

Circulating Estrogens and Postmenopausal Ovarian Cancer Risk in the Women's Health Initiative Observational Study

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Abstract

Background: Hormonal and reproductive factors contribute to the development of ovarian cancer, but few studies have examined associations between circulating estrogens and estrogen metabolites and ovarian cancer risk. We evaluated whether serum estrogens and estrogen metabolite levels are associated with ovarian cancer risk among postmenopausal women in a nested case-control study in the Women's Health Initiative (WHI) Observational Study (OS).

Methods: We selected all 169 eligible epithelial ovarian cancer cases and 412 matched controls from women enrolled in WHI-OS who were not using menopausal hormones at baseline. Baseline levels of 15 estrogens and estrogen metabolites were measured via liquid chromatography/tandem mass spectrometry. Associations with ovarian cancer risk overall and stratified by histologic subtype (serous/nonserous) were analyzed using logistic regression. The mean time from serum collection to cancer diagnosis was 6.9 years.

Results: Overall, we observed modest ovarian cancer risk associations among women with higher levels of estrone [OR (95% confidence interval) quintile (Q)5 vs. Q1: 1.54 (0.82–2.90), $P_{\text{trend}} = 0.05$], as well as 2- and 4-methoxyestron metabolites [2.03 (1.06–3.88), $P_{\text{trend}} = 0.02$; 1.86 (0.98–3.56), $P_{\text{trend}} = 0.01$, respectively]. Associations of estrogens and estrogen metabolites varied substantially by histologic subtype. Associations with serous tumors were universally null, while estrone [2.65 (1.09–6.45), $P_{\text{trend}} = 0.01$, $P_{\text{heterogeneity}} = 0.04$], unconjugated estradiol [2.72 (1.04–7.14), $P_{\text{trend}} = 0.03$, $P_{\text{heterogeneity}} = 0.02$] and many of the 2-, 4-, and 16-pathway metabolites were positively associated with nonserous tumors.

Conclusions: Our study provides novel molecular data showing an association of the parent estrogens and several estrogen metabolites with nonserous ovarian cancers.

Impact: These findings further support the heterogeneous etiology of ovarian cancer. *Cancer Epidemiol Biomarkers Prev*; 25(4): 648–56. ©2016 AACR.

Introduction

Elevated estrogens may be involved in ovarian carcinogenesis; however, the exact nature of the estrogen-ovarian cancer link remains unclear (1). Studies indicated that ovarian cancer risk is higher in women with surrogates of high cumulative estrogen

exposure including early age at menarche, late age at natural menopause, and long-duration exposures to menopausal hormone therapy (2–4). Following the marked reduction in menopausal hormone therapy use around 2002 there was an accelerated decrease in ovarian cancer incidence rates, further supporting a role for estrogens in postmenopausal ovarian cancer risk (5). Conversely, ovarian cancer risk is reduced in women with a history of oral contraceptive use, possibly by reducing the number of lifetime ovulations and by decreasing intra-ovarian estrogen levels (6).

In the premenopausal ovary, estrogen (namely estradiol) is produced via aromatization of androgens within the granulosa cells, and plays a significant role in follicular development (7). However in postmenopausal women, estrone is the predominant estrogen and is produced from peripheral conversion of adrenal androgens (7). Estrogen metabolites are formed from the conversion of estrone and estradiol via hydroxylation at the 2, 4, or 16 position of the carbon ring (Fig. 1; ref. 8). Estrogen effects, whether proliferative or anti-proliferative, are modulated by metabolic conversions occurring systemically and in the target tissues (8). Most experimental research demonstrating differential effects of estrogen metabolites has been conducted in mammary tissue and breast cancer cell lines, with limited research on other hormonally mediated tumors or their target tissues (8). The few experimental studies on ovarian cancer have demonstrated that estradiol can

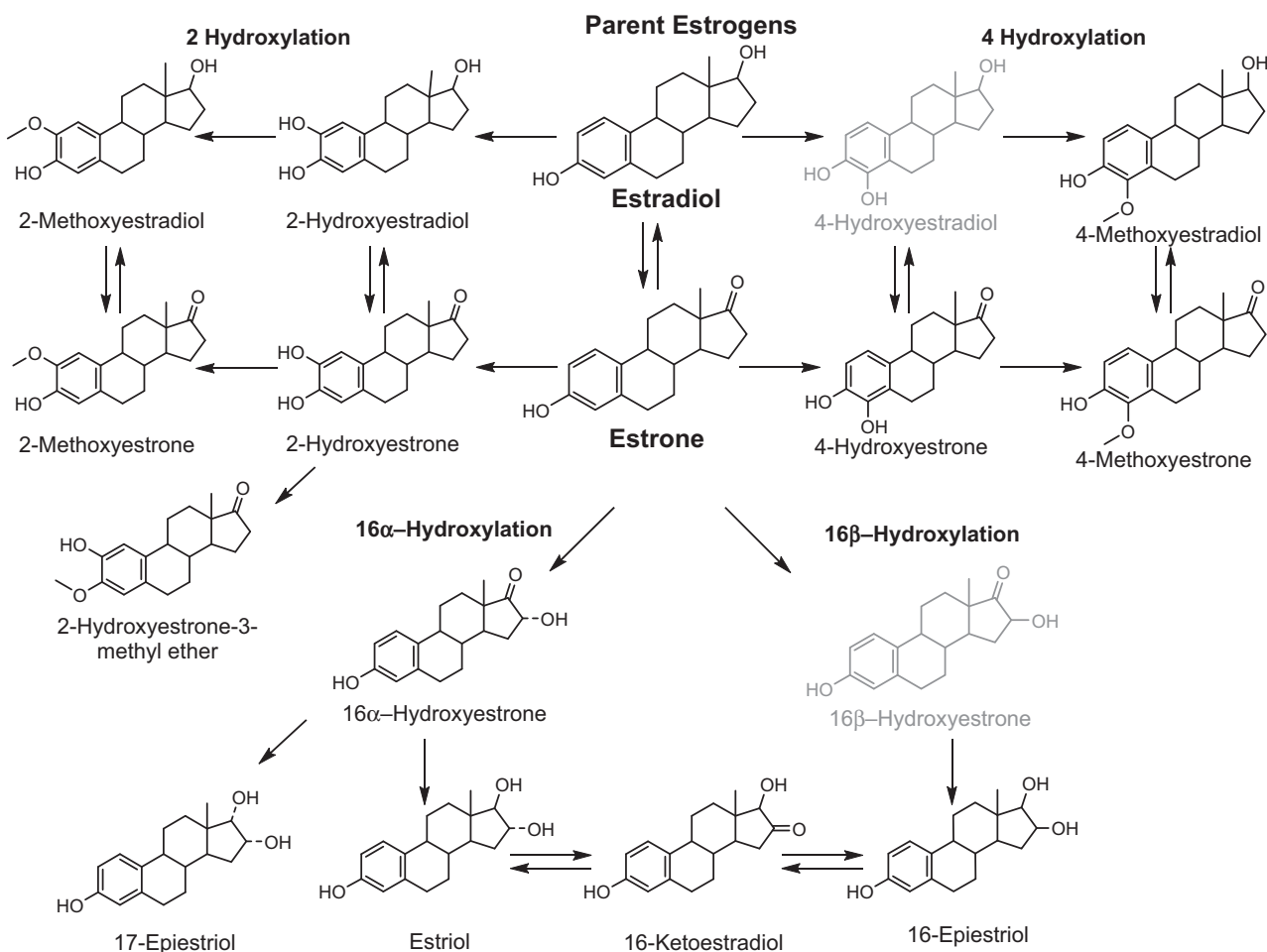
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**Figure 1.**

Formation of 2-, 4-, and 16-hydroxylation pathway estrogen metabolites from parent estrogens. The current serum estrogen metabolite assay measures 15 of the 17 metabolites pictured; 4-hydroxyestradiol and 16 β -hydroxyestrone (in light gray) are not measured with the current assay due to very low abundance in circulation.

stimulate ovarian epithelial cell growth or induce aberrant proliferation with increasing DNA mutations (9–11), but have not evaluated similar effects of estrogen metabolites. The presence of sex steroid hormone receptors in malignant epithelial ovarian tumors supports a potential role for hormones in the origin and promotion of ovarian cancer (12).

Studies evaluating circulating estrogens and ovarian cancer risk are limited. Two prior studies that evaluated associations with prediagnostic circulating estrone and/or estradiol did not find associations with ovarian cancer risk (13, 14), and the one prior study that examined estrogen metabolites was not able to evaluate associations by subtype ($n = 67$ ovarian cancer cases overall) and did not link prediagnostic estrogens to ovarian cancer (15). A study evaluating early pregnancy sex steroid hormones and subsequent cancer risk reported no associations with serous tumors, and an increased risk of endometrioid tumors with higher estradiol concentrations (16). Given the limited number of studies evaluating circulating estrogens and ovarian cancer risk, we conducted a nested case-control study within the Women's Health Initiative (WHI) Observational Study (OS), to evaluate the associations between 15 prediagnostic estrogens and estrogen meta-

bolites and ovarian cancer risk among postmenopausal women not currently using menopausal hormones. With increasing evidence of etiologic heterogeneity of epithelial ovarian cancers, especially with respect to hormonal risk factors (e.g., body mass index (BMI), menopausal hormone therapy use) (17–19), we also evaluated associations by tumor subtype.

Materials and Methods

Study population

The WHI is a large multicenter clinical investigation that enrolled women into either a clinical trial or an observational study to evaluate the predictors and causes of morbidity and mortality in postmenopause. The OS is a prospective cohort that enrolled 93,676 postmenopausal women ages 50 to 79 years at 40 centers through the United States between 1993 and 1998 (20, 21). Women were excluded ($n = 148$) if they had medical conditions with a predicted survival of less than 3 years; if they had adherence/retention issues; or if they were participating in a clinical trial. The present nested case-control study included incident ovarian cancer cases that were diagnosed between study

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initiation and May 2012. Both cases and controls met the following criteria to be eligible: no history of cancer at baseline other than non-melanoma skin cancer (n excluded = 11,674); no current use of exogenous hormones (38,419); no history of bilateral oophorectomy (7,470); and at least 1.1 mL of available pre-diagnostic serum (720). Baseline serum samples were provided for all participants.

Ovarian cancer cases were WHI-OS participants with hospital-confirmed diagnoses of incident primary epithelial ovarian, fallopian tube, or peritoneal cancer. Primary ovarian cancers ($n = 140$) were locally and centrally adjudicated, while fallopian tube ($n = 9$) and peritoneal cancers ($n = 20$) were locally adjudicated. Among the cases, the mean time from sample collection to diagnosis was 6.9 years (SD = 3.8 years; range = 352 days–14.8 years). Controls were eligible WHI-OS cohort members selected from strata defined by age at blood draw (50–54, 55–59, 60–64, 65–69, 70–74, 75–79), year at blood draw (1993–1996, 1997–1998), race/ethnicity (white, black, Hispanic, other/unknown), hysterectomy status at baseline or follow-up prior to event (yes/no), and time since last menopausal hormone use (≤ 1 year, > 1 year). Controls were drawn from the set of eligible cohort members in each stratum containing ovarian cancer cases that were alive at the time of diagnosis of their matched case and were selected with a ratio of at least 2 controls per 1 case per stratum. We included never ($n = 379$) and former menopausal hormone users ($n = 212$, of whom $n = 18$ reported using hormones within 1 year of blood draw) in this analysis and excluded women with unconjugated estradiol concentrations greater than or equal to 184 pmol/L (~ 50 pg/mL; $n = 10$) which is typically indicative of exogenous hormone use. The current study included 169 epithelial ovarian cancer cases and 412 matched controls. Among ovarian cancer cases, 102 were serous tumors and the remaining 67 were nonserous (13 endometrioid, 11 clear cell, 9 mucinous, and 34 other epithelial subtypes). Approval for conducting the study was obtained from human subjects review at the Fred Hutchinson Cancer Research Center (WHI Clinical Coordinating Center), as well as at all 40 clinical centers. Written informed consent was obtained from study participants.

Laboratory assays

Stable isotope dilution LC-MS/MS was used to quantify 15 estrogens and estrogen metabolites including: estrone, estradiol, 2-pathway metabolites (2-hydroxyestrone, 2-methoxyestrone, 2-hydroxyestradiol, 2-methoxyestradiol, and 2-hydroxyestrone-3-methyl ether); 4-pathway metabolites (4-hydroxyestrone, 4-methoxyestrone, and 4-methoxyestradiol); and 16α -pathway metabolites (16α -hydroxyestrone, estriol, 16 -ketoestradiol, 16 -epiestriol, and 17 -epiestriol). Details of the method have been published previously (22). For this study, six labeled internal standards were used: deuterated 2-hydroxyestradiol, 2-methoxyestradiol and estriol (C/D/N Isotopes Inc, Pointe-Claire); deuterated 16 -epiestriol (Medical Isotopes Inc); and ^{13}C -labeled estrone and estradiol (Cambridge Isotope Laboratories).

In serum, this method detects 15 estrogens and estrogen metabolites which circulate, at least in part, as sulfated and/or glucuronidated conjugates to facilitate storage, transport, and excretion. Five of the estrogens (estrone, estradiol, estriol, 2-methoxyestrone, and 2-methoxyestradiol) were also measured in unconjugated forms in circulation. The serum sample was split into two aliquots, one to measure the combined concentration of

each of the 15 metabolites (that is, the sum of conjugated plus unconjugated forms); the other, to measure the unconjugated forms. To measure the combined level of the conjugated plus unconjugated parent estrogen or estrogen metabolite level, an enzyme with sulfatase and glucuronidase activity was added to the samples to cleave any sulfate and glucuronide groups (22). To measure the unconjugated forms only, addition of the enzyme is not included in the sample preparation steps. For those metabolites with both combined and unconjugated measurements, the concentration of the conjugated form was calculated as the difference between the combined estrogen measurement and the unconjugated estrogen measurement, for estradiol that calculation was (conjugated E2 = combined E2 – unconjugated E2). The limit of detection for each estrogen and estrogen metabolite measured using this liquid chromatography/tandem mass spectrometry (LC/MS-MS) assay was 10 fg on column (approximately 0.33–0.37 pmol/L; refs. 22, 23). There were no samples in the current study with undetectable levels for any of the hormones measured. Laboratory coefficients of variation (CV) of masked technical replicates across batches were $< 6.0\%$ for all hormones measured. Intraclass correlation coefficients (ICC) ranged from 0.93 to 0.996 with median value of 0.98.

Statistical analysis

Estrogens and estrogen metabolites were analyzed individually. Individual estrogens were categorized into quintiles based on the distribution in controls. First, we analyzed associations in all ovarian cancer cases (ovarian cancer overall), followed by analyses stratified by subtype (serous/nonserous). Conditional logistic regression models were used to estimate ORs and 95% confidence intervals (CI) of ovarian cancer risk conditioning on matching factors: age at blood draw, calendar year of blood draw, race/ethnicity, hysterectomy status, and time since last menopausal hormone use and further adjusted for *a priori* potential confounding factors: gravidity, BMI (< 25 , 25 – 29.9 , ≥ 30 kg/m²), cigarette smoking status (never, former, current), duration of oral contraception use (never, < 5 , 5 – < 10 , ≥ 10 years), and menopausal hormone use (former/never). Tests for trend were based on the Wald statistic modeling the intracategory quintile median as a continuous parameter. Q-values reflecting the false discovery rates (FDR) were calculated to account for multiple comparisons in the main analysis and analyses by histologic subtype, all other analyses were considered exploratory and therefore not corrected for multiple comparisons. For analyses stratified by case characteristics, we used baseline category polytomous logistic regression models, with the controls as the reference group and adjustment for matching factors and *a priori* selected potential confounding factors (listed above), with one exception: menopausal hormone therapy use and time since last menopausal hormone therapy use were combined (never, ≤ 1 year, > 1 year). We also evaluated associations stratified by time between blood collection and diagnosis (< 5 year, ≥ 5 years) and by age at blood draw. Given the strong associations for many of the estrogens and estrogen metabolites evaluated with non-serous tumors, we evaluated the extent to which unconjugated estradiol (an estrogen strongly correlated with BMI among postmenopausal women) might explain the weak association between BMI and nonserous tumors. We compared the ORs for a 5 kg/m² increase in BMI, with and without further adjustment for unconjugated estradiol. We also conducted the following sensitivity analyses: (i) excluding individuals diagnosed within 2 years of blood draw ($n = 14$),

(ii) excluding potential outliers [greater than five SDs above the median; median excluded subject per hormone measure $n = 7$; (min-max: 3–10)], (iii) excluding women who reported a history of diabetes at baseline ($n = 29$), and (iv) excluding women who reported prior use of menopausal hormones ($n = 210$). All P values were based on two-sided tests, for uncorrected tests P values < 0.05 were considered statistically significant.

Results

Characteristics of cases (overall and by subtype) and controls at baseline are presented in Table 1. Participants were on average 64 years of age at blood draw and predominantly white (90%). Serous

cases were slightly older, on average 64.7 years of age at blood draw compared to 63.2 years of age among nonserous cases. Median levels of estrogens and estrogen metabolites are compared by case-control status and subtype in Supplementary Table S1. Significantly higher levels of estrone and unconjugated 2-methoxyestrone were observed in ovarian cancer cases than controls. The median levels of the remaining estrogens and estrogen metabolites were slightly higher among ovarian cancer cases compared with controls, albeit differences were not statistically significant. There were no differences in median levels of the hormones comparing serous cases and controls. Significantly higher levels of estrone, unconjugated estradiol, and a number of other estrogen metabolites were observed in nonserous cases than controls.

Table 1. Demographic and health characteristics controls and ovarian cancer cases by histologic subtype, nested case-control study within the WHI-OS

Characteristics	Controls	Ovarian cancer cases	Serous	Nonserous
	$n = 412$ Mean (SD)	$n = 169$ Mean (SD)	$n = 102$ Mean (SD)	$n = 67$ Mean (SD)
Age	64.3 (7.2)	64.1 (7.2)	64.7 (7.2)	63.2 (7.1)
Year of blood draw	n^a (%)	n^a (%)	n^a (%)	n^a (%)
1993–1996	254 (61.7)	108 (63.9)	62 (60.8)	46 (68.7)
1997–1998	158 (38.3)	61 (36.1)	40 (39.2)	21 (31.3)
Race/ethnicity				
White	371 (90.0)	151 (89.3)	96 (94.1)	55 (82.1)
Black	17 (4.1)	8 (4.7)	3 (2.9)	5 (7.5)
Hispanic	12 (2.9)	6 (3.6)	2 (2.0)	4 (6.0)
Other	12 (2.9)	4 (2.4)	1 (1.0)	3 (4.5)
Hysterectomy (baseline or follow-up prior to event)				
No	340 (82.5)	140 (82.8)	83 (81.4)	57 (85.1)
Yes	72 (17.5)	29 (17.2)	19 (18.6)	10 (14.9)
Time since last menopausal hormone therapy use				
>1 year	400 (97.1)	163 (96.4)	97 (95.1)	66 (98.5)
≤1 year	12 (2.9)	6 (3.6)	5 (4.9)	1 (1.5)
Family income				
<\$35,000	161 (39.1)	69 (40.8)	41 (40.2)	28 (41.8)
\$35,000–\$74,999	153 (37.1)	65 (38.5)	40 (39.2)	25 (37.3)
\$75,000+	65 (15.8)	24 (14.2)	16 (15.7)	8 (11.9)
Smoking status				
Never	203 (49.3)	81 (47.9)	51 (50.0)	30 (44.8)
Former	168 (40.8)	76 (45.0)	43 (42.2)	33 (49.3)
Current	38 (9.2)	12 (7.1)	8 (7.8)	4 (6.0)
BMI (kg/m ²)				
<25	181 (43.9)	66 (39.1)	43 (42.2)	23 (34.3)
25–29.9	127 (30.8)	52 (30.8)	31 (30.4)	21 (31.3)
30+	103 (25.0)	51 (30.2)	28 (27.5)	23 (34.3)
Age at menarche				
<12	100 (24.3)	37 (21.9)	24 (23.5)	13 (19.4)
12–13	220 (53.4)	100 (59.2)	59 (57.8)	41 (61.2)
14+	89 (21.6)	32 (18.9)	19 (18.6)	13 (19.4)
Ever pregnant				
No	54 (13.1)	25 (14.8)	13 (12.7)	12 (17.9)
Yes	358 (86.9)	144 (85.2)	89 (87.3)	55 (82.1)
Duration oral contraceptive use				
Never	254 (61.7)	106 (62.7)	65 (63.7)	41 (61.2)
<5 years	84 (20.4)	36 (21.3)	23 (22.5)	13 (19.4)
5–<10 years	39 (9.5)	14 (8.3)	5 (4.9)	9 (13.4)
10+ years	35 (8.5)	13 (7.7)	9 (8.8)	4 (6.0)
Age at menopause				
<40	17 (4.1)	4 (2.4)	3 (2.9)	1 (1.5)
40–44	33 (8.0)	14 (8.3)	11 (10.8)	3 (4.5)
45–49	91 (22.1)	34 (20.1)	18 (17.6)	16 (23.9)
50–54	183 (44.4)	83 (49.1)	52 (51.0)	31 (46.3)
55+	66 (16.0)	22 (13.0)	14 (13.7)	8 (11.9)
Menopausal hormone therapy use				
Never	242 (58.7)	129 (76.3)	74 (72.5)	55 (82.1)
Former	170 (41.3)	40 (23.7)	28 (27.5)	12 (17.9)

^aValues may not sum to total because of missing data.

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Table 2. OR and 95% CI for the risk of epithelial ovarian cancer comparing the 5th quintile to the 1st quintile for individual estrogens and estrogen metabolites, nested case-control study within the WHI-OS

	OR ^a (95% CI)	P _{trend} ^b	FDR ^c
Parent estrogens			
Estrone	1.54 (0.82-2.90)	0.05	0.42
Unconjugated estrone	1.47 (0.78-2.79)	0.25	0.48
Conjugated estrone	1.54 (0.81-2.92)	0.09	0.42
Estradiol	0.88 (0.47-1.62)	0.89	0.95
Unconjugated estradiol	1.50 (0.77-2.91)	0.39	0.54
Conjugated estradiol	1.10 (0.59-2.07)	0.95	0.95
2-Hydroxylation pathway			
2-Hydroxyestrone	1.31 (0.68-2.53)	0.33	0.49
2-Hydroxyestradiol	0.93 (0.49-1.79)	0.92	0.95
2-Methoxyestrone	2.03 (1.06-3.88)	0.02	0.25
Unconjugated 2-methoxyestrone	1.44 (0.74-2.79)	0.29	0.49
Conjugated 2-methoxyestrone	1.72 (0.90-3.26)	0.10	0.42
2-Methoxyestradiol	1.55 (0.81-2.97)	0.21	0.48
Unconjugated 2-methoxyestradiol	1.68 (0.88-3.18)	0.21	0.48
Conjugated 2-methoxyestradiol	1.36 (0.72-2.58)	0.30	0.49
2-Hydroxyestrone-3-methyl ether	1.11 (0.57-2.16)	0.53	0.66
4-Hydroxylation pathway			
4-Hydroxyestrone	1.46 (0.76-2.81)	0.17	0.48
4-Methoxyestrone	1.86 (0.98-3.56)	0.01	0.25
4-Methoxyestradiol	1.74 (0.90-3.40)	0.13	0.46
16 α -Hydroxylation pathway			
16 α -Hydroxyestrone	1.29 (0.68-2.45)	0.32	0.49
Estriol	1.61 (0.84-3.07)	0.10	0.42
Unconjugated estriol	1.21 (0.62-2.36)	0.46	0.61
Conjugated estriol	1.35 (0.70-2.58)	0.19	0.48
16-Ketoestradiol	1.32 (0.70-2.50)	0.24	0.48
16-Epiestriol	1.14 (0.58-2.24)	0.92	0.95
17-Epiestriol	1.32 (0.66-2.63)	0.68	0.81

^aOR from model conditioned on matching factors (age at baseline, year of blood draw, race/ethnicity, hysterectomy status, and time since last menopausal hormone use) and adjusted for body mass index, smoking status, gravidity, duration of oral contraceptive use, and never/former menopausal hormone therapy use.

^bP values for trend across quintile (median value of category).

^cFDR *q*-value.

Estrone was positively associated with overall ovarian cancer risk ($P_{\text{trend}} = 0.05$); however, the OR across extreme quintiles of estrone was not statistically significant [Q5 vs. Q1: 1.54 (0.82–2.90); Table 2]. Estradiol was not associated with overall ovarian cancer risk. Only two of the individual estrogen metabolites were significantly associated with overall ovarian cancer risk: compared with women in the lowest quintile, women with the highest quintile of 2-methoxyestrone were at increased risk of ovarian cancer [Q5 vs. Q1: 2.03 (1.06–3.88); $P_{\text{trend}} = 0.02$]. A positive trend across quintile of 4-methoxyestrone was associated with risk ($P_{\text{trend}} = 0.01$); however, the OR comparing extreme quintiles did not reach statistical significance [Q5 vs. Q1: 1.86 (0.98–3.56)]. The ORs for these associations remained elevated in models adjusted for estrone; however, they no longer retained statistical significance (results not shown).

Significant heterogeneity in OR associations for the parent estrogens (estrone and estradiol) as well as the combined measurements of the 2-, 4-, and 16 α -pathway metabolites was observed across histologic subtype of ovarian cancer (Fig. 2). For serous ovarian cancers, associations with the estrogens and estrogen metabolites were universally null, while nonserous tumors were significantly associated with a number of individual estrogen metabolites (Table 3). In analyses restricted to nonserous cases, positive trends and/or significant top versus bottom quintile OR associations were observed for 17 of the 25 individual estrogen measures evaluated. Estrone [2.65 (1.09–6.45), $P_{\text{trend}} = 0.01$, $P_{\text{heterogeneity}} = 0.04$] and unconjugated estradiol [2.72 (1.04–

7.14), $P_{\text{trend}} = 0.03$, $P_{\text{heterogeneity}} = 0.02$] were positively associated with nonserous tumors. For metabolites where the $P_{\text{heterogeneity}}$ was ≤ 0.01 , nonserous ovarian cancer risk was at least tripled for the following: 2-hydroxyestrone [Q5 vs. Q1: 4.27 (1.52–12.01)], 2-methoxyestrone [Q5 vs. Q1: 4.26 (1.68–10.81)], 16 α -hydroxyestrone [Q5 vs. Q1: 3.33 (1.17–9.50)], and 16-epiestriol [Q5 vs. Q1: 3.46 (1.18–10.15)].

Some previous studies have shown associations of increased BMI with nonserous ovarian cancers. Accordingly, nonserous tumors were positively associated with BMI (OR (95% CI) per 5 kg/m² increase in BMI: 1.21 (1.00–1.48)), while serous tumors were not associated with BMI in the current study. Given the strong positive associations for many of the estrogens and nonserous tumors, and the known correlation between unconjugated estradiol and BMI, we evaluated the extent to which unconjugated estradiol might explain the association between BMI and nonserous tumors (Table 4). After adjustment for unconjugated estradiol, the association between BMI and nonserous tumors was substantially attenuated and no longer statistically significant [per 5 kg/m² increase in BMI: 1.10 (0.88–1.38)]. The associations between unconjugated estradiol and nonserous tumors remained statistically significant after adjustment for potential confounding effects of BMI [OR (95% CI) for a doubling in unconjugated estradiol without adjustment for BMI: 1.40 (1.13–1.73); and with adjustment for BMI: 1.36 (1.08–1.71)].

Results were not significantly modified by time between blood draw and diagnosis or age at blood draw (Supplementary

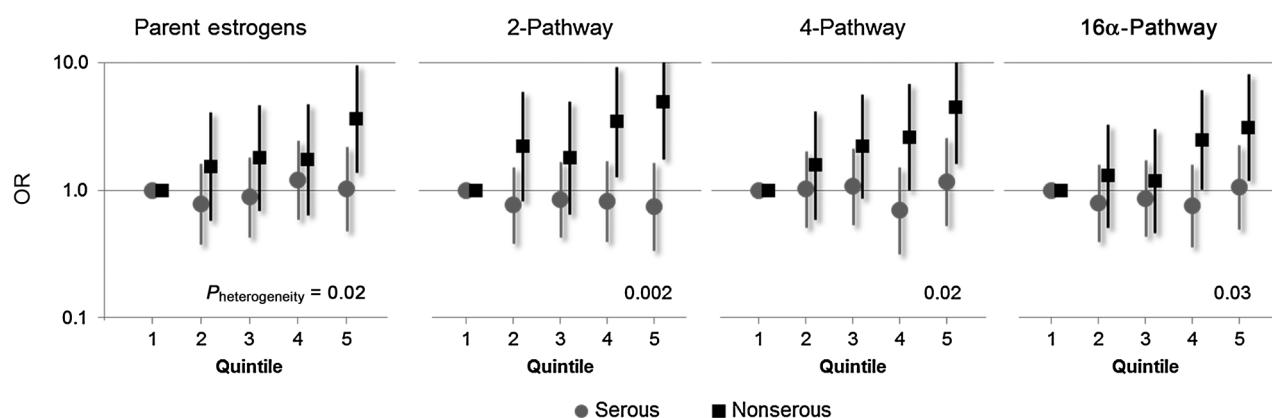


Figure 2.

Quintile-specific ORs and 95% CIs for the risk of epithelial ovarian cancer subtypes with estrogen metabolite pathways, nested case-control study within the WHI-OS.

Tables S2 and S3, respectively). In sensitivity analyses excluding cases diagnosed within two years of blood draw, excluding outliers for individual estrogen measurements, excluding individuals with a history of diabetes at baseline, or excluding former hormone users, effect estimates were largely unchanged (results not shown).

Discussion

There is strong *in vitro* and *in vivo* evidence, as well as some epidemiologic data, suggesting that estrogens (and potentially other sex steroids) play a role in ovarian carcinogenesis. However, previous studies did not show associations between circulating

Table 3. OR and 95% CIs for the risk of epithelial ovarian cancer subtypes comparing the 5th quintile to the 1st quintile for individual estrogens and estrogen metabolites, nested case-control study within the WHI-OS

	Serous <i>n</i> = 102		Nonserous <i>n</i> = 67		<i>P</i> _{het} ^c	FDR	
	OR ^a (95% CI)	<i>P</i> _{trend} ^b	OR ^a (95% CI)	<i>P</i> _{trend} ^b		Trend ^d	Het ^d
Parent estrogens							
Estrone	1.11 (0.52–2.33)	0.56	2.65 (1.09–6.45)	0.01	0.04	0.02	0.05
Unconjugated estrone	1.14 (0.52–2.50)	0.67	1.95 (0.86–4.41)	0.01	0.01	0.02	0.04
Conjugated estrone	1.03 (0.48–2.21)	0.71	3.15 (1.24–8.01)	0.01	0.03	0.02	0.05
Estradiol	0.60 (0.29–1.23)	0.42	1.79 (0.71–4.51)	0.22	0.09	0.23	0.09
Unconjugated estradiol	0.97 (0.44–2.16)	0.58	2.72 (1.04–7.14)	0.03	0.02	0.04	0.04
Conjugated estradiol	0.81 (0.38–1.72)	0.41	1.63 (0.67–3.96)	0.27	0.16	0.27	0.16
2-Hydroxylation pathway							
2-Hydroxyestrone	0.76 (0.35–1.65)	0.72	4.27 (1.52–12.01)	0.01	0.01	0.02	0.04
2-Hydroxyestradiol	0.71 (0.33–1.55)	0.42	1.70 (0.68–4.26)	0.11	0.02	0.13	0.04
2-Methoxyestrone	1.27 (0.59–2.73)	0.58	4.26 (1.68–10.81)	0.001	0.01	0.01	0.04
Unconjugated 2-methoxyestrone	0.87 (0.39–1.96)	0.85	3.20 (1.15–8.92)	0.02	0.01	0.03	0.04
Conjugated 2-methoxyestrone	0.98 (0.45–2.14)	0.86	3.66 (1.46–9.19)	0.003	0.02	0.02	0.04
2-Methoxyestradiol	1.11 (0.51–2.39)	0.99	2.36 (0.94–5.94)	0.03	0.05	0.04	0.06
Unconjugated 2-methoxyestradiol	1.23 (0.56–2.70)	0.94	2.96 (1.18–7.44)	0.02	0.04	0.03	0.05
Conjugated 2-methoxyestradiol	0.89 (0.41–1.90)	0.74	2.32 (0.93–5.79)	0.03	0.05	0.04	0.06
2-Hydroxyestrone-3-methyl ether	0.76 (0.36–1.64)	0.80	2.36 (0.84–6.65)	0.13	0.08	0.14	0.09
4-Hydroxylation pathway							
4-Hydroxyestrone	1.00 (0.46–2.16)	0.80	3.13 (1.21–8.06)	0.01	0.02	0.02	0.04
4-Methoxyestrone	1.21 (0.57–2.58)	0.32	3.25 (1.31–8.10)	0.001	0.03	0.01	0.05
4-Methoxyestradiol	1.20 (0.55–2.65)	0.99	2.91 (1.11–7.64)	0.01	0.02	0.02	0.04
16α-Hydroxylation pathway							
16α-Hydroxyestrone	0.76 (0.35–1.65)	0.76	3.56 (1.32–9.64)	0.02	0.01	0.03	0.04
Estriol	1.10 (0.51–2.36)	0.60	3.76 (1.41–10.01)	0.01	0.02	0.02	0.04
Unconjugated estriol	0.85 (0.37–1.93)	0.47	2.37 (0.96–5.85)	0.01	0.001	0.02	0.03
Conjugated estriol	0.99 (0.46–2.14)	0.80	2.55 (1.02–6.40)	0.02	0.04	0.03	0.05
16-Ketoestradiol	0.91 (0.42–1.95)	0.96	2.98 (1.16–7.63)	0.01	0.02	0.02	0.04
16-Epiestradiol	0.65 (0.29–1.48)	0.25	3.46 (1.18–10.15)	0.05	0.01	0.06	0.04
17-Epiestradiol	0.96 (0.44–2.09)	0.74	2.85 (0.97–8.34)	0.12	0.05	0.14	0.06

^aOR from model adjusting for matching factors (age at baseline, year of blood draw, race/ethnicity, hysterectomy status), recency of menopausal hormone therapy use (never, ≤1, >1), body mass index, smoking status, gravidity, and duration of oral contraceptive use.

^b*P* values for trend across quintile (median value of category).

^c*P*_{heterogeneity}.

^dFDR *q*-value corresponding to the *P*_{trend} among nonserous tumors and the test for heterogeneity across subtype.

Table 4. OR and 95% CIs evaluating the independent effects of body mass index (BMI) and unconjugated estradiol on ovarian cancer risk

		All ovarian cancer OR ^b (95% CI)	Serous OR ^b (95% CI)	Nonserous OR ^b (95% CI)
BMI (per 5 kg/m ²)	Not adjusted for estradiol	1.09 (0.94-1.27)	1.02 (0.84-1.23)	1.21 (1.00-1.48)
	Adjusted for estradiol	1.07 (0.89-1.27)	1.06 (0.85-1.31)	1.10 (0.88-1.38)
Unconjugated estradiol ^a	Not adjusted for BMI	1.12 (0.97-1.30)	0.98 (0.81-1.17)	1.40 (1.13-1.73)
	Adjusted for BMI	1.10 (0.93-1.30)	0.96 (0.78-1.18)	1.36 (1.08-1.71)

^aLog-2 transformed unconjugated estradiol, OR associated with a doubling in circulating level of unconjugated estradiol.

^bOR adjusting for matching factors (age at baseline, year of blood draw, race/ethnicity, hysterectomy status), recency of menopausal hormone therapy use (never, ≤1, >1), smoking status, gravidity, and duration of oral contraceptive use. Additional adjustment for unconjugated estradiol or BMI as listed in second column.

estrogens and overall ovarian cancer risk. In the current study, we demonstrate significant heterogeneity of associations between estrogen metabolites and ovarian cancer subtypes. We did not find associations with serous cancers, the most common and fatal subtype, and consequently, we did not see strong associations with overall ovarian cancer risk. However, for the first time, we provide evidence suggesting that estrogen and estrogen metabolites in postmenopausal women are associated with nonserous ovarian cancers.

In a previous study of early pregnancy hormones and subsequent cancer risk, Shock and colleagues (16) evaluated associations by histologic subtype and reported that higher levels of unconjugated estradiol in pregnancy were associated with increased risk of endometrioid ovarian tumors. They also reported elevated ORs (T3 vs. T1 OR ranging from 1.54 to 1.80) for the association between estradiol and other nonserous subtypes including borderline and invasive mucinous and clear cell tumors, while ORs for serous tumors were null (OR borderline serous: 0.87; invasive serous: 0.98; ref. 16). Although pregnancy levels of estrogens are high and may not correlate with non-pregnancy levels or levels among postmenopausal women, the increased risk associated with higher concentrations of the parent estrogens in our nonserous cases is consistent with the findings from the maternity cohort.

The three other prospective studies evaluating estrogens and overall ovarian cancer risk all reported null associations for the parent estrogens (13–15). This could be due to relatively limited number of available cases [$n = 31$ (13), $n = 132$ across 3 cohorts (14), $n = 67$ (15)], but is most likely due to the predominance of serous tumors in these studies, as we observed null associations with parent estrogens and a very limited number of positive associations with estrogen metabolites in our analysis of overall ovarian cancer risk. In the only other study of estrogens metabolites to date, 4-methoxyestrone was associated with a significant increased risk of overall ovarian cancer in a case-cohort study ($P_{\text{trend}} = 0.04$), however the other 2-, 4-, and 16-pathway metabolites evaluated were not significantly associated with overall risk (15). We note a similar elevated risk with 4-methoxyestrone in our overall ovarian cancer analysis ($P_{\text{trend}} = 0.01$), which is likely driven, in part, by substantially elevated risk among non-serous tumors (OR = 3.25). The findings from these early studies (13, 14) are not directly comparable with our results, given inclusion of pre- and perimenopausal women and measurement of parent estrogens using RIA, which have recognized limitations. While the findings from the recent case-cohort study are directly comparable, given measurement of estrogens using the same assay and that both study populations are primarily Caucasian, postmenopausal women with similar ages at blood draw and diagnosis (15).

The subtype-specific associations we observed in the current study are consistent with positive associations between BMI and nonserous tumors, and null associations between BMI and serous tumors, reported in prospective cohort studies (17–19). This is not surprising given that adipose tissue is a source of circulating estrogens via peripheral conversion of androgens in postmenopausal women (7). Ovarian cancer associations for other hormonal risk factors demonstrated more variability across the studies, but the extent to which these variables correlate with endogenous estrogen levels in postmenopausal women has not been well characterized. Associations with exogenous hormones, namely unopposed estrogen or estrogen plus progestin menopausal hormone therapy, suggest increased risks of both serous and nonserous subtypes (17, 18). Thus, the lack of association between circulating estrogens and serous tumors in the current study is at odds with results from the exogenous estrogen data. However, in women taking menopausal hormones, circulating levels of estrogens are severalfold higher than in postmenopausal women not currently taking exogenous hormones and have only limited applicability to the endogenous measures in the current study. Combined with the accumulating evidence of etiologic heterogeneity of epithelial ovarian cancers, particularly for hormonally related factors, the subtype-specific associations we observed in the current study support the evaluation of other circulating sex steroid hormones, namely androgens and progesterone, by ovarian cancer subtypes to further clarify the hormonal mechanisms that underlie the etiology of ovarian cancer.

The current study has several important strengths. The WHI-OS cohort is a large prospective study with standardized prediagnostic specimen collection and storage. The five estrogens and estrogen metabolites found in circulation in both unconjugated and combined forms and 10 measured in combined form provide a novel phenotypic characterization of individual patterns of estrogen metabolism. We were able to measure estrogens among all postmenopausal women in our study population, including those with relatively low levels, through the use of an LC/MS-MS assay with high sensitivity (22, 23). This study also has some limitations. Although we included all available cases, the study was still limited in power, which affected our ability to evaluate specific subtypes of nonserous tumors. While somewhat imprecise, the associations with nonserous tumors were robust to correction for multiple comparisons. We measured circulating estrogens in a single baseline serum sample, which may not accurately reflect long-term exposure and/or intraovarian levels. Prior research by our group suggests that among postmenopausal women reproducibility was moderate-to-high for most of the estrogens and estrogen metabolites (e.g., 3-year ICCs greater than 0.65 and 0.72 for estrone and estradiol, respectively and on

average greater than 0.25, 0.29, and 0.45 for the 2-, 4-, and 16-pathway metabolites, respectively; ref. 24).

Our study provides novel molecular data that supports a role for estrogens and estrogen metabolites in nonserous ovarian cancer, and suggests etiologic heterogeneity across histologic subtypes of ovarian cancer. Furthermore, the association of obesity with nonserous tumors appears to be explained, in part, by circulating estradiol. Additional investigation in a large prospective study is needed to clarify the risk of individual ovarian cancer subtypes for specific estrogen metabolites.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

Authors' Contributions

Conception and design: L.A. Brinton, R.T. Falk, B.J. Fuhrman, N. Wentzensen
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Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): L.A. Brinton, G.L. Anderson, B.J. Fuhrman, X. Xu
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): B. Trabert, L.A. Brinton, R.M. Pfeiffer, R.T. Falk, H.D. Strickler, L.H. Kuller, N. Wentzensen
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Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): B. Trabert, L.A. Brinton
Study supervision: B. Trabert

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For a list of all the investigators who have contributed to WHI science, please visit: <https://www.whi.org/researchers/Documents%20Write%20a%20Paper/WHI%20Investigator%20Long%20List.pdf>

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