomized to receive combination cisplatin/doxorubicin with or without the addition of tamoxifen (Patel & Rothenberg 1994). However, tamoxifen was only administered during chemotherapy (median time: 36 weeks), and >54% of patients had residual tumor \geq 2 cm. Additionally, hormone receptor status was only determined in 72% of patients. The same clinical question was asked in the recurrent disease setting. In one trial with 50 patients with recurrent disease, treatment with a platinum agent and tamoxifen showed a RR of 50% (Benedetti Panici *et al.* 2001). The high RRs in platinum-resistant disease lead other investigators to conduct similar trials. However, similar results have yet to be replicated, and in one trial of 14 patients with platinum-resistant disease, there were no objective responses to combination carboplatin/tamoxifen (Johnson *et al.* 1993, Markman *et al.* 2004b).

Randomized trials of combination progestin therapy and chemotherapy vs chemotherapy alone have also been plagued by small patient sample size, unknown tumor hormone receptor status, and heavily pretreated disease. The development of chemoresistant disease is a big clinical problem, and while still little understood, increased expression of P-glycoprotein, a transmembrane energy-dependent drug efflux pump, contributes to resistance in some tumor models (Johnson *et al.* 1993, Patel & Rothenberg 1994). Although several preclinical models showed the ability of progestins to overcome P-glycoprotein-mediated multidrug resistance (Yang *et al.* 1989, Fleming *et al.* 1992, Wang *et al.* 1994, Panasci *et al.* 1996, Tansan *et al.* 1997), there were no objective responses observed in 44 patients with paclitaxel-refractory ovarian cancer treated with combination paclitaxel and megestrol acetate (Panasci *et al.* 1996). Unfortunately, response to combination MPA-cytotoxic chemotherapy as first-line therapy also showed no difference in RR or survival compared with chemotherapy alone in 71 patients with advanced ovarian cancer (Tansan *et al.* 1997).

Finally, a number of biological agents are being studied for the treatment of ovarian cancer, some in combination with hormonal therapy. There is recent *in vitro* evidence that tamoxifen resistance may be related to EGFR overexpression in solid tumors like breast (Shou *et al.* 2004) and that combination treatment with gefitinib (Iressa), a signal transduction inhibitor of EGFR tyrosine kinase, and tamoxifen was more effective than either agent alone (Hiscox *et al.* 2004). Additionally, there is evidence that EGFR overexpression is seen in 30–70% of cases of platinum-resistant ovarian tumors (Scambia *et al.* 1995, Bartlett *et al.* 1996). Therefore, a phase II trial of combination tamoxifen and gefitinib was conducted in 56 patients with ovarian cancer refractory or resistant to platinum- and taxane-based therapy. Although no tumor responses were seen, 16 patients had SD (Wagner *et al.* 2007). Gefitinib has also been used in combination therapy with the AI anastrozole; in patients with asymptomatic Müllerian cancer, there was a RR of 3% (1/35 patients) (Krasner 2007). Although this number of responders is disappointingly low, the authors comment that the single responder had a durable CR and an additional patient had SD for more than 600 days.

Ongoing clinical trials

At the time of writing of this review, we identified one clinical trial involving the use of hormonal modulation for the treatment of ovarian cancer, which is actively enrolling patients. NCT01273168 is a phase I clinical trial sponsored by the National Cancer Institute (NCI) of endoxifen, a potent and active metabolite of tamoxifen that may have clinical utility as an antiestrogen (Wu *et al.* 2009). It is administered to adults with hormone receptor-positive solid tumors including breast, dermoid, and gynecologic tumors that did not respond to standard treatment. Study participants will take daily oral endoxifen in escalating doses at 28-day intervals to determine safety, tolerability, pharmacokinetics, and determine the maximum tolerated dose. Additionally, an investigational radiolabeled imaging agent, 16- α -[(18)F]-fluoro-17 β estradiol, will be used with positron emission tomography (PET) to determine *in situ* tumor ER activity before and after treatment with endoxifen. Another NCI trial, NCT00445887, is not examining the role of hormonal therapy in the treatment of ovarian cancer but instead is evaluating the effect of oral levonorgestrel on prevention of ovarian carcinoma in high-risk patients.

Several other clinical trials in recurrent ovarian cancer patients have completed enrollment; however, results have not yet been published, including NCT00003865, assessing toremifene citrate, a SERM, and NCT00181688, a phase II trial of combination of AI anastrozole and gefitinib.

Conclusion: endocrine treatment in ovarian cancer

In summary, there is strong *in vitro* and *in vivo* evidence, and epidemiological data showing that estrogens (and potentially other steroids) regulate ovarian carcinogenesis. In addition, ER is highly expressed in subsets of ovarian tumors. Clinical trials testing endocrine therapy in ovarian cancer to date have been poorly designed and not properly evaluated to allow solid conclusions. Many did not select on ER status, and few correlated ER status with response. While the overall RR to hormonal therapy in ovarian cancer appears to be modest, there are clearly subgroups of patients that respond very well. As hormonal agents have the benefit of easy administration, low cost, and data on long-term effects of prolonged use, we must identify such subgroups of patients! Thus, the identification of biomarkers predicting endocrine treatment response and their introduction into clinical practice for both selection of patients for enrollment in clinical trials and subsequently for personalized endocrine treatment are critical. To accomplish such goals, we must continue to investigate mechanism of action of hormones in ovarian cancer, we must perform additional well-designed clinical trials using biomarkers, and we must collect tissue.

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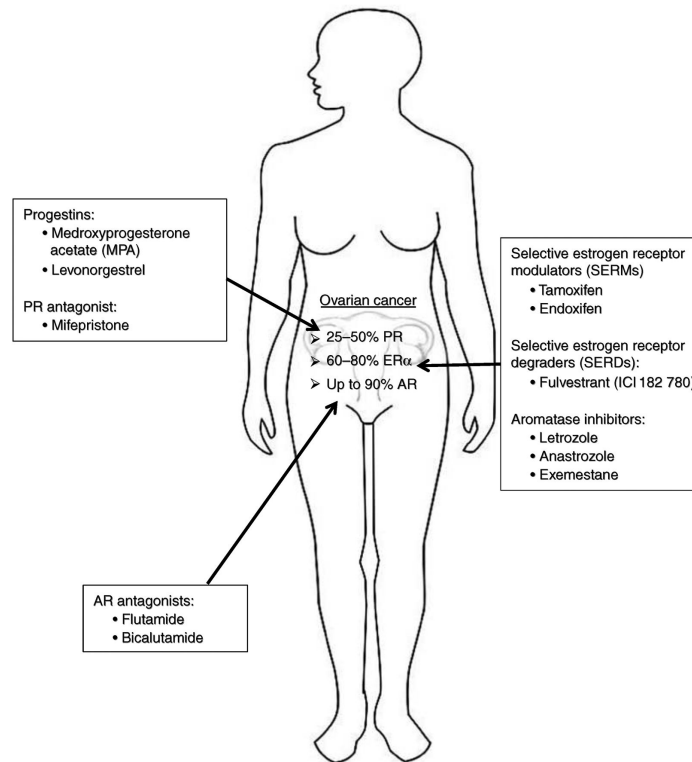


Figure 1. Steroid receptors as clinical targets in ovarian cancer. Estimates of % tumors expressing PR, ER α , or AR (without considering subtypes of ovarian cancer due to lack of information). Also listed are approved drugs or drugs used in previous or currently ongoing ovarian cancer trials.

Table 1

Hormonally related epidemiological factors associated with EOC risk

	Strength of relationship
Factors associated with increased EOC	
Reproductive factors	
Nulliparity	+++
Exogenous hormone use	
Combined HRT	+
Estrogen-only HRT	++
Androgens	+
Reproductive disorders	
Endometriosis	++ ^a
PCOS	+/0
Infertility	++
Other	
Age	+++
Factors associated with decreased EOC risk	
Reproductive factors	
Pregnancy	---
Breast-feeding	-
Twinning and other non-singleton births (note 3)	-
Tubal ligation	---
Hysterectomy	--
Exogenous hormone use	
Oral contraceptive use	---
High vs low progestin OCs	-
High vs low estrogen OCs	0
Factors not associated with EOC risk	
Reproductive factors	
Early menarche	0
Late menopause	0
Exogenous hormone use	
Fertility drug use	0

+, positively associated with EOC; -, negatively associated with EOC; 0, no association with EOC.

^aEndometrioid and clear cell subtype only.

Table 2

Response rates of tamoxifen (TAM) in persistent or recurrent EOC

References	Trial	n	Drug dosage	Number of patients responding (%)					Other
				CR	PR	SD	PD		
Marth <i>et al.</i> (1997)	Nonrandomized phase II	65	TAM 30–40 mg PO daily	2 (3)	2 (3)	50 (77)	11 (17)	Recurrent disease, refractory 1 chemotherapy regime	
Gennatas <i>et al.</i> (1996)	Nonrandomized series	50	TAM 40 mg PO daily	2 (4)	26 (52)	NA	NA	50% previously untreated, 50% refractory or progressive disease	
Jager <i>et al.</i> (1995)	Unblinded randomized controlled trial	33	TAM 30 mg PO daily	0	0	2 (6)	NA	Progressive disease	
Van Der Velden <i>et al.</i> (1995)	Nonrandomized series	30	TAM 40 mg PO daily	2 (7)	NA	10 (33)	NA	Refractory or recurrent disease	
Ahlgren <i>et al.</i> (1993)	Nonrandomized series	29	TAM 80 mg PO×30 d, then 20 mg PO daily	2 (7)	7 (24)	18 (62)	6 (21)	Prior cytotoxic chemotherapy	
Losa <i>et al.</i> (1993)	Randomized trial comparing hormonal therapy (168 patients in three arms)	NA	TAM 40 mg PO daily	0	1	22	32	Prior multi-agent cytotoxic chemotherapy	
Hatch <i>et al.</i> (1991)	Nonrandomized phase II	105	TAM 40 mg PO daily	10 (10)	8 (8)	40 (38)	47 (45)	Recurrent or persistent disease	
Osborne <i>et al.</i> (1988)	Nonrandomized series	51	TAM 100mg/m ² PO over 24 h then 40 mg PO daily	0	1 (2)	0	50 (98)	Disease refractory to 1 chemotherapy regime	
Weiner <i>et al.</i> (1987)	Nonrandomized series	31	TAM 160mg/m ² PO×7d, then 20 mg PO daily	1 (3)	2 (6)	6 (19)	22 (71)	Prior cytotoxic chemotherapy	
Slevin <i>et al.</i> (1986)	Nonrandomized series	22	TAM 20 mg PO daily	0	0	1 (5)	21 (95)	Disease refractory to 1 chemotherapy regime	
Hamerlynck <i>et al.</i> (1985b)	Nonrandomized series	36	TAM 40 mg PO daily	0	2 (6)	7 (19)	NA	Prior cytotoxic chemotherapy	
Landoni <i>et al.</i> (1985)	Nonrandomized series	19	TAM 40 mg PO daily	0	0	7 (37)	NA	Prior cytotoxic chemotherapy	
Shirey <i>et al.</i> (1985)	Nonrandomized series	23	TAM 20–40 mg PO daily	0	0	19 (83)	NA	Disease refractory to 1 chemotherapy regime	
Schwartz <i>et al.</i> (1982)	Nonrandomized series	13	TAM 20 mg PO daily, increase to 40 mg PO daily if disease progression	0	1 (8)	4 (31)	8 (62)	Rapidly advancing recurrent disease	

PO, oral; CR, complete response; PR, partial response; SD, stable disease; PD, progressive disease; NA, not available.

Table 3

Response rates of aromatase inhibitors in persistent or recurrent EOC (all are phase II trials)

References	n	Drug	Number of patients responding (%)				ER status
			CR	PR	SD	PD	
Bowman <i>et al.</i> (2002)	54	Letrozole	0	5 (9)	14 (26)	30 (56)	Mixed ER-/ER+ tumors
Papadimitriou <i>et al.</i> (2004)	27	Letrozole	1 (4)	3 (11)	5 (19)	18 (67)	Mixed ER-/ER+ tumors
Gourley <i>et al.</i> (2006)	33	Letrozole	0	3 (9)	14 (42)	16 (49)	NA
Smyth <i>et al.</i> (2007)	42	Letrozole	0	7 (17)	11 (26)	NA	ER+
Ramirez <i>et al.</i> (2008)	33	Letrozole	0	1 (3)	7 (21)	23 (70)	ER+
Del Carmen <i>et al.</i> (2003)	53	Anastrozole	0	1 (2)	22 (42)	30 (57)	Mixed ER-/ER+ tumors
Verma <i>et al.</i> (2006)	22	Exemestane	0	0	8 (36)	NA	NA

Table 4

Response rates to progestins (MA and MPA) and antiprogestins in persistent or recurrent EOC (all are phase II trials)

References	n (patients)	Drug	Responders (%)		
			CR	PR	SD
Mangioni <i>et al.</i> (1981)	33	MPA	0	5 (15)	2 (6)
	30	MPA	0	0	2 (7)
Slayton <i>et al.</i> (1981)	19	MPA	0	0	1 (5)
Aabo <i>et al.</i> (1982)	27	MPA	0	1 (4)	0
Tropé <i>et al.</i> (1982)	25	MPA	0	1 (4)	9 (36)
Geisler (1985)	23	MA	7 (30)	4 (17)	0
Hamerlynck <i>et al.</i> (1985a)	41	MPA	0	1 (2)	7 (17)
Sikic <i>et al.</i> (1986)	47	MA	1 (2)	3 (6)	5 (11)
Belinson <i>et al.</i> (1987)	33	MA	0	0	12 (39)
Ahlgren <i>et al.</i> (1993)	32	MA	0	0	13 (41)
Malfetano <i>et al.</i> (1993)	24	MPA	0	1 (4)	9 (38)
Veenhof <i>et al.</i> (1994)	54	MA	0	1 (2)	9 (17)
Wiernik (1998)	30	MA	0	0	0
Rocereto <i>et al.</i> (2000)	44	Mifepristone	3 (9)	6 (17)	NA
Wilailak <i>et al.</i> (2001)	36	MA	3 (8)	4 (11)	NA
Rocereto <i>et al.</i> (2010)	22	Mifepristone	0	1	3

MA, megestrol acetate; MPA, medroxyprogesterone; CR, complete response; PR, partial response; SD, stable disease; PD, progressive disease; NA, not available.