

Progesterone and Breast Cancer

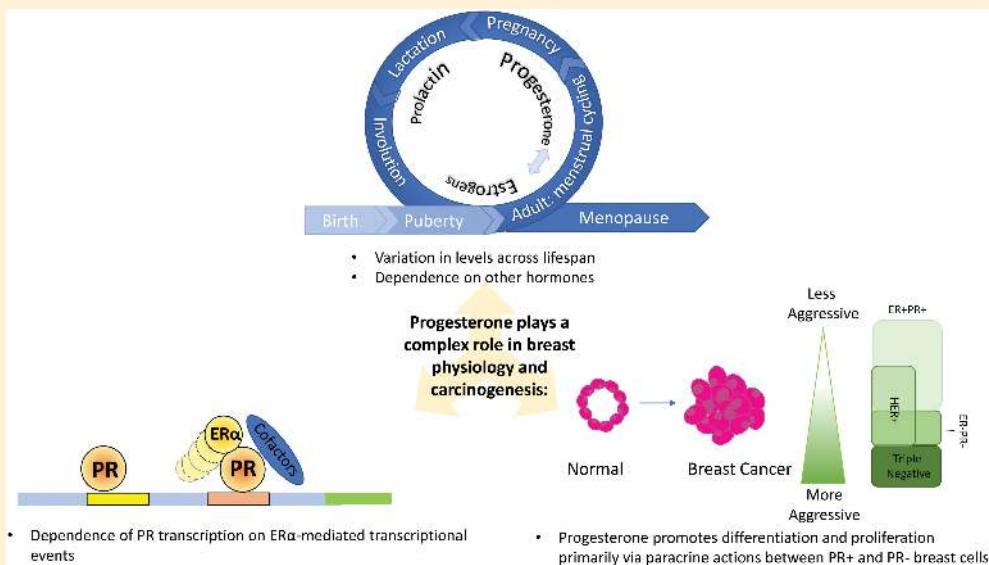
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ABSTRACT Synthetic progestogens (progestins) have been linked to increased breast cancer risk; however, the role of endogenous progesterone in breast physiology and carcinogenesis is less clearly defined. Mechanistic studies using cell culture, tissue culture, and preclinical models implicate progesterone in breast carcinogenesis. In contrast, limited epidemiologic data generally do not show an association of circulating progesterone levels with risk, and it is unclear whether this reflects methodologic limitations or a truly null relationship. Challenges related to defining the role of progesterone in breast physiology and neoplasia include: complex interactions with estrogens and other hormones (eg, androgens, prolactin, etc.), accounting for timing of blood collections for hormone measurements among cycling women, and limitations of assays to measure progesterone metabolites in blood and progesterone receptor isotypes (PRs) in tissues. Separating the individual effects of estrogens and progesterone is further complicated by the partial dependence of PR transcription on estrogen receptor (ER) α -mediated transcriptional events; indeed, interpreting the integrated interaction of the hormones may be more essential than isolating independent effects. Further, many of the actions of both estrogens and progesterone, particularly in “normal” breast tissues, are driven by paracrine mechanisms in which ligand binding to receptor-positive cells evokes secretion of factors that influence cell division of neighboring receptor-negative cells. Accordingly, blood and tissue levels may differ, and the latter are challenging to measure. Given conflicting data related to the potential role of progesterone in breast cancer etiology and interest in blocking progesterone action to prevent or treat breast cancer, we provide a review of the evidence that links progesterone to breast cancer risk and suggest future directions for filling current gaps in our knowledge. (*Endocrine Reviews* 41: 320 – 344, 2020)

GRAPHICAL ABSTRACT



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ESSENTIAL POINTS

- Estrogen and progesterone are sequentially involved in pubertal breast development primarily via a paracrine mechanism
- Preclinical and clinical evidence support differentiation and proliferative roles of progesterone in the adult breast through primarily paracrine actions between PR+ and PR- breast cells
- RANKL expression and high serum progesterone levels are highly correlated in the human breast; further, RANKL expression is required for progesterone-induced proliferation in the breast
- To understand the role of progesterone in initiation and progression of breast cancer, it is prerequisite to understand how epithelial cells of the mammary gland control their fate.
- Hormones are pivotal in controlling physiologic proliferation of normal breast epithelium, and therefore progesterone may influence early events in breast carcinogenesis.
- In breast cancer, the proliferative effect of progesterone is mediated primarily by PR-B; extranuclear signaling actions of PR are also mediated predominantly by PR-B
- Given recent improvements in assay sensitivity, research is needed to evaluate the association between endogenous progesterone/progesterone metabolites and breast cancer risk in the general population and among high-risk women (eg, *BRCA 1* and *BRCA 2* mutation carriers)

The critical importance of ovarian sex steroid hormones (estrogens and progesterone) in promotion and maintenance of the growth of breast cancers was suggested more than 100 years ago when George Beatson demonstrated that bilateral oophorectomy resulted in the remission of breast cancer in a premenopausal woman (1). Subsequently, recognition that many estrogen receptor-positive (ER+) breast cancers are estrogen dependent led to the development of highly effective adjuvant and chemopreventive agents for these tumors. Epidemiologic evidence has also demonstrated that some exogenous synthetic progestogens (progestins) administered with estrogen as menopausal hormone therapy or as contraception increase breast cancer risk. The finding that estrogen+progestin menopausal hormone therapy increases breast cancer risk led to a major decline in long-term use of these agents to alleviate postmenopausal symptoms (2). In contrast to the pharmacologic effects of progestins that have been clearly linked to breast cancer risk, the etiologic role of endogenous progesterone in the development of breast cancer is uncertain.

Mechanistic studies implicate progesterone in the development of breast cancer, whereas limited epidemiologic data have not provided strong support for a risk relationship with circulating levels. Data generated primarily by the Wiebe laboratory (3) suggest that progesterone metabolites have pro- and anti-carcinogenic effects, and that the balance of these factors may contribute to breast cancer risk, but this hypothesis has received limited attention in population-based research to date, primarily due to the lack of available assays. Finally, studies implicate progesterone signaling in the pathogenesis of breast cancers among *BRCA1* mutation carriers

(4), suggesting that chemoprevention to interrupt downstream signaling may be protective.

Complex factors contribute to the challenge of defining the role of progesterone in breast physiology and neoplasia, including: 1) dependence on and interaction with estrogens and other hormones (eg, androgens, prolactin, etc.), 2) variation in exposure levels, eg, during the menstrual cycle or pregnancy, 3) availability of sensitive assays, 4) limited data about the roles of progesterone metabolites, and 5) difficulties in assessing progesterone receptor (PR) isotypes in routinely prepared clinical specimens. Determining individual effects of estrogens and progesterone is complicated because transcription of *PR* is driven partly, but not exclusively, by estrogen receptor (ER) α -mediated transcriptional events (5, 6). Further, many of the actions of both estrogens and progesterone, particularly in “normal” breast tissues, are driven by paracrine mechanisms in which ligand binding to receptor-positive cells evokes secretion of factors that influence cell division of neighboring receptor-negative cells. Thus, levels in tissues and blood may differ and biological effects may be under local control.

Given these conflicting data and the interest in exploring whether blocking progesterone/PR signaling has utility in breast cancer prevention and treatment, we critically review the evidence that links progesterone to breast cancer risk and provide future directions for filling existing gaps in our knowledge. The aims of this review article are as follows: 1) provide background on the formation, transport, and metabolism of progesterone; 2) provide a summary of pharmacologic differences among exogenous progesterone and progestins; 3) describe what is known about physiologic levels of progesterone and breast development across the reproductive lifespan, including during pregnancy; 4) review the

molecular mechanisms related to progesterone action; 5) summarize the current state of knowledge regarding progesterone and breast carcinogenesis; 6) summarize breast cancer risks associated with oral

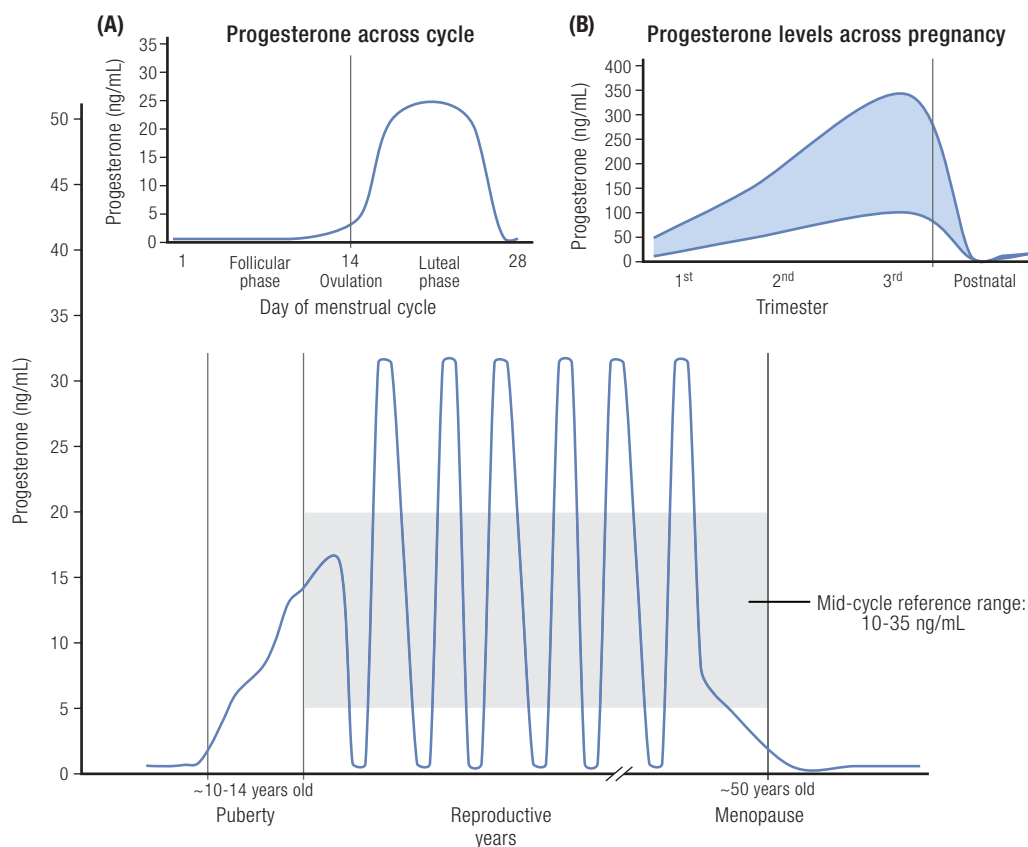
contraceptives and menopausal hormone therapy; and 7) provide a summary of unanswered questions, challenges, and future directions.

Progesterone Formation, Transport, and Metabolism

Progesterone is a 21-carbon steroid that exerts its primary physiological functions through binding to the progesterone receptors A and B (PR-A and PR-B), which initiate transcription of targeted genes resulting in conversion of proliferative endometrium to secretory endometrium in an estrogen-primed uterus. The physiological role of progesterone is primarily confined to the peri- and post-ovulatory phases of the menstrual cycle, and to pregnancy. In the menstrual cycle, progesterone is produced by the corpus luteum beginning in the early postovulatory phase. Its production rate increases from about 1 mg/day in the follicular phase to about 25 mg/day in the luteal phase (Fig. 1)

(7). The biosynthesis of progesterone by the corpus luteum requires continued luteinizing hormone (LH) stimulation. During the follicular phase of the cycle serum progesterone levels are <1 ng/mL (8). The ovarian and adrenal contributions to the total progesterone production rate appear to be equal at that time. Subsequently, progesterone levels rise to about 1–2 ng/mL on the day of the LH surge and continue to rise, reaching a plateau of approximately 10–35 ng/mL during the mid-luteal phase; then levels decline until the end of the menstrual cycle (8). Most women have on average 35 years of predictable menstrual cycles with an average 14-day follicular phase and 14-day luteal phase.

Figure 1. Progesterone levels across the lifespan: [Inset A) across the menstrual cycle, B) across pregnancy]. Legend: Progesterone levels increase from <1 ng/mL before age 10 to 10–12 ng/mL around puberty. Throughout the reproductive years levels range from 3 to 35 ng/mL in the luteal phase, decreasing in the late luteal phase, and are <1 ng/mL in the follicular phase (Inset A). In postmenopausal women, progesterone levels wane to <0.2 ng/mL. In pregnant women (Inset B), progesterone levels increase from 10–35 ng/mL in the 1st trimester to 100–300 ng/mL in the 3rd trimester; progesterone levels then drop to low levels postnatally as prolactin levels increase to facilitate lactation and return to average reproductive age ranges when ovulation returns.



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In contrast to the menstrual cycle, progesterone production is considerably higher in pregnancy. During the first 10 weeks of pregnancy, maternal serum progesterone levels are in the luteal phase range of 10–35 ng/mL. The placenta then begins to secrete sizable amounts of progesterone, resulting in a steady rise throughout the remainder of pregnancy. The production rate of progesterone during the late stage of pregnancy is as high as 300 mg/day. At term, serum progesterone levels range from 100 to 300 ng/mL (8).

As women age, ovarian follicle loss increases exponentially with accelerated loss as a woman enters the menopausal transition. The menopausal transition is characterized by increasing menstrual cycle irregularity, including lengthy periods of anovulation with fluctuation in hormone levels, as women progress over a period of 2–6 years towards having their final menstrual periods (aged 51.4 years on average) (reviewed in (9, 10)). In brief, FSH levels begin to rise on average 6 years before the final menstrual period (11). Estrogen levels remain relatively constant through the early years of the menopausal transition, and then begin to fall about 2 years prior to the final menstrual period, reaching the low stable levels characteristic of postmenopausal women about 2 years after the final menstrual period (9, 12). In contrast to relatively consistent estrogen levels, urinary pregnanediol glucuronide levels in ovulatory cycles (representing progesterone excretion) slowly decline during the menopausal transition, about 7% annually, signaling progressive luteal dysfunction as a characteristic of the menopausal transition (12). Early follicular inhibin B levels, which reflect the diminishing quantity and quality of ovarian follicles, and anti-Müllerian hormone levels consistently decline prior to the menopausal transition, becoming nondetectable approximately 4–5 years before the final menstrual period (9).

In postmenopausal women, the major source of progesterone is the adrenals, which produce less than 1 mg/day resulting in serum progesterone levels that are generally <0.2 ng/mL.

In blood, progesterone is bound to corticosteroid-binding globulin (CBG) and to albumin (13). In cycling women only about 20% of progesterone is bound to CBG, and the rest predominantly to albumin; about 2–3% of progesterone is unbound (free). However, in pregnancy CBG increases considerably and about 40% of progesterone is CBG-bound. It is plausible that the difference in bound and unbound progesterone between pregnant and non-pregnant women could explain some of the breast cancer risk reduction associated with

pregnancy, although to our knowledge this has not been evaluated in epidemiologic studies.

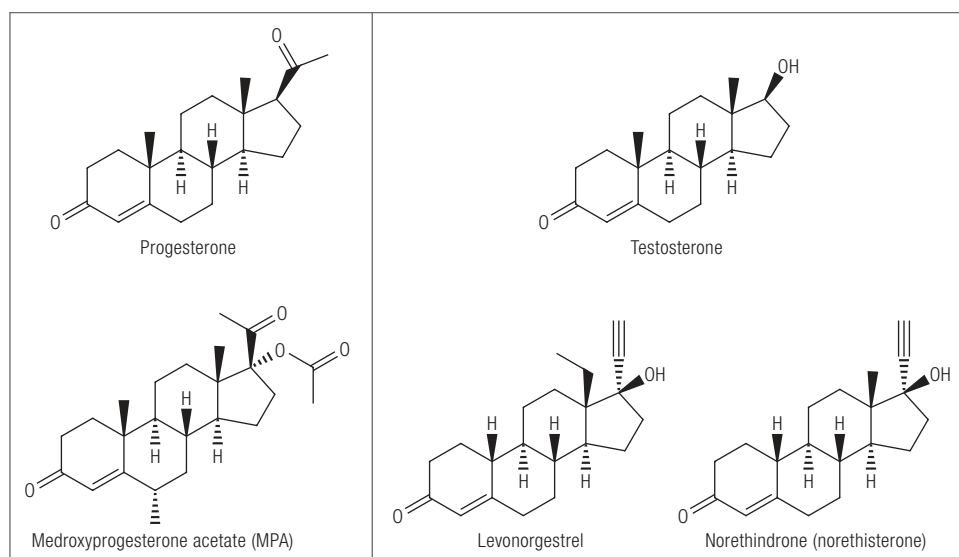
Progesterone is highly vulnerable to enzymatic action by hepatic reductases and hydroxysteroid dehydrogenases because its structure contains two ketone groups and a double bond (14). Thus, the molecule is transformed to two isomers of dihydroprogesterone, four pregnanolone isomers, and eight isomers of pregnanediol. In addition, progesterone can undergo hydroxylation by cytochrome p450 enzymes. Subsequently, all progesterone metabolites with a hydroxyl group can be sulfated and glucuronidated, and their conjugated products are then excreted primarily in urine, but also in feces. Theoretically, progesterone can have over 100 metabolites when the unconjugated, sulfated, and glucuronidated metabolites are combined (summarized in (15)).

In general, progesterone acts on the reproductive tract to prepare it for initiation and maintenance of pregnancy. The major physiologic roles of progesterone are mediated in the uterus and ovary. After ovulation the corpus luteum releases progesterone which stimulates maturation of the uterine lining to facilitate implantation; progesterone maintains pregnancy through stimulation of uterine growth and differentiation and suppression of myometrial contractility.

Exogenously Administered Progesterone and Progestins

Orally administered progesterone has limited biological effects because it is poorly absorbed, even in micronized form, and is metabolized extensively during the hepatic first pass. For this reason and because more potent progestational compounds were needed for effective antifertility treatment, progestins were developed. A variety of progestins are now available not only for contraception, but also for menopausal hormone therapy to prevent endometrial hyperplasia and lessen the risk of endometrial cancer in estrogen-treated postmenopausal women. The progestins can be classified, based on their chemical structures, as those related to progesterone (C-21 progestins) and those structurally related to testosterone (C-19 progestins) (16). An example of a C-21 progestin is medroxyprogesterone acetate (MPA), whereas norethindrone (norethisterone) and levonorgestrel are C-19 related progestins (Fig. 2). Differences in pharmacologic properties of progesterone and some of the commonly used progestins are summarized in Table 1.

Figure 2. Chemical structures of progesterone and testosterone and for comparison common C-21 and C-19 related progestins.



Progesterone's Role in Breast Development

Much of what is known about breast development has been derived from studies of rodents, which provides a useful, albeit imperfect, model for breast development in women. The breast develops at puberty with the establishment of menstrual cycling (17, 18). Estrogen and progesterone are sequentially involved in pubertal breast development primarily via a paracrine mechanism (19–21). Specifically, *in vivo* studies of hormonal ablation via ovariectomy and subsequent hormone replacement suggest that estrogens drive the first stage of pubertal development in the breast, whereas both estradiol and progesterone are responsible for cellular proliferation in the mammary gland (reviewed in (22); references to “mammary” gland throughout this manuscript refer to animal model data). Cell proliferation in the mammary epithelium ceases in ovariectomized mice; however, reintroduction of estradiol is sufficient to induce epithelial cell proliferation and restores ductal outgrowths consistent with intact pubertal animals. Estradiol alone is not sufficient to induce mammary gland cellular proliferation in pregnant animals (23). Similarly, progesterone is not sufficient to stimulate cellular proliferation in the absence of estrogen, as demonstrated by experimental studies conducted in ovariectomized mice showing that PR expression in mammary epithelium is dependent on interactions of estrogen with ER α to induce PR transcription (24–26). As is emphasized throughout this review, progesterone acts primarily through a paracrine mechanism, which makes cell culture experiments challenging. We are not aware of any co-culture systems for luminal progenitor cells that enable testing

amplification of paracrine progesterone signaling. Further, progesterone's paracrine signaling makes clinical translation a challenge as well, given that levels of other circulating hormones/growth factors and their cognate receptors likely influence the response to or effects of progesterone.

The dependence on hormones in mouse models is also relevant in humans, as the development of breast buds (thelarche) corresponds with increasing ovarian estrogen secretion preceding menarche and the introduction of cyclic ovarian progesterone secretion via menstrual cycling (27). In normal breast development in humans, it has been demonstrated that estrogen and progesterone are required for ductal elongation and side branching, respectively, across the reproductive years (28). However, the mouse and human differ substantially in several ways, including different levels of circulating hormones and hormone metabolites, processing by different liver enzymes, and different regulation of aromatase (29).

Androgens, on the other hand, inhibit breast development (reviewed in (30)). Higher absolute levels of androgens (regardless of estrogen level) suppress breast development, while low circulating testosterone relative to normal estradiol levels has been shown to stimulate breast development.

The mature breast is characterized by ducts and side branches (extralobular terminal ducts and lobular buds), but the highly branched architecture and terminal end buds characteristic of pregnancy are generally absent (18). Thus, pregnancy is another important developmental stage for the breast. During early pregnancy, the breast epithelium undergoes rapid and pronounced

Adapted with permission from Stanczyk, F.Z., et al., Progesterone used in postmenopausal hormone therapy: differences in their pharmacological properties, intracellular actions, and clinical effects. *Endocr Rev*, 2013. 34(2): p. 171–208.

Table 1. Pharmacologic difference among progestogens

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|-----------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 1. Progesterone | <ul style="list-style-type: none"> • A natural progestogen (21 carbons) • Undergoes extensive metabolism during hepatic first pass • Approximately 20% is bound to CBG • Has low bioavailability (<5%) • Half-life is 16–18 hours after oral dosing • Binds with relatively high affinity to the PRs and MR • Typical daily dose used for endometrial protection is 200 mg |
| 2. Medroxyprogesterone acetate | <ul style="list-style-type: none"> • Structurally related to progesterone • Acetate group at the carbon 21 position limits metabolism (steric hinderance) • Does not bind to CBG or SHBG • Has high bioavailability (>90%) • Its half-life is ~22 hours • Binds with high affinity to the PRs • Has relatively high receptor/ligand binding affinity (RBA) for the AR but its androgenic activity is controversial • Has high RBA for the GR, displays higher binding affinity for the GR than cortisol • Common daily doses for endometrial protection are 2.5 and 5 mg |
| 3. Norethindrone (Norethisterone) | <ul style="list-style-type: none"> • Structurally related to testosterone • Ethinyl group at carbon 17 limits metabolism; can be converted to ethinyl estradiol • Binds to SHBG • Its bioavailability is ~64% and half-life is ~8 hours • Binds with relatively high affinity to the PRs and has some androgenic activity |
| 4. Levonorgestrel | <ul style="list-style-type: none"> • Structurally related to testosterone • Metabolism is very limited during hepatic first pass • Binds to SHBG • Its bioavailability is 89–99% and half-life 10–13 hours • It binds with high affinity to the PRs and AR, and has substantial androgenic activity |
| 5. Desogestrel | <ul style="list-style-type: none"> • Structurally related to testosterone; it is prodrug that is rapidly converted to etonogestrel during hepatic first pass • Etonogestrel binds to SHBG, has a bioavailability of 62–76% and half-life of 12–24 hours, and has a high affinity for the PRs |
| 6. Drospirenone | <ul style="list-style-type: none"> • Structurally related to spironolactone • Does not bind to SHBG or CBG • Has bioavailability of ~66% and half-life of ~32 hours • Has a relatively low RBA for the PRs but a very high RBA for the MR; exhibits both anti-androgenic and anti-mineralocorticoid properties |

expansion and physiologic differentiation to form the highly branched architecture (lobular buds into intralobular ducts and ductules) of the breast. This expansion facilitates further differentiation of terminal duct lobular units (TDLUs) during late pregnancy to prepare for high levels of milk production resulting in lactation (31, 32). Similarly, studies of human breast tissues donated for research show that parous women have a greater density of breast lobules than nulliparous women and that this effect may be greatest within 10 years of a live birth (33).

Progesterone is critical for inducing ductal side-branching of the mammary gland which is essential for lobuloalveolar development during pregnancy (reviewed in (22)). PR null mouse mammary glands fail to undergo alveolar morphogenesis during pregnancy (28); in contrast, forced over-expression of PR results in increased ductal side-branching in adult virgin mice (34). It has further been demonstrated in selective knockout models in mice that PR-B is the primary mediator of progesterone's proliferative effects during pregnancy (35, 36). Additional data indicate that progesterone mediates the secretion of

Wnt4 from PR+ mammary cells, further supporting progesterone's paracrine mode of action to regulate side-branching during early pregnancy (37). Additional data also suggest that progesterone acts directly on the ER α /PR+ progenitor cell population by inducing receptor activator nuclear transcription factor kappa B ligand (RANKL) secretion from these mammary cells. This factor then directly binds to its receptor, RANK (also referred to as tumor necrosis factor superfamily 11A (TNFRSF11A)), to induce side branching and alveolar development (35, 38, 39). As mentioned previously, given that progesterone acts primarily through a paracrine mechanism, cell culture experiments are challenging. Co-culture systems for luminal progenitor cells are needed to allow testing amplification of paracrine progesterone signaling. Inevitably, as research evolves to test context-dependent signaling and transcriptional events involving multiple receptors and complex mixtures of ligands, our understanding of the role of hormones in breast development and cancer will improve. This information will be necessary to translate the existing laboratory and epidemiologic research into clinically actionable treatments and provide context on whether progesterone agonists or antagonists may be more useful.

Molecular Mechanisms Related to Progesterone Action

Progesterone Receptors

Steroid hormones, including progesterone, function by binding to cytoplasmic and nuclear proteins. Progesterone binds to two predominant PR isoforms, PR-A and PR-B. These isoforms are transcribed from the same gene (*PGR*) by two distinct promoters, which have different transcriptional and functional activities. PR-B has an additional stretch of amino acids located at the amino-terminus of the receptor. PR-B can encode a transactivation function that is specific to the PR-B protein, which plays an essential role in specifying target genes that can be activated by PR-B protein but not PR-A protein (6, 40, 41).

Reproductive tissue levels of PR-A and PR-B and their ratio varies based on developmental stage and hormonal status of the tissue (42, 43). When PR-A and PR-B are equivalent in cells, the receptors can dimerize and bind DNA as three species: A:A, B:B, and A:B (heterodimer) (44). PR-A and PR-B are capable of binding progesterone, dimerizing, and interacting with the progesterone response element and transcriptional machinery to regulate gene expression (41, 45). Mediation of

the regulatory effects of progesterone depends on differential transactivation properties contributed to these complexes by the PR-B-specific domain (45–47). Further, the ratio of PR-A to PR-B in target cells as well as dimerization likely predict the overall cellular response to progesterone (45, 48).

Complicating research in this area, isoforms PR-A (94 kDa) and PR-B (114 kDa) can be readily distinguished by western blot due to their significant size difference of 20 kDa, but owing to similarities in structure, antibodies have failed to reliably distinguish them by immunohistochemical methods. Antibodies originally found to immunohistochemically detect the PR-A isoform alone have been found later to detect both PR-A and PR-B, and sample preparation methods appear to be important for PR-B epitope exposure. Recently, a monoclonal antibody Ab-6 (Thermo Fisher) has been described to uniquely detect PR-B due to its likely binding to the extra-stretch at the N-terminal (49) although this remains to be validated in larger studies including demonstrating consistency of the results across pathology laboratories.

Ligand-occupied PR binds to DNA and recruits pro-regulatory proteins (coactivators or corepressors) that interact with the transcription apparatus to modulate gene expression. When bound to ligand, PR controls the transcription of genes, which in the breast include genes for ER, insulin receptors, epidermal growth factor (EGF) and its receptors, and transforming growth factor (TGF)- α . Thus, progesterone is theorized to indirectly affect proliferation by modulating levels of growth factors and their cognate receptors (6). Progesterone may also modulate transcriptional activity via interaction with other transcription factors or by preventing access of transcriptional activators to DNA regulatory regions (squenching). Transcription of PR target genes can be affected by synthetic PR ligands through selective recruitment or blockade of regulatory cofactors. This action can result in differential regulation of gene expression in various progesterone target tissues (50). It has been shown in reproductive cancer cell lines that PR target genes, including *CCND1*, *CDKN1A*, *DUSP1*, *EGFR*, and *PGR*, are dependent on *MAPK* activity (51).

Progesterone Signaling via the PR in the Normal Breast

Role of PR-A and PR-B

As demonstrated in mouse models, estrogen and progesterone act sequentially on mammary epithelium to signal mammary gland development.

With respect to hormone receptor signaling, estradiol and epithelial ER α signaling are necessary for ductal elongation during early puberty (52–54); further, it has been demonstrated that progesterone/PR are not necessary for this stage of early breast development (55). Increased estrogen levels induce PR expression; this is known as ‘estrogen priming’ and is common to most progesterone target tissues. PR signaling by progesterone is required in the epithelial compartment for side branching and alveologenesis (28, 55–57). Recent studies demonstrate different roles of PR-A and PR-B in estrogen signaling and ER chromatin binding (58–60). More specifically it has been demonstrated that PR-B is uniquely required for the latter stage of breast development (ie, side branching and alveologenesis) (35, 36). Thus, as the mouse reaches maturity, cyclic progesterone exposure results in ductal development and dichotomous branching that fills the mammary fat pad. During pregnancy, progesterone signaling via PR and prolactin signaling via the prolactin receptor (PrlR) in the epithelial compartment are required for branching and alveolar proliferation and differentiation (61, 62).

Paracrine mechanism

Specific progesterone effects that are distinct from estrogen and vice versa remain to be delineated; the co-expression and functional interdependence of these hormones poses challenges for distinguishing their respective functions. A paracrine mechanism is suggested to mediate estrogen and progesterone-induced proliferation of mammary epithelial cells. Studies supporting a paracrine mechanism for progesterone have demonstrated that approximately 40% of adult mouse mammary epithelial cells are PR+ and largely non-proliferative. These PR+ epithelial cells are located next to or near PR- cells that are proliferative (63, 64). Direct evidence derives from a study that demonstrated the return of proliferation in transplanted PR knockout mammary epithelial cells that were in close proximity to wild type PR+ mammary epithelial cells (28). The majority of PR+ epithelial cells are also ER α + in the adult mouse mammary. It has been demonstrated in the mouse model and suggested in humans that PR upregulated target gene expression of ER and estradiol are required to maintain high expression of PR in mammary epithelium. Recently, it was also shown that PR-B dominantly activates more ER target gene expression than PR-A, whereas PR-A represses PR-B, ER, and other steroid receptors (reviewed in (41)). Further, excess expression of either PR-A or PR-B has been shown to

disproportionately affect mammary gland development leading to increased lateral ductal branching or inappropriate lobulo-alveolar growth, respectively (34, 64).

Preclinical and clinical evidence support differentiation and proliferative roles of progesterone in the adult human breast through primarily paracrine actions between PR+ and PR- breast cells. Evidence from mouse models also suggests that expansion of the mammary stem cell population is driven by progesterone and PR signaling (20, 65). Consistent with this finding, research shows the expansion of stem cells, measured as an increase in the number of bipotent cells, after progesterone stimulation of normal human breast cells in matrix-embedded culture (66–68). However, it is not clear whether expression of PR isoforms differentially affects stem cell expansion in humans; in mouse models it is suggested that PR-A expression is restricted to luminal cells and PR-B expression is present in both luminal cells and a proportion of basal and mammary stem cells (20). Thus, the experimental data provide support that regulated expression of both PR-A and PR-B is critical for the mammary gland (and potentially human breast tissue) to respond appropriately to progesterone. This further highlights the need for accurate methods to quantify PR-A and PR-B expression in tissue that can be utilized in a clinical setting.

RANK pathway

The RANK pathway acts as the primary mediator of progesterone-driven proliferation in mammary epithelial tissues. Selective induction of high level *RANKL* expression in mammary epithelial tissues is associated with the high progesterone phase of the reproductive cycle and at this stage co-expression of PR is found in nearly 100% of *RANKL*+ mammary cells (69). *RANKL* was believed to work solely through *RANK* to provide developmental signaling that promotes mammary alveologenesis essential for lactation (70–72). Recently, leucine-rich repeat-containing G-protein coupled receptor 4 (*LGR4*), a receptor for R-spondins (RSPOs), has been shown to be a second *RANKL* receptor (73). *LGR4* knockout mouse phenocopies *RANK* knockout, ie, mammary glands of *LGR4* knockout mice have delayed ductal development, fewer terminal end buds, and decreased side-branching. It has been suggested that loss of *LGR4* leads to decrease in Wnt signals important for alveologenesis (74). Both loss and overexpression of *RANK* in mice prevents development of pregnancy-induced milk-secreting structures, suggesting a possible dual role of *RANK* signaling in regulation of alveologenesis

in pregnant mice (23, 70, 71, 75, 76). During pregnancy, *RANKL* expression is upregulated in mammary epithelial cells and is essential for development of the lobulo-alveolar mammary structures and the formation of a lactating mammary gland (35, 70, 75).

Hormones can regulate the expression of *RANKL* protein or mRNA in the human breast specifically; this regulation has been demonstrated for progesterone, estradiol, and prolactin (70, 77, 78). Increased *RANKL* mRNA expression and high serum progesterone levels are highly correlated in fine-needle aspirate samples from normal human breast; further *RANKL* expression is required for progesterone-induced proliferation in the breast (78). *RANKL* protein or mRNA expression in normal human breast tissue is higher in high progesterone conditions, ie, during pregnancy and during the luteal phase of menstrual cycle, as well as in women on combined hormone therapy. Expression of both *RANKL* mRNA and protein is induced by prolactin and parathyroid hormone protein-related peptide; *RANKL* expression is higher in luminal mammary cells of pregnant mice vs. virgin mice (19). However, estrogens, (either in the presence of a natural estrogen-rich environment (70) or via injection of estradiol and progesterone (20)) in conjunction with progesterone are required to induce *RANKL* mRNA and protein expression in mammary tissue (mouse). In normal human breast tissue, higher estradiol levels induced *RANKL* expression in the context of both high (luteal phase) and low (midcycle) progesterone levels (78), further supporting the complicated regulatory mechanisms of increased *RANKL* expression with respect to menstrual cycle hormonal fluctuations. Evaluating characteristics or patterns of altered *RANKL* expression may help clarify the divergent breast cancer risk across women with normal menstrual cycles. Osteoprotegerin is a soluble member of the TNF receptor superfamily that regulates osteoclastogenesis (79); it is secreted by osteoblasts and acts like a cytokine. Osteoprotegerin is a decoy receptor for *RANKL*, and has been shown to be produced by breast cancer tumors and to promote tumor growth and metastasis (reviewed in (80)). *In vitro* osteoprotegerin binds to *TRAIL* as a decoy receptor and prevents cell death. Higher circulating osteoprotegerin levels have been associated with increased risk of receptor negative breast cancer and a possible inverse relationship with receptor positive breast cancer (81), although limited to one study. It has been suggested that the inverse relationship with hormone receptor positive breast cancers is due to interference with *RANKL* signaling. This

hypothesis is supported by *in vitro* data showing similar results in human breast tumors samples exposed to osteoprotegerin and *RANKL* (78). Further, lower circulating levels of osteoprotegerin has been observed in *BRCA* carriers in general, while a recent study reported an increased breast cancer risk with low osteoprotegerin levels in 206 *BRCA* carriers among whom 18 incident breast cancers were observed (82, 83).

Membrane-Bound Progesterone Receptors (“non-genomic” mechanisms of progesterone action)

The nuclear receptors described above (section IV.A.) are the key mediators of the biologic effects of progestogens. However, progesterone has also been shown to elicit its biologic effects via non-genomic mechanisms through activation of signal transduction pathways that are mediated by cell membrane associated PRs distinct from the classical PR (84, 85). The “non-classical” effects of progesterone can be elicited rapidly in various tissues, in contrast to the classical PR-mediated effects which require time to induce transcription and translation of genes into protein products. Among the rapid non-nuclear signaling pathways known to be activated by progesterone are the following: extracellular signal-regulated kinase (ERK) pathways, cyclic AMP/protein kinase A (PKA) pathway, cyclic GMP/protein kinase G (PKG) pathway, Ca⁺⁺ influx/protein kinase C (PKC) activation pathway, and the phosphoinositide 3-kinase (PI3K)/Akt pathway (86).

Two types of distinct cell surface associated proteins unrelated to the classical PRs have been identified: membrane PRs (mPRs) and the progesterone receptor membrane component 1 (PGRMC1). The mPRs have a molecular mass of approximately 40 kDa and are comprised of different subtypes. In contrast to the mPRs, PGRMC1 is part of a multiprotein progesterone-binding complex that has multiple functions including activating cytochrome P450 enzymes that metabolize steroid hormones (87–89). PGRMC1 is expressed in breast tissue and several other tissues and is suggested to mediate the antiapoptotic effects of progesterone (88–92). PGRMC1 expression in blood cells does not vary across the menstrual cycle but was found to be downregulated in postmenopausal women, women with premature ovarian failure, and women with polycystic ovary syndrome (93).

PGRMC1 is overexpressed in breast cancer. It has also been reported that MPA, but not progesterone, increases cell viability and proliferation

in the context of low estradiol levels, and that the proliferative effects are differential by PGRMC1 expression, with a more pronounced effect observed for PGRMC1 overexpressing cells compared with low PGRMC1 expressing cells (94, 95). However, it is unclear if this can entirely explain the absence of increased breast cancer risk with the menopausal hormone therapy formulations, estrogen-progesterone and estrogen-dydrogesterone (structurally similar to progesterone) compared to other menopausal estrogen-progestin formulations that are associated with transient increases in breast cancer risk among current and recent users (96).

Progesterone and Stem Cell Fate in the Breast

To understand the role of progesterone in the initiation and progression of breast cancer, it is prerequisite to understand how epithelial cells of the mammary gland control their fate. Mammary development starts from an embryonic mammary anlage that differentiates through coordinated stages to generate a diverse set of self-renewing, transitional, and terminal cell types (97, 98). By adulthood the normal adult female mammary gland has become a continuous, branching epithelial structure with a distinct outer basal/myoepithelial layer and an inner luminal epithelial layer. The ability to enzymatically dissociate these cells into a suspension of viable single cells and separate them by multicolor flow sorting into subpopulations has led to a major leap in our understanding of the biology of the mammary gland (99, 100).

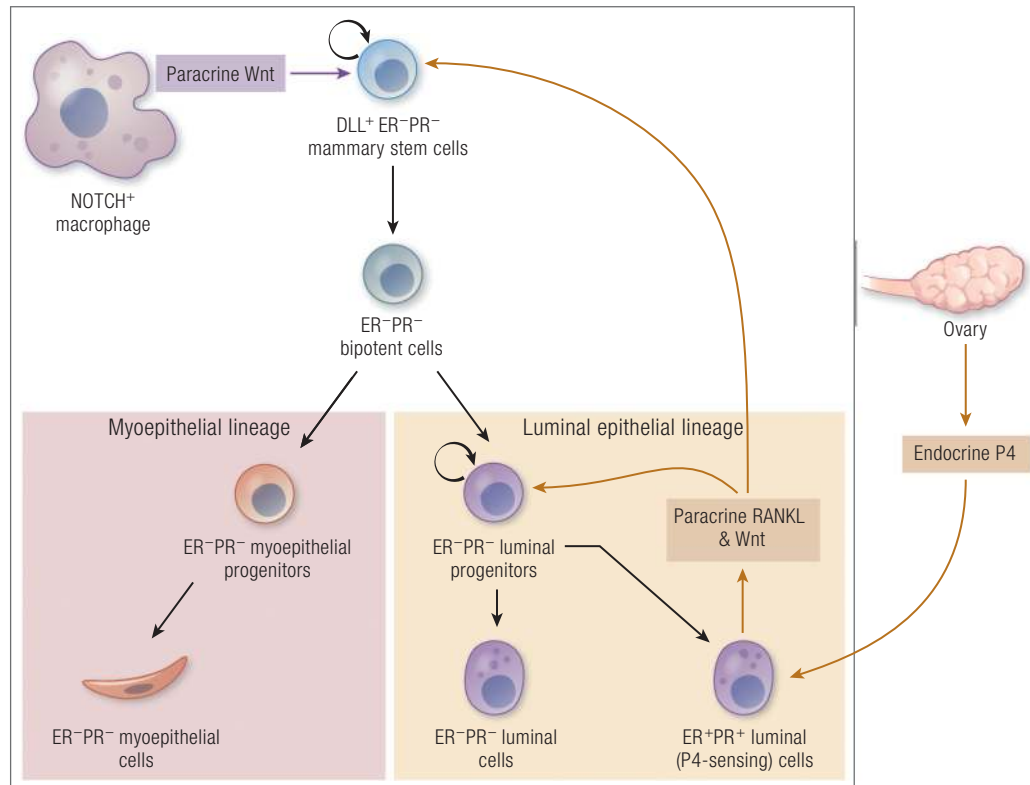
In humans, the CD49f+/EpCAM-low basal cells contain all of the stem/bipotent/myoepithelial progenitor activities as well as differentiated myoepithelial cells (101). The CD49f+/EpCAM+ luminal progenitor cells contain luminal lineage-restricted progenitors at an approximately 30% purity with the other 70% of the cells in this fraction having undefined activities. The entire CD49f-/EpCAM+ luminal cell fraction is devoid of cells with clonogenic activity (102). A growing number of studies suggest that luminal progenitors are naturally at risk for acquiring cancer-predisposing mutation(s) due to the presence of a number of mechanisms that are contributors of genome instability, namely chronic oxidative stress (103), telomere dysfunctional state (102), and elevated R-loop structures (104).

In humans, immunomagnetically purified *in vitro* transformed EpCAM+ luminal cells and CD10+ basal/myoepithelial cells were shown to generate ER+ and ER- subtypes and a metaplastic subtype, respectively, which resembled human breast cancers (105). With fluorescence-activated

cell sorting (FACS) purified subsets of human breast epithelial cells, it was further shown that a single oncogene was sufficient to de novo transform basal cells and luminal progenitor subsets and not terminal luminal cells in highly immunodeficient mice providing new insight into the mechanism of breast cancer initiation (106). In mice, activation of the mutant *PIK3CA* gene in a cytokeratin-5 expressing basal/myoepithelial lineage generated only luminal ER+PR+ tumors, whereas its expression in cytokeratin-8 expressing luminal lineage cells generated both luminal ER+PR+ and basal-like ER-PR- tumors (107). Interestingly, neither stem cell-rich basal nor luminal progenitor-rich fractions express PR. The ER and PR are commonly found to be co-expressed within a developmentally mature subset of non-cycling cells in the luminal cell-fraction. According to the stem cell model, the ER+PR+ luminal cells act as a conduit for ovarian steroids to regulate more primitive mammary cells in the basal cells and luminal progenitor fraction and in turn control their own numbers (20). The ER+PR+ luminal cells, also referred to as P4 sensors, directly respond to progesterone and express mitogenic signals namely RANKL and Wnt, which activate RANK and Wnt pathways in more primitive mammary cells (19, 20, 108) (Fig. 3), while progesterone-mediated RANKL/Wnt signals are necessary but not sufficient to drive mammary development (109). A recent mouse study found that the NOTCH proteins in mammary-resident macrophages interact with their Delta-like ligand (DLL) expressed by mammary stem cells and produce several Wnt signals additionally required to reach the necessary threshold for mammary cell expansion during the menstrual cycle (109, 110). The highly complex multistep control of mammary cell expansion mediated by multiple cell types, using endocrine and paracrine signals, suggests that the risk mechanisms associated with breast cancers need to be investigated in humans to understand the full impact of progesterone signaling on cancer-susceptible primitive human breast cells. Owing to the recently acquired knowledge of the complexity of paracrine mechanisms through which progesterone controls mammary tissue turnover, the downstream gene targets of indirect effector NFκB specifically regulated by progesterone has not been resolved yet. Understanding these gene targets may be useful in identifying possible antibodies/treatments that could be tested in the context of either prevention or treatment/prognosis.

Recent studies examining the epithelial subpopulations in breast-cancer-susceptibility

Figure 3. Progesterone regulation of functionally-defined mammary epithelial cell hierarchy.



gene *BRCA1* mutation carriers have found abnormally high numbers of functionally intact subsets of luminal progenitor cells with RANK expression and derailed progesterone regulation, and with high levels of persistent DNA damage. This abnormality, linked to breast cancer risk in *BRCA1* carriers, is attributed to progesterone mediated RANKL expression by the terminal ER+PR+ luminal cell fraction (111, 112). Therefore, dismantling luminal cell/luminal progenitor communication via progesterone/RANKL with RANKL-neutralizing antibody denosumab (Genentech) is perceived as a strategy to control luminal progenitor numbers and possibly prevent breast cancer in *BRCA1* mutation carriers (111, 113). An international trial to repurpose denosumab as a breast cancer precision prevention drug in a genetically pre-defined subset of high-risk patients (namely *BRCA1/2*) is currently underway. While this is a highly promising development, more caution is warranted since the knowledge of the extent of denosumab exposure and its response among the *BRCA1/2* carriers with distinct mutations will determine the outcome of the first precision prevention trial for breast cancer.

Progesterone and Lobule Involution Among Women

Breast lobules are microscopic structures that represent the main source of breast cancer precursors and with aging these structures undergo simplification and obsolescence (33). Studies of women who have undergone benign breast biopsies demonstrate that reduced levels of TDLU involution are associated with increased breast cancer risk, independent of other breast cancer risk factors (32, 114, 115). Analyzing the association between serum hormone concentrations among premenopausal women and levels of TDLU involution of normal breast tissue donated for research demonstrated that higher progesterone levels in the luteal phase were associated with fewer lobules per unit area [odds ratio (OR) 0.80, 95% confidence interval (CI) 0.72–0.95; $p < 0.0001$] (116). However, this analysis was based on 237 premenopausal women whose samples were evaluated with chemiluminescent immunometric assays, which may lack the sensitivity and specificity of newer LC-MS/MS assay methods (117). In the same study (116), analyses of postmenopausal women were limited by a smaller sample size ($n=148$) and low assay sensitivity (large number of samples below assay detection limit), which precluded quantitation.

Progesterone Signaling via the PR in Breast Cancer Development

Equal amounts of PR-A and PR-B are present in healthy adult breast tissue and the same is true in benign breast lesions, whereas this ratio is altered in breast cancer (reviewed in (118)). In the breast, it is suggested (and supported with mouse model data) that progesterone stimulates normal human breast epithelium through a paracrine mechanism (48, 118) and is a risk factor for breast cancer, because it promotes pre-neoplastic progression via stimulation of cyclical proliferation of mammary stem cell pools or occult tumor initiating cells in the mature breast epithelium (118). It is further suggested that cancer progression is a result of progesterone/PR signaling and a switch from paracrine to autocrine regulation of proliferation (57, 118, 119). In breast cancer, the proliferative effect of progesterone is mediated primarily by PR-B. Extranuclear signaling actions of PR are also mediated predominantly by PR-B (120). It is unclear if extranuclear signaling actions of PR occur in normal breast tissue, or if this represents a mechanism by which progesterone and PR in breast cancers overtake signaling pathways normally used by growth factors or other cell surface receptors (118). In contrast, PR-A does not efficiently mediate rapid activation of the protein kinase signaling pathway, making it ineffective at extranuclear signaling (120–122). In human breast cancer cell lines, PR-B was found to regulate gene expression of more genes than PR-A, with modest overlap in regulated genes by both receptors (123), further supporting the important role of PR-B in breast carcinogenesis. In breast cancer cell lines, PR-A is needed for appropriate responsiveness to progesterone, whereas PR-B-induced target gene expression has been shown to enhance the carcinogenicity of tamoxifen (124). Further, it has been suggested that antiprogesterins may be more effective for tumors expressing PR-A (125).

Recent studies suggest that the PR's role in breast cancer is more than just as a marker of ER activity (126, 127). Specifically, studies suggest PR acts as a binding partner as well as a modifier of ER activity that targets gene selection; however, findings on this topic have been inconsistent. Mohammed and colleagues (126) suggest that studies evaluating progesterone in addition to an antiestrogen may be informative, while Singhal and colleagues (127) supported the use of a selective PR modulator/antagonist to co-target both ER and PR in receptor positive breast cancers as a means for enhancing treatment, as compared with individual therapies. Even more complex interactions were suggested in

a study from the Lange laboratory that supported not only targeting ER/PR in hormone receptor positive cancers, but also the insulin-like growth factor 1 receptor (IGF1R) (128). As such, progesterone and PR are not the only receptors/ligand combination that likely play a role in breast cancer. Steroid receptor crosstalk, including an overview of ER/PR crosstalk in the context of primarily breast cancer, was recently reviewed by Truong and Lange (129).

DNA Repair Enzyme Mutations and Hormonally-Related Breast Cancers

Pathogenic mutations in DNA repair genes such as *BRCA1*, *BRCA2*, *CHEK2*, *PALB2*, *ATM*, *P53* and others are associated with elevated risk for breast cancers. While some gene mutations are linked to triple negative breast cancers (eg, *BRCA1*) others are linked to ER+ breast cancer (eg, *CHEK2*). The developmental and mechanistic origin of these tumors are poorly described in the literature. A recent study that investigated *BRCA1* mutation carriers concluded that *BRCA1* mutation could drive aberrant luminal progenitor expansion independent of progesterone (130). A more recent study from the same group has shown that *BRCA1* mutation carriers have deregulated progesterone signaling leading to higher proliferation and DNA damage in a progesterone-sensitive RANK+ luminal progenitor subset (111). An independent study of *BRCA1* mutation carriers observed that luminal progenitors displayed an aberrant differentiation program but failed to find expanded luminal progenitors in their samples (131). Furthermore, data from a mouse model provided evidence to suggest that aberrant expansion of *BRCA1* null luminal progenitor is linked to the replication-stress associated DNA damage response, where proliferation of mammary progenitors is perpetuated by damage-induced, autologous NF- κ B signaling (112). There are other functions of *BRCA1* relevant to genomic stability that are compromised in patients carrying a deleterious mutant allele of this gene. Recent studies show that haploinsufficiency of *BRCA1* leads to accumulation of R-loop, a DNA-RNA hybrid structure associated with transcriptional regulation and genomic instability in luminal progenitors (104). Chromosomal aberrations have been associated with mammary progenitors in *BRCA1* insufficiency as well as experimental models of human mammary progenitors mimicking loss of *BRCA1* function (132, 133). Together, human and murine studies provide credence to the notion that the origin of *BRCA1*-mutation associated triple negative breast cancers is in the ER- luminal progenitors. Such detailed examination of cancer risk mechanisms at the

level of cellular differentiation is simply not available for other DNA repair genes.

There is some evidence to support higher circulating levels of luteal phase estrogen and progesterone in *BRCA* carriers compared with noncarriers (134). Specifically, it has been suggested that the increased hormone levels observed in carriers are due to a defect in ovarian hormone biosynthesis that leads to an increased risk of breast and ovarian cancer (134). However, the link between premenopausal luteal phase circulating hormones and ovarian cancer risk is unclear. Additional research is required to determine whether hormones influence the potential mutagenic effects of *BRCA* mutations (or possibly other DNA repair enzyme mutations) and thus provide data to support organ specific penetrance.

Progesterone and Breast Carcinogenesis

Progestins Versus Progesterone

Breast cancers arise from the epithelial cells of the breast. It is now well accepted that pharmacologic and physiologic concentrations of estradiol increase the mitogenic activity of epithelial cells, while the influence of progesterone continues to be debated. Inconsistent evidence suggests that progesterone can increase, decrease, or have no effect on mitotic activity and proliferation in breast epithelial cells (135–138). In contrast, a relative consensus has been reached that long-duration exposure to pharmacologic progestogen levels combined with estrogen, either through use of contraceptives (transient, albeit small elevation in risk with current use) or menopausal hormone therapy (primarily with estrogen+progestin formulations, but not estrogen+progesterone or dydrogesterone), increases breast cancer risk (139–143). In addition to possible progestational effects, the progestins used in menopausal hormone therapy and oral contraception can have antiandrogenic, proandrogenic, glucocorticoid, and antiminerlocorticoid effects (14, 16) (summarized in Table 1). Thus, the potential effects of endogenous levels of progesterone in breast cancer cannot be inferred from studies of exogenous hormones, and epidemiologic studies evaluating circulating levels of progesterone are limited.

Progesterone and Proliferation

In vitro studies suggest that progesterone decreases mitotic activity of normal human breast epithelium and partially inhibits estrogen-induced proliferation in human cancer cell lines (135, 144). These data are limited and do not appear to be consistent with the *in vivo* effects of progesterone

on mammary cells (145). Recent discoveries suggest that functionally distinct epithelial and non-epithelial mammary cells are involved in mediating progesterone/RANKL-induced paracrine signaling to promote expansion and differentiation of mammary stem and progenitor cells, and these are not adequately captured in the *in vitro* studies. It may therefore be challenging to reproduce the mammary-specific effects of progesterone *in vitro*, especially in 2-dimensional cell culture systems. The 3-dimensional primary tissue-organoid based short-term culture system has been shown to capture some proliferative effects of progesterone on mammary tissue *ex vivo* but more studies are required (111).

Maximum breast epithelial proliferation measured in surgical biopsies has been reported for early follicular (136), early luteal (137), and late luteal phases (138, 146), with only the latter suggesting potential mitogenic activity for progesterone. A new model has been proposed to explain the delay in response. According to this model, mammary stem cell numbers oscillate during the menstrual cycle, in response to oscillations of serum progesterone and macrophage numbers in the breast (109). Further investigation can shed light on the link between progesterone and macrophage coordination to increase local Wnt concentration in order to increase mammary stem cell expansion.

Contrarily, it has also been suggested that increased progesterone tissue concentrations are correlated with decreased mitotic activity in normal breast epithelium in women (145). Specifically, treatment with hydro-alcoholic gel provided very limited changes in plasma concentrations of estradiol and progesterone but produced markedly different levels of steroid accumulation in the breast tissue that was consistent with the treatment (eg, higher concentrations of progesterone in the progesterone and estradiol+progesterone groups, and higher concentrations of estradiol in the estradiol and estradiol+progesterone treated groups). Mitotic activity was highest in the estradiol-only groups, and lower in the progesterone-only and estradiol+progesterone group, providing evidence that adding 10–13 days of progesterone to estradiol reduced the proliferative effect of estradiol alone (145). However, it is unclear whether the change in progesterone concentrations reflect pharmacologic or physiologic levels during a relevant period of the menstrual cycle (147). One study showed that breast cell proliferation is higher in the luteal than follicular phase (characterized by higher serum progesterone relative to estradiol levels) (137). At physiologic levels, it is unclear whether

this enhanced proliferation is due to progesterone exerting a mitotic effect or a weak antimitotic effect that is insufficient to block the mitotic effect of increased estradiol (147). As such, the association between progesterone and malignant transformation of breast epithelial cells may be dependent on the levels of estradiol, or other sex steroid hormones.

Hormones exert important effects on proliferation of normal breast epithelium, and therefore, progesterone is a plausible inducer of proliferation in the early phases of breast cancer development. In one model (148), risk of breast cancer was postulated to be determined by cumulative exposure of breast tissue to estrogen, but this same hypothesis could be relevant for the combined effects of estrogen and progesterone exposure over the menstrual cycle, and thus over a woman's reproductive years. Indirect evidence supporting this model included increased risk of breast cancer with events that increase estrogen exposure, but also plausibly increase progesterone exposure (possibly relative to estrogen) over the menstrual life course, including early age at menarche, shorter menstrual cycle length, late first full-term pregnancy, and late menopause. Inverse associations have also been demonstrated for factors related to potentially reduced exposure to estrogen and/or progesterone including oophorectomy and early menopause. Cumulative effects of these exposures seem to be associated with more accurate risk assessment than individual factors. Improved accuracy could also be due to more precise characterization of total exposure to progesterone that reflects exposure of TDLU epithelium that has not undergone differentiation induced by pregnancy and lactation, with resulting post-lactational involution (and "reset" or turnover of damaged cells). The model described above by Pike and colleagues (148) treated breast cancer as a composite outcome. Given advances in the understanding of etiologic heterogeneity in age-specific incidence and risk factor associations for breast cancer subtypes, it is realistic that this model should be refined by major subtype and/or hormone receptor status (149). Further, the cell of origin of the breast cancer subtype (eg, basal or luminal) may be differentially sensitive to progesterone. For example, if the cell of origin of basal breast cancer is more stem-like and sensitive to progesterone, this could explain the epidemiologic risk factor of high parity (higher lifetime progesterone exposure) for basal-like tumors. In contrast, the cell of origin for luminal cancers is likely less sensitive to progesterone, and epidemiologic risk factors like parity are associated with reduced risk.

Factors Contributing to Variations in Progesterone

Individual variations in progesterone exposure may also be attributed to other factors, and specifically this could help explain the paradoxical association of obesity (ie, high body mass index (BMI)) with pre- and postmenopausal breast cancer risk. Obese BMI is inversely associated with premenopausal breast cancer risk overall, and positively associated with postmenopausal breast cancer risk (150). However, the association among premenopausal women is heterogeneous by hormone receptor status (151, 152). Premenopausal obesity is associated with increased risk of ER-, PR-, and triple negative breast cancers (ER-/PR-/HER2-) and decreased risk of hormone receptor positive tumors (151, 152). However, the increased breast cancer risk with obesity in postmenopausal women was not heterogeneous across hormone receptor status (151, 152). The association in postmenopausal women is likely a result of increased estradiol due to higher aromatase activity related to increased adipose tissue in overweight/obese women. However, the reduced risk of hormone receptor positive tumors in premenopausal women is more likely explained, in part, by reduced progesterone exposure due to dampening of progesterone peaks in premenopausal women with high BMI that can result in longer cycle length and increased frequency of anovulatory cycles, thereby reducing breast epithelial cell proliferation (153). Alternatively, or complementary, to the contribution of increased circulating estrogen concentrations with increased adiposity, insulin resistance could also be a mechanism by which obesity is associated with increased breast cancer risk. Insulin is a modest growth factor that promotes cell growth, division, and migration, and inhibits apoptosis. While insulin is a weaker mitogen than other prominent growth factors (eg, insulin-like growth factors, platelet-derived growth factor, vascular endothelial growth factors, etc.) its specific mitogenic action potentiates cellular responsiveness to other growth factors, in a sense amplifying its mitogenic capabilities (reviewed in (154)).

Plausible Mechanisms

In terms of potential mechanisms for progesterone's ability to cause breast cancer, we have emphasized data throughout the review supporting that both estrogen and progesterone are required to elicit substantial cellular proliferation. Research supporting genotoxic effects of progesterone is very limited. Studies evaluating blood DNA damage via the comet assay, dominant lethal

assays in mice, chromosomal aberrations in rats, and DNA double strand breaks in mice and rats were all null (155–157). Individual studies have demonstrated that treatment with progesterone induces an increase in chromosomal aberrations in embryonic human fibroblasts or aneuploidy by a non-disjunctional mechanism (158, 159). No direct DNA damage (via measurement of DNA adducts) was demonstrated in the liver after treatment with progesterone (160, 161). Global gene expression in human breast tissues (non-cancer) donated for research from 20 premenopausal women revealed that 255 genes were differentially expressed in a study comparing tissues collected during the follicular and luteal phase (162). Of the 255 genes, 221 were increased in the luteal phase with functions related to cell cycle, mitosis, and DNA damage and repair; 3 paracrine factors were also identified, including RANKL, WNT4, and epiregulin (162). Currently, there is no direct evidence to suggest that progesterone has the capability to initiate tumors via inducing enzymes and proteins involved in nucleic acid synthesis or through activation of oncogenes. The expansion of a cancer-susceptible luminal progenitor population is the proposed indirect developmental effect of progesterone that may contribute to tumor initiation. The discovery of widespread telomere dysfunction in ostensibly normal luminal progenitors in the cancer-free breast (102) and widespread telomere fusion observed in early breast cancer lesions (163) have provided strong support for the specific cell-of-origin theory for breast cancers.

Other Hormones and Breast Carcinogenesis

Elevated circulating androgen levels are consistently associated with increased breast cancer risk, but the underlying mechanism of action is unclear (164). Androgens may increase risk indirectly through aromatization to estrogens, whose role in breast cancer etiology is well established (165), or decrease risk by exerting antiestrogenic and antiproliferative effects via androgen receptor (AR) signaling (164, 166). Androgenic activity via the AR is responsible for healthy functioning of many organs in women (166) including the breast, where estrogens stimulate while androgens inhibit development, and the balance between these regulates breast development. In breast tumors, AR expression has been detected in up to 85% of cases (164), although this varies by breast cancer subtype; while almost all ER+ cancers express AR, only 10–35% of triple negative (ER-/PR-/HER2-) breast cancers express AR (167).

Elevated levels of circulating postmenopausal androgens (androstenedione and testosterone) and androgen precursors (dehydroepiandrosterone (DHEA) and its sulfated form (DHEAS)) increase breast cancer risk, with women in the top quartile of these hormones at 2–3 fold increased risk versus women in the lowest quartile (165, 168). Because these androgens are the obligate precursors for estrogens, it is not known whether their observed effect is independent of the estrogen association. In addition, epidemiologic studies have not examined the functional role of androgen metabolites synthesized within the breast. Indeed, the normal breast contains the steroidogenic enzymes needed to convert circulating precursors to biologically active forms (169), with the derived 5 α -reduced androgens, particularly dihydrotestosterone (DHT), being the most potent. In women, however, levels of circulating DHT are often below assay detection, and do not reflect peripheral 5 α -reductase activity (170). Rather, androsterone glucuronide (ADT-G), a distal metabolite of DHT, together with androstanediol glucuronide (found as 2 isomers: 5 α -androsterone-3 α ,17 β diol-3-glucuronide (3 α -diol-3G) and 5 α -androsterone-3 α ,17 β diol-17-glucuronide (3 α -diol-17G)), have been shown to reflect total tissue-level androgenic activity better than the proandrogens (eg, testosterone, androstenedione, etc.) (170, 171). Additional research is necessary to understand the interplay between progesterone, estrogen, and androgens in breast cancer risk with careful attention to potential interactive effects of the hormones and possible differences in associations by breast cancer subtypes.

Unlike female breast cancer, breast cancer in males is exceptionally rare (less than 1% of female breast cancer risk) and does not plateau after age 50. The slowed increase in incidence in women at the menopausal transition likely reflects in part the substantial decrease in circulating levels of a number of hormones including estrogen and progesterone. Further support for the importance of estrogen in both male and female breast cancer etiology is lent by the association with anthropometry, reproductive factors, and circulating hormones (172, 173). Obesity increases postmenopausal breast cancer risk and male breast cancer risk by similar magnitudes, likely as a result of higher levels of bioavailable estrogen via peripheral conversion of androgens to estrogens (172). Increased risk for male breast cancer with higher circulating levels of estradiol and a null association with ADT-G (173) suggests that the substantially lower rate of these cancers among men is likely due

in part to much lower lifetime estrogen exposure than among women. Progesterone levels across the life course are also substantially lower in men than women, but they have not been evaluated in the context of male breast cancer risk to date.

Prolactin is another hormone (described above) that interacts with progesterone and their respective receptors to influence ductal and luminal epithelial cell proliferation during pregnancy (reviewed in (174)). High circulating prolactin is related to breast cancer risk factors, such as nulliparity and high mammographic density. Further, higher levels of circulating prolactin are associated with increased risk of predominantly receptor positive pre- and postmenopausal breast cancer risk (175). It has been suggested that both progesterone and prolactin may influence the epithelial cell hierarchy via RANKL feedback through mammary stem cells and luminal progenitors; however, more research is needed to understand the shared signaling pathways and whether they could serve as potential therapeutic targets (174).

Breast Cancer Risks Associated With Endogenous and Exogenous Progesterone/Progesterin Exposure

Endogenous Progesterone Exposure

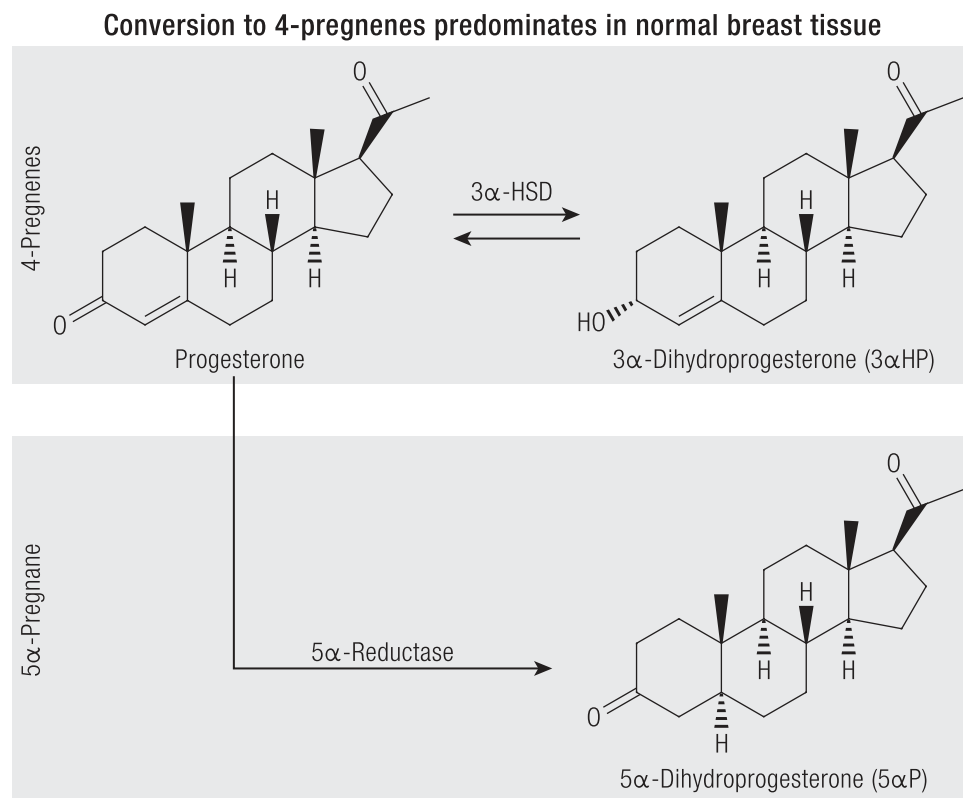
Few population-based studies have assessed circulating progesterone levels and breast cancer risk. One study of postmenopausal women including 322 breast cancer cases and 643 matched controls (176) showed no association between prediagnostic circulating progesterone levels and breast cancer risk. However, the study was limited because almost 30% of samples had undetectable levels using an RIA assay, with preceding organic solvent and Celite column chromatography steps, and a limit of detection of 3 ng/dL. Thus, the influence of endogenous progesterone on breast cancer risk among postmenopausal women has been infrequently studied primarily due to low levels of circulating progesterone, the large sample volume required for assays, and inadequate sensitivity of available assays.

Studies of circulating progesterone in premenopausal women and subsequent breast cancer risk include 6 published studies, with the majority measuring progesterone in luteal phase blood samples (177–182). Of these studies, 4 included a limited number of cases ($n \leq 104$) and generally reported null results (177, 178, 182) or a modest reduction in breast cancer risk with higher levels of circulating progesterone (179). The 2 largest studies evaluating progesterone concentrations in premenopausal women utilized either direct

chemiluminescent immunoassays on automated platforms (180) or direct RIA using a commercial kit (181). Among 501 breast cancer cases and 1030 matched controls from the Nurses' Health Study II, luteal phase progesterone was not associated with breast cancer risk (180). There was no difference in the association by menopausal status at cancer diagnosis (180). No association with progesterone among 801 premenopausal breast cancer cases and 1132 matched controls, irrespective of timing of the blood draw in the menstrual cycle, was seen in the European Prospective Investigation into Cancer and Nutrition (EPIC) (181). Analyses limited to progesterone levels from 232 cases and 323 controls measured during the luteal phase of the menstrual cycle also suggested no association (181).

Circulating progesterone levels are significantly lower among postmenopausal women as compared to premenopausal women, posing a challenge for immunoassay techniques (117). Furthermore, available commercial kits measure only progesterone and do not measure relevant major metabolites which may also play a role in the carcinogenic process. A number of progesterone metabolites have been characterized in normal breast and cancerous breast cancer tissue and broadly fall into 2 groups, 1) 4-pregnenes: metabolites that retain their double bond, and 2) 5 α -pregnanes: metabolites in which 5 α -reductase has reduced the double bond (3). Different relative distributions in the 4-pregnenes and 5 α -pregnanes have been demonstrated in normal and cancerous breast tissue (Fig. 4). In normal breast tissue the conversion of progesterone to pregnene metabolites exceeds or predominates the conversion of progesterone to 5 α -pregnane metabolites, whereas the opposite is true in breast cancer tissue. Thus, it has been hypothesized, and laboratory studies have confirmed, that the promotion of breast cancer may be related to relative changes in concentrations of cancer-inhibiting 4-pregnenes (eg, 3 α -dihydroprogesterone (3 α HP)) and cancer-promoting 5 α -pregnanes (eg, 5 α -dihydroprogesterone (5 α P)) (183). Further, 5 α P and 3 α HP have been shown to act with equal efficacy on all breast cell lines tested, regardless of their receptor status, estrogen sensitivity, and tumorigenicity (184). While the ovary is the primary source of progesterone in premenopausal women, smaller amounts are produced in the adrenals and adipose tissue, and local production of 3 α HP and 5 α P metabolites may occur in the breast (183). However, to date, no epidemiologic studies have evaluated the role of circulating progesterone metabolites and breast cancer risk in premenopausal or postmenopausal women.

Figure 4. Primary pathways of progesterone metabolism in breast tissue. Legend: In normal tissue, pregnenes (progesterone is a pregnene) are the predominant compounds. All of the 4 pregnenes (not shown) can be irreversibly converted to 5 α -pregnane (respectively) via 5 α -reductase. The 2 metabolites, 3 α HP and 5 α P, show the greatest differences between tumorous and nontumorous tissue; the ratio of 5 α P-to-3 α HP is more than 10-fold higher comparing breast tumors to normal breast tissue. Metabolic activities of the hormones are similar in pre- and postmenopausal women, by age, and by ER-status.



Exogenous Progesterone and Progestin Exposure

Information about the risk of breast cancer associated with isolated progesterone or progestin use is scant. Although oral and parenteral progestin-only formulations are used for contraception by premenopausal women, clinical breast cancer data on the progestin-only effect is limited because the reproductive age group of contraceptive users has a low baseline risk of breast cancer.

Oral contraceptives are widely used but the majority contain an estrogen combined with a progestin (combined oral contraceptives (COC)). Progestin-only pills (POP) are also available. Their primary use is for contraception in women in which estrogen is contraindicated. COCs act by inhibiting ovulation, whereas POPs primarily prevent conception by developing thick, hostile cervical mucus and atrophy of the endometrium. In postmenopausal women, progestogens are widely used in combination with an estrogen to protect the uterus. The result of adding a progestogen to

estrogen in COCs or menopausal hormone therapy creates a complex and confusing situation because there is no consensus on the background effect of the estrogen itself and the effect of its dose, delivery system, and treatment time. Additionally, in premenopausal women the addition of a progestogen may prevent ovulation, thus both the direct effect of the progestogen on the breast and indirect effects associated with anovulation are inseparable.

Sparse data on progestin-only contraceptive usage and breast cancer risk

To date, a limited number of studies have been published on POPs and breast cancer risk (reviewed in (185–187)). The most recent, and largest, included a prospective population-based cohort in Denmark with contraceptive information on 1.8 million premenopausal women, of which 11,517 developed breast cancer during an average of 11 years of follow-up (187). Current or recent use of the levonorgestrel POP was associated with

increased breast cancer risk compared to never use of hormonal contraception [levonorgestrel POP RR (95% CI): 1.93 (1.18–3.16)]. However, the number of women exposed to levonorgestrel POP in this analysis was relatively small. Other formulations of POPs (norethindrone or desogestrel) with larger proportions of users were not associated with breast cancer risk (187). In this same study, the levonorgestrel-releasing intrauterine system was also associated with increased breast cancer risk: 1.21 (1.11–1.33), whereas progestin-only implants and injections (DMPA) were not associated with risk (187). A recent review of the literature relating contraceptives to breast cancer risk estimated that progestin exposure at 6 months of levonorgestrel-releasing intrauterine system use would be roughly half of the progestin exposure with estrogen+progestin menopausal hormone therapy (ie, 2.5 mg of MPA), which is similarly associated with increased breast cancer risk and is thus a plausible association (188). They also observed that the current levonorgestrel-releasing intrauterine system use in the Danish cohort study did not take into account prior contraceptive use with its associated known increases in risk, which could have contributed to the increased breast cancer incidence (188). Further the Danish cohort study had too little exposure to DMPA, contraceptive implants, and most oral POPs to report precise risk estimates for those progestin-only methods, which are more widely used in the United States (188). A comparable increased breast cancer risk [Standardized Incidence Ratio (95% CI): 1.19 (1.13–1.25)] with the levonorgestrel-releasing intrauterine system was observed in a cancer registry linkage analysis using data from the prescription National Reimbursement Registry among 93,843 Finnish women; and in an updated analyses reported that the increased risk was apparent for both ductal [SIR (95% CI): 1.20 (1.14–1.25)] and lobular breast cancer [SIR (95% CI): 1.33 (1.20–1.46)] (189, 190). For the latter study, the same limitation of not being able to account for prior COC use is relevant since the prescription reimbursement registry does not capture lifetime exposure history, and thus the associations with current levonorgestrel-releasing intrauterine system use could be confounded by prior COC use.

In a large, US population-based case-control study in which POP use accounted for 0.5 percent of total oral contraceptive use, POP use was not associated with breast cancer [OR 0.9 (95% CI not reported)] (191). Progestin-only injections or implants were also not associated with breast

cancer risk in the same study population (192). The Norwegian and Swedish Women's Lifestyle and Health cohort study included 1008 primary breast cancers diagnosed among 103,027 women. Current use of POPs (which could have included former use of COCs) was associated with increased breast cancer risk [RR (95% CI): 1.6 (1.0–2.4)], while exclusive use of POPs (regardless of duration of use) was not associated with breast cancer risk [RR (95% CI): 1.1 (0.8–1.7) 29 cases out of 3435 vs. never, 261 cases out of 28,171] (193). In a study from the large French prospective cohort study (Etude Epidémiologique de Femmes de la Mutual Générale de l'Education Nationale, referred to as the E3N study or cohort), ever use of POPs after the age of 40 was not associated with breast cancer risk. Some elevations were reported in subgroup analyses, suggesting that prolonged use of oral progestin (more than 4.5 years of continuous use) was associated with increased breast cancer risk among older women (194). Earlier studies, specifically 8 case-control studies conducted between 1986 and 1996, reviewed in an IARC monograph, reported no risk of breast cancer among women using progestin-only contraceptives (pill or injectable DMPA) compared to non-users (186).

Combined oral contraceptives and breast cancer risk

COCs have been associated with a small magnitude increase in breast cancer risk. The Collaborative Group on Hormonal Factors in Breast Cancer was set up in 1992 to bring together and reanalyze data from 54 studies in 25 countries to assess the relationship between the risk of breast cancer and use of COCs (195). The study involved data from a total of 53,297 women with breast cancer and 100,239 women without breast cancer. The results provide strong evidence for the following conclusions. First, there is a small increase in the relative risk (RR) of having breast cancer diagnosed in current users (RR: 1.24) and in the 10 years following termination of COC use (RR: 1.16 after 1–4 years and RR: 1.07 after 5–9 years). Second, there is no significant excess risk of having breast cancer diagnosed 10 or more years after stopping use. In addition, the data show that the cancers diagnosed in women who had used COCs were less advanced clinically than in never-users. Increased risk among women currently taking COCs and diminishing risks after cessation of use, suggest a promotional, rather than an initiating, effect of progestin in these medications on breast cancer risk.

Menopausal hormone therapy and breast cancer risk

The relationship between risk of breast cancer and use of menopausal hormone therapy has been investigated in many epidemiologic studies. A 1997 review by the Collaborative Group on Hormonal Factors in Breast Cancer brought together and reanalyzed data from 51 epidemiologic studies that represented about 90% of the worldwide evidence on this topic (140). The main analyses of the relation between risk of breast cancer and use of hormone therapy included 53,865 postmenopausal women (17,949 cases and 35,916 controls) with known age at menopause and use of menopausal hormone therapies. The results show that the increase in the relative risk of breast cancer associated with each year of use in current and recent users is small. Therefore, inevitably some studies show significant associations and others do not. However, combination of the results across many studies has the advantage of reducing such fluctuations. The increased risk of breast cancer was not significant until 5 years of menopausal hormone therapy use. For women who used menopausal hormones for 5 years or longer, the relative risk was 1.35 (95% CI: 1.21–1.49). The risk was reduced after discontinuing menopausal hormone therapy use and largely, if not completely, disappeared after about 5 years. Information about the hormonal constituents of the menopausal hormone therapy preparations used the most was available for only 4,640 (39%) of eligible women. The women in the main analysis had their breast cancers diagnosed on average in 1985, when the type of hormone used was predominantly estrogen alone; only 12% of the women used estrogen and progestogen combinations. There was no significant variation in the relative risk of breast cancer according to the formulation or dose of the estrogen used by most women, and no evidence of marked differences between menopausal hormone preparations containing estrogen alone and those containing both estrogen and progestogen.

In a later review (196), data from 8 randomized controlled trials and 19 epidemiologic studies were used to determine whether there is an association between menopausal hormone therapy use and breast cancer. The review also addressed a number of other questions, including whether data exist to show a differential effect from use of menopausal hormone therapy products, doses, regimens, and routes of administration. The results show that the average risk of breast cancer with use of estrogen alone was 0.79 (95% CI: 0.61–1.02) in 4 randomized

trials involving 12,643 women, whereas the risk with estrogen-progestin use was 1.24 (95% CI: 1.03–1.50) in 4 randomized trials involving 19,756 women. In contrast, the average relative risks reported in the epidemiologic studies were higher in current users: 1.18 (95% CI: 1.01–1.38) for estrogen alone and 1.70 (95% CI: 1.36–2.17) for estrogen-progestin; however, the risks were not as strong in ever users: 1.08 (95% CI: 0.97–1.20) and 1.31 (95% CI: 1.12–1.53), respectively (196).

For the analyses of the association of breast cancer risk by hormone formulation, dose, regimen and route of administration, the number of studies used in each group varies. For progestogen types, only 2 studies included risks with current exposure (139, 197). The analyses compared a C-21 related progestogen, namely MPA, with 2 C-19 related progestogens, norethindrone and levonorgestrel. Breast cancer risks with C-21 and C-19 related progestogens were virtually the same (2.14, 95% CI: 1.18–3.87 and 2.14, 95% CI: 1.68–2.72, respectively) (196). Only one study reported information on progestogen dose; the Women's Health Study evaluated MPA dosages over the ranges of <5 to 10 mg/day; the trend from lowest to highest dose was not significant (198).

Although the 2 major reviews of data reanalysis just described include a substantial number of studies (51 epidemiologic studies in one review, and 8 randomized controlled studies and 19 epidemiologic studies in the other review), none of these studies include progesterone data. However, this changed with the publication of results from the E3N study to investigate risk factors for cancer in women, including exogenous hormones (142, 199). Although oral progestins were used most widely in the E3N study, there was also substantial use of exogenous progesterone. The progestins included dydrogesterone, medrogestone, chlormadinone acetate, cyproterone acetate, promegestone, nomegestrol acetate, norethindrone acetate, and MPA.

In an updated report on the E3N cohort evaluating 80,377 postmenopausal women 40–65 years of age at enrollment and followed for up to 12 years, use of estrogen alone was associated with a 29% increased breast cancer risk (95% CI: 1.02–1.65) (199). This increased breast cancer risk with use of estrogen alone (almost exclusively estradiol compounds and mostly administered transdermally) differs from that of the WHI estrogen-alone trial which found a decreased risk with oral conjugated equine estrogens (200).

The association of estrogen-progestogen combinations with breast cancer risk varied

significantly according to the type of progestogen (199). The RR was 1.00 (95% CI: 0.83–1.22) for estrogen-progesterone, 1.16 (95% CI: 0.94–1.43) for estrogen-dydrogesterone, and 1.69 (95% CI: 1.50–1.91) for estrogen combined with other progestogens, which included medrogestone, chlormadinone acetate, promegestone, norgestrel acetate, norethindrone acetate, and MPA; the increased breast cancer risk was consistent across these latter estrogen-progestogen formulations. Estrogen alone, estrogen-progesterone and estrogen-dydrogesterone did not differ significantly from one another in breast cancer risk, but the risks were all significantly lower than that of estrogen combined with the other progestogens. It should be noted that the chemical structure of dydrogesterone differs from that of progesterone only in that the methyl group at the carbon 10 is in the alpha-orientation and there is a double bond between carbons 6 and 7. The E₃N study was the first epidemiologic study to show that estrogen-progesterone and estrogen-dydrogesterone combinations may be the least harmful menopausal hormone therapies with respect to postmenopausal breast cancer risk. However, more evidence is required to make firm clinical recommendations for use of these formulations, compared to other formulations, in managing menopausal symptoms.

Challenges, Future Directions, Unanswered Questions

There are many gaps in our current knowledge of the cellular hierarchy model of mammary gland development and regulation of its cellular content and glandular function throughout life. Though it is assumed that the progesterone-sensor luminal cells constitute a developmentally terminal 'post-mitotic' state, the following observations raise doubts that cannot be ignored: 1) Lack of sufficient evidence in the literature that functionally and molecularly-defined ER+PR+ luminal cells are direct products of ER-PR- luminal progenitor cells; 2) In humans, the telomere length of the terminal luminal cell-fraction is longer than luminal progenitor cells in younger women and gradually declines with age (102); 3) A study looking at the synthetic nucleoside labeling cycling cells within defined mammary subsets in a mouse model has argued that luminal progenitor and luminal cells could represent 2 distinctly self-sustained lineages with shared luminal features (201); 4) It appears

that the post-mitotic ER+ luminal cells which generally co-express PR can be induced to grow and expand *in vitro* (202). If ER+PR+ are indeed self-sustaining luminal cell lineages, then our understanding of how progesterone controls mammary cells may have to be revisited. Hence, the current mammary stem cell model requires further scrutiny. There is much to be learned about PR and its ligands with respect to downstream signaling and genomic and cellular targets. Better antibodies are needed to detect PR isoforms and cell systems to capture paracrine signaling events. Advances in these areas will provide a broad spectrum of exciting new discoveries that will improve not only the biologic understanding of progesterone in breast cancer, but also the clinical relevance.

There are many unanswered questions pertaining to progestogens and breast cancer risk. Epidemiologic evaluation of the role of endogenous progesterone, including newly identified progesterone metabolites, in the etiology of premenopausal or postmenopausal breast cancer risk is limited. The latter is primarily due to limited assay sensitivity while the former is due to limited annotation of the menstrual phase of blood draw or urine collection in prospective epidemiologic studies that include breast cancer outcomes. Studies have either included samples collected from women across the menstrual cycle or limited to luteal phase samples. It is not clear if this is the correct approach to determining risk, whether the maximum level from the luteal cycle is the appropriate measure, or whether the usual level during the early follicular phase could be more relevant to risk. Further, it could be the change in progesterone levels from follicular to luteal, or the concentration relative to estradiol that is most relevant to risk. Finally, cumulative exposure over a lifetime or levels during the final years of the menopausal transition could be important. At present, it is not clear what the appropriate comparison should be, and typical sample collections are limited to only one time point during the menstrual cycle, which would preclude the ability to evaluate the change in progesterone level over a menstrual cycle. Given recent advances in assay technology (LC-MS/MS assays can detect circulating progesterone concentrations ~0.1 ng/dL (117)), clarification of an association between endogenous progesterone and any relevant metabolites with breast cancer risks among premenopausal and postmenopausal women should be resolved in the near term. These studies should consider whether associations differ by tumor characteristics, most importantly

molecular subtype and receptor status, as well as by race, BMI, and likely other genetic and non-genetic factors (eg, other circulating hormones). Studies are also needed to evaluate the association between progesterone/progesterone metabolites and breast cancer risk in higher risk groups, including women with ductal carcinoma in situ, a history of benign breast disease, or increased mammographic breast density, and specifically among *BRCA1* mutation carriers, as antiprogesterin therapy may be particularly relevant to this subpopulation.

Questions remain as to whether POPs or progestogen-only menopausal hormone therapy are related to breast cancer risks. In addition, breast cancer risks with low-dose and newer formulations of oral contraceptives and menopausal hormone therapy remain unresolved. Thus, continued monitoring of breast cancer associations with COCs, new generation progestin-containing contraceptives, and progestogen-containing menopausal hormone therapy use is needed. Further, it is unclear if progesterone and progestins relate to breast cancer risk through direct or indirect proliferative effects, as well as the potential for genotoxicity, DNA damage, tumor-initiating effects via induction of enzymes and proteins involved in nucleic acid synthesis, or through activation of oncogenes.

In this review we did not extensively summarize the expansive literature about the possible role of progesterone or PR in breast cancer prognosis and/or treatment. In section IV.F, we reference recent support for a more active role of PR in hormone receptor positive breast cancers. These

studies individually made additional therapeutic recommendations beyond treatment with tamoxifen alone. As a result, multiple clinical trials have been developed to test combination hormone therapies targeting both PR and ER in breast cancer treatment. It is clear in this context that the clinical trials are outpacing the basic science in that there is more to learn regarding steroid receptor cross-talk in breast cancer (129), yet the clinical trials are actively testing combination therapies that include PR targets. At a minimum we will learn clinically relevant information from these trials. If the emerging laboratory and epidemiologic data support a role of progesterone in the etiology of breast cancer, it is plausible that similar trials (focused on prevention) could be planned to evaluate the use of combination therapies in women at high risk of breast cancer who would be given tamoxifen as current standard clinical practices.

Although many unanswered questions about the role of progesterone in breast physiology and carcinogenesis remain, there have been multiple advances over recent years. It is likely that continued progress in our understanding of the potential proliferative role of progesterone in the adult breast, including advances in stem cell technology, may clarify the role of progesterone in the initiation and progression of breast cancers. Finally, development of sensitive LC-MS/MS assay technology will likely facilitate evaluation of progesterone and progesterone metabolites in large epidemiologic studies of breast cancer that will ultimately address some of the data gaps outlined in this review.

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