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Fat/Fiber intakes and sex hormones in healthy premenopausal women in USA

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Abstract

The mechanisms by which diet affects breast cancer (BC) risk are poorly understood but a positive relationship between fat and a negative association with fiber intake and BC risk have been demonstrated. Here we study the association between dietary fat/fiber ratio and estrogen metabolism. Fifty women were recruited, 22 were included in the low fat/high fiber and 22 were in the high fat/low fiber and 6 did not meet our criteria. Estrogens (determined in plasma, urine and feces) and dietary records were collected during 3 following days. All data were collected in winter and in summer. The high fat/low fiber group had significantly higher urinary total estrogens, estriol-3-glucuronide, 2-hydroxyestradiol, 16 α -hydroxyestrone, and a higher 2-hydroxyestrone/4-hydroxyestrone ratio. Total fat intake correlated significantly with plasma estrone, estradiol, urinary 2-hydroxyestrone, 2-hydroxyestradiol, 2-hydroxyestrone/4-hydroxyestrone ratio, and total urinary estrogens, even after adjustment for total fiber intake. The high fat/low fiber diet was associated with high values both for catechol and 16 α -hydroxylated estrogens and a high 2-hydroxyestrone/4-hydroxyestrone ratio, but 2-hydroxyestrone/16 α -hydroxyestrone ratio was not different between the groups. Our results suggest that fat affects estrogen metabolism more than does fiber and that one mechanism resulting in high estrogen values is an increased reabsorption of biliary estrogens.

Keywords

Diet; Estrogens; Breast cancer risk; Women

1. Introduction

Breast cancer (BC) incidence is significantly lower in Asian than in Western women [1]. One explanation is that traditional Asian diets may be protective because they are often rich in soy

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products and lower in fat [2]. The mechanism by which diet influences BC risk is unclear although it has been shown that changes in dietary habits may increase or decrease BC risk [3]. A meta-analysis and a pooled study [4,5] demonstrated positive relationship between fat and BC risk, but negative results have also been reported [6,7].

More recently, the Women's Health Initiative Dietary Modification Trial (WHI-DMT, [8]) found a statistically non significant 9% reduction of BC incidence in the low-fat intervention group compared to a control group. That study also reported significant differences between the intervention and control groups for body weight, total energy, and vegetable fiber intake; factors which are also thought to influence BC risk. Alcohol intake was similar and unchanged in both groups.

It is believed that fiber consumption protects against BC by binding enteral carcinogens, and estrogens during their enterohepatic circulation thus facilitating their fecal elimination [9,10]. Reduction of intestinal β -glucuronidase activity, necessary for hydrolysis of the estrogen conjugates before absorption, is another mechanism diminishing estrogen levels [11]. In premenopausal Finnish women total fiber or grain fiber intake/kg body weight showed a statistically significant negative association with the excretion of 10 of 13 estrogens measured in urine [12]. An inverse relationship between high fiber intake and BC risks was noted in a Swedish prospective study [13]. However, in most other prospective studies the associations between dietary fiber and BC were null [14]. In a meta-analysis, Wu et al. demonstrated that 4/14 interventional studies showed that a high level of dietary fiber combined with a low level of fat had an effect on BC risk [15].

Evidence from both epidemiologic and biomolecular studies indicate that estrogens are implicated in human BC [16–18]. Some cohort studies have revealed a strong relationship between endogenous estrogen levels and BC risk [19,20]. In one study [21] but not in another [22], women who were switched from a high-fat (40% of calories) to a low fat (25%) showed a significant decrease of the level of 16α -OHE1 which is recognized to be a potent estrogen receptor agonist [23]. It has been demonstrated that the level of 16α -OHE1 is elevated in women with BC [24]. In addition it was suggested that 2-OHE1 is only weakly estrogenic and possibly antiestrogenic [24] and it has been assumed that a high level could reduce BC risk. However, it has also been shown that in postmenopausal women a high catechol estrogen concentration is associated with increased BC risk [25]. Catechol estrogens are thought to be carcinogenic because of their capability to form DNA adducts [17]. Furthermore, it has been suggested that the ratio 2-OHE1/ 16α -OHE1 may serve as a marker of BC risk [26]. Early evidence indicated that postmenopausal vegetarians, known to consume twice the fiber than omnivores, had significantly lower blood estrogen and androgen levels [27–29]. Cross-sectional studies among postmenopausal women with low fiber intakes reported no association between fiber and endogenous estrogen levels [30–32]. Recently, Monroe et al. demonstrated that fiber intake was significantly negatively associated with a decrease of serum E1 and E2 in postmenopausal Mexican-American women [33].

The aim of this study was to examine the association between, on the one hand, a high fat/low fiber Western-type diet, and on the other hand, an ideal low fat/high fiber diet with the sex hormone pattern in plasma and urine, including some catechol estrogens, in premenopausal healthy women living in USA. To our knowledge no such study on subjects consuming their habitual diets has been carried out. To obtain information about the enterohepatic circulation of estrogens we also included assays of estriol-3-glucuronide in urine, which is a marker of conjugation in the mucosal cells of the gastrointestinal tract.

2. Materials and methods

2.1. Subjects

The participants were premenopausal healthy volunteer women and were recruited in the Boston area (USA). Women who had maintained their habitual diet, including omnivores, lacto-ovo-vegetarians and vegans, for at least 2 years were recruited by advertisements in local newspapers, bulletins and magazines. Subjects were excluded if they had: a BMI over 120 percent or under 90 percent of ideal body weight, history of heart disease, history of cancer, arthritis, hypertension, diabetes mellitus, thyroid problems, renal or liver disease, irregular menstrual cycle, regular consumption of alcohol (≥ 15 g/d) or using prescribed medications (oral contraceptives, corticosteroids or antibiotics).

For the purpose of this study, subjects were divided by the percent of fat of energy intake and the amount of fiber consumed per day in two groups. Based on their dietary intakes, the two groups included: a low fat/high fiber group consisting of women who consumed less than 21% of energy/day as fat and more than 21g/day of fiber, and a high fat/low fiber group made up of women with $>21\%$ calories from fat and less than 21 g fiber per day. Among the 50 premenopausal subjects studied, 22 women were classified in low fat/high fiber group and 22 women in high fat/low fiber group (6 women could not be included in our analysis because their food intakes did not match the inclusion criteria used).

2.2. Collection of samples

Subjects were studied for 3 consecutive days on 2 occasions (one visit in the summer and one in the winter season). An average of the winter and summer values was used in the statistical analyses to counteract the seasonal effect. In fact, it is recognized that diet during the seasons are different [34]. Each of these visits included a 3 times 24-h collection of urine and feces (totally 6 days of collection), 3 different blood samples in heparinized tubes during the morning after urine collection and 3-day food records. The 3-day blood samples were pooled to reduce the number of analyses. All samples were collected during the mid-follicular phase of the menstrual cycle [days (3)5–7(9)]. To prevent oxidation of the hormones and stop bacterial growth, we added as preservative 0.1 % of ascorbic acid and 0.1 % of sodium azide in heparinized tubes. One g of ascorbic acid powder per liter of volume was added to the plastic bottle before urine collection. Feces were collected in plastic containers and after the first defecation 10 ml of 1 % ascorbic acid was poured on the sample. The urine and feces samples were stored in a cold place (in the winter outside) and brought to the hospital every day and then 0.1 % sodium azide added. The samples for 3 days were pooled and feces homogenized. Suitable volumes of the urine and feces samples for analysis were transferred to new tubes. The storage was at -20 °C. The samples were sent to Helsinki frozen in dry ice.

2.3. Hormones

The plasma samples were collected from fasting subjects. All analyses in duplicate were carried out within a year of sample collection. If the results of the duplicates deviated more than 10 % from the mean value the analysis was repeated. Quality control samples based on pooled samples that had been analyzed ten times, were included in each batch.

Plasma estrogens, estrone (E1) and estradiol (E2), were determined by radioimmunoassay (RIA) after chromatographic separation on an LH-20 column as described in detail [29,35]. After addition of tritiated internal standards, the plasma was extracted with 5 volumes of 20 per cent ethyl acetate in petroleum ether (vol/vol) in one step by freezing. After chromatography on Sephadex LH-20 with methanol-toluene 9:1 (vol/vol), the separated estrogens were determined by radioimmunoassay, with free and bound ligand separated by dextran-coated charcoal [35]. The coefficient of variation (CV%), which was calculated from the differences

between duplicate analyses of a control serum pool, was 5.1 % for E1 and 3.9 % for E2. The corresponding inter-assay CV% calculated from the mean values of these duplicates during a 3-year period (n = 15) was 2.0 % for E1 and 1.4 % for E2 [29].

Plasma testosterone (T) and androstenedione (A) were determined by radioimmunoassay as described in detail by Kuoppasalmi et al. [36]. Free testosterone (Free T) was measured according to Bergink et al. [37] and sex hormone binding globulin (SHBG) as described by Rosner [38] with slight modifications [39]. Dehydroepiandrosterone-sulfate (DHEAS) was measured with a commercial kit (Wien Laboratories,inc. Succansunna, NJ,USA). The intra-assay CV% were for T 8.3 %, for A 5.7 %, for DHEAS 6.2 % for SHBG 8.8 %, and for free T 7.1 %. The inter-assay CV% were for T 12.7 %, for A 9.2 %, for DHEAS 7.5 %, for SHBG 11.1 %, and for free T 8.5 %.

Urinary estrogen metabolites were measured by GC/MS in the selected ion monitoring mode as described by Fotsis et al. [40]. All details including validation of the procedure were published [40]. Urinary “immunoreactive” estriol-3-glucuronide (E3-3G), a biomarker of enterohepatic circulation of estrogens, was determined by radioimmunoassay as described [41] using reagents kindly provided by Drs A. Kellie and P. Samarajeewa. This estriol conjugate is formed exclusively in the mucosal cells of the intestine and thereafter excreted without further metabolism in the urine and may, therefore, serve as a biomarker of the reabsorption of biliary estrogens in the intestine.

Fecal analyses of unconjugated E1, E2 and E3 were carried out by RIA using a method previously described in detail [42]. The inter-assay CV% for E1 is 5.8 %, for E2 5.9 % and for E3 1.7 %.

All the methods have been validated in regard to the accuracy, sensitivity and reproducibility. The E3-3G method probably also measures small amounts of other 16-hydroxylated estrogen-3-glucuronides, but this does not invalidate its use as a biomarker of the extent of enterohepatic circulation of estrogens, because the intestinal mucosa is the only organ where 3-glucuronidation of polar estrogens occur. Plasma or serum for SHBG analyses cannot be stored for long periods of time because this leads to the combination of two SHBG molecules (dimerization) and these dimers cannot be detected in the immunoassay resulting in low values. This problem was avoided in the present study. Our own studies and discussion on the methodological issues have been published [43].

2.4. Dietary intake

Subjects were instructed to maintain their usual diets throughout the three days of the dietary recordings in each season [44]. Each subject was provided with a postal scale balance and was instructed on how to complete the dietary records. It has been demonstrated that a 3-day dietary record is valid to estimate dietary intakes in adults without cognitive impairments [44]. Dietary analyses were completed by a nutritionist using the Nutrient Data Base at the University of Massachusetts at Amherst. Dietary fiber intake was calculated individually by nutritionist using the values of Paul and Southgate [45].

2.5. Statistical analysis

Results are presented as means \pm SD. Normality of distribution was determined using the Kurtosis test. Data were log-transformed if abnormally distributed. Homogeneity of variances was tested using the Levene test. We used an independent sample t-test to compare groups for all variables and a General Linear Model (GLM) multivariate analysis with total Kcal intake as the covariable. Finally, we used Pearson and partial correlation tests to examine the relation

between dietary components and sex hormone levels. $P \leq 0.05$ was considered significant (SPSS 15.00).

3. Results

We observed no significant differences in the anthropometric data between the 2 groups (Table 1). For dietary intake, we found significant differences between the groups for all the dietary variables except for protein intake and PUFA (Table 2). The high fat/low fiber group consumed more fat, SFA and MUFA and had higher energy intake. The fat/fiber ratio was, related to the grouping of the subjects, significantly higher in the high fat/low fiber group compared to the low fat/high fiber group.

Plasma E1 ($p = 0.069$), E1+E2 ($p = 0.067$) and SHBG ($p = 0.051$) were lower in the low fat/high fiber group compared to the high fat/low fiber group (Table 3).

We found significantly higher values in the high fat/low fiber group for the urinary E3 ($p = 0.025$), total estrogens ($p = 0.013$), E3-3G ($p = 0.002$), 2-OHE2 ($p = 0.004$), 16 α -hydroxylated-estrogens (16 α -OHE1+E3; $p = 0.032$), 2-OHE1/4-OHE1 ratio ($p = 0.045$), 16 β -hydroxyestrone ($p = 0.023$), 16-*epi*estriol ($p = 0.033$), and a tendency for the urinary 16 α -OHE1 ($p = 0.055$), 2-OH-estrogens ($p = 0.078$) and 16-keto-estradiol ($p = 0.067$) (Table 4).

We carried out a GLM multivariate analysis for all the plasma sex hormones and urinary estrogen metabolites using total energy intake as a covariable (Table 5). We observed significantly higher values in the high fat/low fiber group for the plasma E1+E2 ($p = 0.046$), plasma SHBG ($p = 0.045$), urinary E3-3G ($p = 0.024$), 16 α -OHE1 ($p = 0.024$) and 16 β -hydroxyestrone ($p = 0.024$). We found a trend toward higher values in the high fat/low fiber group for the plasma E1 ($p = 0.052$), plasma E2 ($p = 0.078$), urinary E2 ($p = 0.068$), 2-OHE2 ($p = 0.054$), 16-keto-estradiol ($p = 0.061$) and 16 α -OH-estrogens ($p = 0.072$).

Total energy intake correlated significantly and positively with the plasma E2, E1+E2, and urinary 2-OHE2, 2-OHE1/4-OHE1 ratio, 2-OHE and total urinary estrogens (Table 6).

Total fat intake correlated significantly and positively with the plasma E1, E2, E1+E2, and the urinary E2, E3-3G, 2-OHE1, 2-OHE2, 2-OHE1/4-OHE1 ratio, 2-OHE, 16 α -OH-estrogens, 16 β -hydroxyestrone, 16-*epi*estriol, 17-*epi*estriol and total urinary estrogens (Table 6). Total carbohydrate intake correlated significantly and negatively with the urinary E3-3G, 16 α -OHE1 and 16 β -OHE1 and positively with fecal E1 and E2 (Table 6). Total SFA and MUFA correlated significantly and positively with the plasma E1, E2, E1+E2, and the urinary E2, E3, E3-3G, 2-OHE1, 4-OHE1, 2-OHE2, 2-OHE1/4-OHE1 ratio, 4-OHE1/E1 ratio, 2-OHE, 16 α -OH-estrogens, 16 β -OHE1, 16-keto-estradiol, 16-*epi*estriol and total urinary estrogens. Fat/fiber ratio correlated significantly and positively with the plasma E1, E1+E2 and urinary E2, E3, E3-3G and 2-OHE2. Total fiber intake correlated significantly but negatively with the urinary E3, total estrogens, E3-3G, 16 α -OHE1, 16 α -OH-estrogens and 16 β -OHE1 (Table 6).

Urinary E3 (-0.360 , $p = 0.018$) and E3-3G (-0.370 , $p = 0.015$) still correlated negatively and significantly with the total fiber intake even after adjustment for the total fat intake. In addition, we found that the total fat correlated significantly with the plasma E1 (0.447, $p = 0.003$), E2 (0.358, $p = 0.018$), E1+E2 (0.432, $p = 0.004$), the urinary 2-OHE1 (0.499, $p = 0.007$), 2-OHE2 (0.646, $p = 0.001$), 2-OHE1/4-OHE1 ratio (0.540, $p = 0.003$), 2-OHE (0.552, $p = 0.002$) and all the urinary estrogens (0.563, $p = 0.002$) even after adjustment for total fiber intake. These results suggest that fat affects estrogen metabolism more than does fiber.

4. Discussion

The aims of this study were to investigate the hormonal pattern in healthy premenopausal women on a high fat/low fiber Western-type diet, which is associated with high BC risk, and to compare the results with women on a low fat/high fiber diet.

Our main results showed that women who consumed more fat and less fiber had significantly higher level of urinary E3-3G which is known as a reliable marker for the enterohepatic circulation [46,47], and higher catechol estrogens (particularly 2-OHE2), 2-OHE1/4-OHE1 ratio and 16 α -hydroxylated-estrogens, than women who ingested less fat and more fiber. Our results confirm and extend the results obtained previously by Goldin et al. [29,48]. In another study vegetarians had lower plasma E2, and fiber intake correlated negatively with plasma E2 [28]. Furthermore, another study [49] and an intervention study [50] indicated that fiber reduces plasma estrogens. Fat seems to have the opposite effect [15].

The inhibitory effect of dietary fiber on breast cancer could differ according to the type of fiber [51]. Approximately 50% of the conjugated estrogens in the liver are excreted into the bile and reabsorbed in the intestine [47]. It was postulated that insoluble fiber from cereal might increase the fecal excretion of estrogens by several mechanisms (see below). Fruit and vegetable fibers that are mainly soluble may not affect the enterohepatic circulation of the estrogens to the same extent as insoluble fiber. It is important to note that the major source of dietary fiber differs between countries [52]. For instance, cereal fiber accounts for approximately 70% of total fiber in Sweden while vegetables are a major source of dietary fiber in USA [53]. These differences could explain the controversial results in the literature.

An important observation is the significant difference between the groups for E3-3G that is higher in the high fat/low fiber group. This finding in combination with the tendency to low fecal estrogens compared to the low fat/high fiber group indicates that the reabsorption of biliary estrogens is more extensive in the high fat/low fiber group compared to the low fat/high fiber group. Estriol-3-glucuronide is exclusively formed in the intestinal mucosal cells and excreted unchanged in urine without further metabolism [54]. The increased enterohepatic circulation of estrogens shown by the results for E3-3G explains the higher total plasma (E1 +E2) and urinary estrogen levels and lower fecal estrogen levels in this group. The 16 α -hydroxylation takes place mainly in liver as demonstrated by the very low estriol formation in the cases of Gilbert's syndrome with poor liver uptake of bilirubin and probably also of estrogens [46]. The principal metabolites in bile are 16-oxygenated [55–57] being transported from the small intestine to the colon where they are hydrolyzed, reabsorbed and reconstituted in the mucosal cells [55,58]. The fecal E1, E2 and E3 values are rather similar suggesting that the 16 α -hydroxylated estrogens are more readily reabsorbed. In the present study a significant ($p \leq 0.01$) negative correlation was found between plasma E1 and fecal E3, and plasma E1 + E2 and fecal E3 ($p \leq 0.05$), and fecal total estrogens and plasma E1 ($p \leq 0.05$) (data not shown). Several possible mechanisms have been suggested. All the biliary estrogens are conjugated [59] and have to be hydrolyzed before absorption. An increase of β -glucuronidase activity in feces associated with a high fat and low fiber diet [29] leads to increased reabsorption. The other mechanism proposed to explain this result, is that the estrogens are eliminated by the fecal route because fiber binds estrogens [9,33,51] and fiber speeds up intestinal transit [60]. Each of these mechanisms could account for the results obtained.

In a large number of studies a high androgen level has been found to be a risk factor for BC [20,61–65]. We found no differences between the groups in regard to plasma androgens (Table 3). In a previous study using identical hormone analyses in American postmenopausal women [27], we found significantly higher androstenedione and testosterone values in subjects with BC compared to vegetarians, but the omnivores had values in between. Our present results are

in accordance with some studies that showed no difference between dietary factors and serum androgens [30,66,67]. Androgens do not participate to any great extent in the enterohepatic circulation which could explain these results. The only significant result that we obtained was a negative correlation between the protein intake and plasma androstenedione (Table 6). The altered androgen levels in women with high BC risk may, therefore, not be due to dietary factors but to changes in endogenous hormone production caused by gene polymorphisms changing androgen production in the adrenals or by ovarian disease such as polycystic ovaries.

Catechol estrogens and 2-OHE1/4-OHE1 ratio were higher in the high fat/low fiber group. This ratio is in accordance with a study in Finnish women [43] who showed a high ratio in omnivores and BC women compared to vegetarians. The results are also in accordance with another previous study showing much higher catechol estrogen values and higher 2-OHE1/4-OHE1 ratio in Finnish premenopausal women consuming western diet compared to recent oriental immigrants to Hawaii [68] who had a very low fat diet. Because we observed the high levels of 2-OHE2 and 16 α -OHE1 and no difference of the 2-OHE1/16 α -OHE1 ratio in the high fat/low fiber group compared to the low fat/high fiber group, our results are not in concordance with those of Mutti et al. [69] who suggested that in women a high level of 2-OHE1 and a high 2-OHE1/16 α -OHE1 ratio prevents BC, because a high fat/low fiber diet is associated with an increased BC risk.

The lower 16 α -OHE1 in the low fat/high fiber group is in agreement with a study showing that a low fat diet in women decreases urinary 16 α -OHE1 [21]. However, our results are not in agreement with the observation of an increase in catechol estrogens when women are on a low-fat diet [21].

16 α -OHE1 binds covalently to primary amino groups in proteins and specifically to the estrogen receptors, and its effect is, therefore, prolonged [23]. The 2-hydroxylated estrogens have been shown to have a low estrogenic activity and this forms the basis for the hypothesis that a high 2-OHE1/16 α -OHE1 ratio means diminished BC risk. However, the activation of the estrogen receptor by 4- and 2-hydroxyestradiol is qualitatively and quantitatively comparable to that of estradiol [70]. In the present study both the 2-hydroxylated, particularly 2-OHE2, and 16 α -OHE1 were higher but the 4-OHE1 not in the high fat/low fiber group compared to the low fat/high fiber group. At the time of this study we were not able to measure 4-OHE2 and 2-MeOE2. Low values for the 2-hydroxylated estrogens in the low fat/high fiber group were not reflected in the values for 4-OHE1. This is the reason for the high 2-OHE1/4-OHE1 ratio in the high fat/low fiber group compared to the other group.

The catechol estrogens are easily auto-oxidized to semiquinones and further to quinones and these metabolites are both electrophiles capable of covalently binding to nucleophilic groups on DNA [16]. The quinones are initiators of breast and other cancers and it seems that catechol O-methyltransferase (COMT) inhibits this process by methylating the catechol estrogens [71, 72]. Both 16 α -OHE1 and the catechol estrogens are carcinogenic.

The protection against formation of DNA adducts by COMT is interesting because 2-hydroxylation and methylation of the 2-hydroxy group seem to be a prerequisite for the stimulation of production of SHBG by estradiol [73] at least as judged from *in vitro* studies. A higher SHBG is also protective with regard to BC, because this reduces the concentration of free estradiol and particularly that of free testosterone. In the present study SHBG was significantly lower in the low fat/high fiber group compared to the high fat/low fiber group which is difficult to explain. In one study we found that plasma free fatty acids are positively associated with SHBG [74]. We did not measure free fatty acids, but it is likely that they are high in the high fat/low fiber group, which could explain the high SHBG. However, the

mechanism by which free fatty acids could influence SHBG values is not known, but it could also be a methodological problem.

Because 2-OHE1/16 α -OHE1 ratio is not significantly different between the groups, this result does not fit with the hypothesis that a higher ratio is associated with lower BC risk [69]. The finding that urinary E1, E2 & E3, the catechol estrogens, 16 α -OHE1 and plasma estrogen levels are higher in the high fat/low fiber group leads to the conclusion that a high fat/low fiber diet causes a general increase of all estrogens (Table 5). Because both the 16 α -hydroxylated and the catechol estrogens are carcinogenic both of the groups of estrogens probably play a role in increasing BC risk. Because of the demonstrated carcinogenicity of the 4-hydroxylated estrogens [75,76], it is important to note that there was no difference for the values of 4-OHE1 between the groups and that Finnish women have 100% higher excretion of 4-OHE1 in urine compared to oriental women [68] with a low BC risk. Because 2-OHE1 is high during a high fat/low fiber diet and 4-OHE1 does not change and is low in a population with a low BC risk the 2-OHE1/4OHE1 ratio may be a good biomarker of BC risk.

We observed a significant difference in carbohydrate intake between the groups. A high carbohydrate intake is seen as necessary to maintain caloric intake in a low fat diet. A high carbohydrate diet seems to reduce sex hormone levels and to reduce BC risk [77,78]. However, one study did not find an association between carbohydrate intake and overall BC risk in pre- and postmenopausal women [79].

In conclusion, the high fat/low fiber diet reflects the Western diet compared to low fat/high fiber diet that is closer to the Asian diet. The results show that the Western diet is associated with high hormonal values and more specifically with the estrogens that might increase BC risk. Both catechol estrogen and 16 α -hydroxylated estrogen levels are significantly higher but no difference for the 2-OHE1/16 α -OHE1 ratio between the groups could be seen. The significantly higher 2-OHE1/4-OHE1 ratio in the high fat/low fiber group compared to the low fat/high fiber group that was seen previously in Finnish premenopausal women with BC [43], suggest that in premenopausal women this ratio is a better predictor of BC risk compared to the 2-OHE1/16 α -OHE1 ratio. Our results also suggest that dietary fat has a greater influence on estrogen metabolism than does dietary fiber.

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Table 1

Anthropometric characteristics of the groups

Variables	high fat & low fiber group (n = 22)	low fat & high fiber group (n = 22)	p values
Age (yrs)	25 ± 3	26 ± 3	0.46*
Weight (kg)	57.1 ± 7.5	54.8 ± 6.6	0.28*
BMI (kg/m ²)	20.75 ± 1.94	20.46 ± 1.87	0.62*

Means ± SD.

Significant $p \leq 0.050$.

* p values obtained using log values.

Independent sample t-test.

BMI = body mass index.

Table 2

Dietary intake of the groups

Variables	high fat & low fiber group (n = 22)	low fat & high fiber group (n = 22)	p values
Total calorie intake (Kcal)	1813 ± 327	1599 ± 345	0.041
Fat intake (g/d)	84 ± 19	44 ± 19	0.001
Carbohydrate intake (g/d)	182 ± 36	250 ± 56	0.001
Protein intake (g/d)	65 ± 14	56 ± 15	0.080
SFA (g/d)	32 ± 10	12 ± 8	0.001
MUFA (g/d)	28 ± 6	13 ± 7	0.001
PUFA (g/d)	14 ± 5	11 ± 6	0.164
Fat/Fiber ratio (g/d)	7.8 ± 2.9	1.4 ± 0.6	0.001
Fiber intake (g/d)	11 ± 3	31 ± 9	0.001

Means ± SD.

Significant $p \leq 0.050$.

Independent sample t-test.

SFA = saturated fatty acids; MUFA = mono-unsaturated fatty acids; PUFA = poly-unsaturated fatty acids.

Table 3
Plasma and fecal sex hormone levels of the groups

Variables	high fat & low fiber group (n = 22)	low fat & high fiber group (n = 22)	p values
Estrone (pmol/l)	306 ± 60	264 ± 84	0.069
Estradiol (pmol/l)	249 ± 80	221 ± 84	0.148*
E1 + E2 (pmol/l)	556 ± 126	486 ± 159	0.067*
E1 / E2 ratio	1.27 ± 0.22	1.24 ± 0.30	0.618*
Androstenedione (nmol/l)	5.06 ± 1.20	4.94 ± 1.30	0.752
Testosterone (nmol/l)	1.55 ± 0.60	1.40 ± 0.44	0.375*
Free testosterone (pmol/l)	15.66 ± 7.19	15.64 ± 6.32	0.856*
SHBG (nmol/l)	54.47 ± 15.96	44.03 ± 18.47	0.051
DHEAS (µmol/l)	6.78 ± 3.13	6.96 ± 4.31	0.570*
F-E1 (nmol/24h)	2.30 ± 2.10	3.03 ± 2.72	0.226*
F-E2 (nmol/24h)	1.38 ± 1.32	1.77 ± 1.75	0.162*
F-E3 (nmol/24h)	2.15 ± 1.95	3.10 ± 2.96	0.177*
Total F-Estrogens (nmol/24h)	5.84 ± 5.23	7.90 ± 7.03	0.123*

Means ± SD.

Significant $p \leq 0.050$.

* p values obtained using log variables values.

Independent sample t-test.

E1 = estrone; E2 = estradiol; E3 = estriol; SHBG = sex hormone binding globulin, DHEAS = Dehydroepiandrosterone-sulfate; F = Fecal measures; Total F-estrogens = F-E1 +F-E2 +F-E3.

Table 4
Urinary estrogen metabolite levels of the groups

Variables	high fat & low fiber group (n = 22)	low fat & high fiber group (n = 22)	p values
E1 (nmol/24h)	22.70 ± 7.49	20.64 ± 8.11	0.266*
E2 (nmol/24h)	13.78 ± 6.87	10.85 ± 5.06	0.123*
E3 (nmol/24h)	26.74 ± 11.48	19.42 ± 10.38	0.025*
E3-3G (nmol/24h)	41.50 ± 17.98	27.34 ± 14.00	0.002*
2-OHE1(nmol/24h)	47.54 ± 19.45	37.16 ± 19.13	0.161
4-OHE1 (nmol/24h)	8.15 ± 2.02	8.75 ± 3.12	0.861*
2-OHE2 (nmol/24h)	12.61 ± 4.99	7.44 ± 3.93	0.004
16 α -OHE1 (nmol/24h)	8.61 ± 4.47	5.58 ± 3.64	0.055
2-OHE1/4-OHE1 (nmol/24h)	5.87 ± 2.37	4.17 ± 1.78	0.045*
2-OHE1/16 α -OHE1 (nmol/24h)	8.36 ± 8.68	10.16 ± 7.63	0.709*
2-OHE1/E1 (nmol/24h)	2.08 ± 1.12	1.69 ± 0.57	0.478*
4-OHE1/E1 (nmol/24h)	0.36 ± 0.17	0.45 ± 0.20	0.202*
16 α -OHE1/E1 (nmol/24h)	0.37 ± 0.22	0.30 ± 0.24	0.295*
E3/16 α -OHE (nmol/24h)1	4.78 ± 4.53	5.09 ± 3.17	0.663*
2-OHE (2-OHE1+ 2-OHE2; nmol/24h)	60.16 ± 23.63	44.60 ± 22.00	0.078
16 α -OHE (16 α -OHE1+E3; nmol/24h)	38.09 ± 13.34	27.07 ± 12.76	0.032
2-OHE/16-OHE (nmol/24h)	1.74 ± 0.85	2.14 ± 1.45	0.993*
2-MeO-E1 (nmol/24h)	7.90 ± 2.91	7.26 ± 3.25	0.590
15 α -OHE1 (nmol/24h)	0.86 ± 0.59	0.70 ± 0.39	0.386
16 β -OHE1 (nmol/24h)	3.02 ± 1.43	1.98 ± 1.90	0.023*
16-ketoE2 (nmol/24h)	5.76 ± 2.43	3.96 ± 2.58	0.067
16-epiE3 (nmol/24h)	6.40 ± 3.62	3.89 ± 2.18	0.033*
17-epiE3 (nmol/24h)	2.27 ± 1.53	1.48 ± 0.81	0.123*
Total urinary estrogens (nmol/24h)	174 ± 47	132 ± 38	0.013

Means ± SD.

Significant $p \leq 0.050$.

* p values obtained using log variables values.

Independent sample t-test.

E1 = estrone; E2 = estradiol; E3 = estriol; Total estrogens = E1+ E2 +E3; E3-3G = estriol-3-glucuronide; OHE1 = hydroxyestrone; OHE2 = hydroxyestradiol; OHE = hydroxylated estrogens, all urinary estrogens = total estrogens + all estrogens metabolites (except E3-3G).

Table 5
Hormonal differences between groups with total kcal intake as covariable

Variables	high fat & low fiber group (n = 22)	low fat & high fiber group (n = 22)	p values
<u>In plasma:</u>			
Estrone (pmol/l)	294 ± 20	240 ± 19	0.052
Estradiol (pmol/l)	234 ± 15	200 ± 14	0.078 *
E1+E2 (pmol/l)	529 ± 34	440 ± 31	0.046 *
E1/E2 ratio	1.25 ± 0.06	1.22 ± 0.06	0.587 *
Testosterone (nmol/l)	1.81 ± 0.16	1.53 ± 0.15	0.289 *
SHBG (nmol/l)	62.11 ± 5.29	47.20 ± 4.89	0.045
<u>In urine:</u>			
E1 (nmol/24h)	23.56 ± 2.47	21.68 ± 2.28	0.562 *
E2 (nmol/24h)	16.04 ± 1.68	11.96 ± 1.55	0.068 *
E3 (nmol/24h)	28.6 ± 3.2	22.12 ± 2.96	0.127 *
E3-3G (nmol/24h)	41.28 ± 4.32	29.97 ± 3.99	0.024 *
2-OHE1 (nmol/24h)	45.18 ± 5.66	38.24 ± 5.22	0.457
4-OHE1 (nmol/24h)	8.15 ± 0.81	8.66 ± 0.75	0.745 *
2-OHE2 (nmol/24h)	11.61 ± 1.21	8.25 ± 1.12	0.054
16 α -OHE1 (nmol/24h)	9.05 ± 1.19	5.36 ± 1.10	0.024
2-OHE1/4-OHE1 (nmol/24h)	5.47 ± 0.58	4.45 ± 0.53	0.271 *
2-OHE (2-OHE1+ 2-OHE2; nmol/24h)	56.79 ± 6.59	46.50 ± 6.09	0.317
16 α -OHE (16 α -OHE1+E3; nmol/24h)	37.65 ± 3.97	27.48 ± 3.67	0.072
16 β -OHE1 (nmol/24h)	3.01 ± 0.52	2.03 ± 0.48	0.024 *
16-ketoE2 (nmol/24h)	5.81 ± 0.76	4.04 ± 0.70	0.061 *
16-epiE3 (nmol/24h)	6.01 ± 0.85	4.30 ± 0.79	0.133
Total urinary estrogens (nmol/24h)	167 ± 12	138 ± 10	0.099

Means ± SE.

Significant $p \leq 0.050$.

* p values obtained using log variables values.

GLM multivariate with total Kcal as covariable.

Table 6
Pearson correlation between dietary components (g/d) and plasma, fecal or urinary sex hormones in premenopausal women

Variables	Total Kcal	Total Fat	Total CHO	Total Proteins	Total SFA	Total MUFA	Total PUFA	Fat/Fiber Ratio	Total Fibers
<u>Plasma:</u>									
E1	0.283	0.409**	-0.096	0.511	0.448**	0.462**	0.271	0.331**	-0.052
E2	0.302*	0.347*	0.027	0.099	0.367*	0.365*	0.150	0.220	0.000
E1+E2	0.313*	0.417**	-0.051	0.102	0.450**	0.456**	0.236	0.298*	-0.041
E1/E2 ratio	-0.061	0.067	-0.196	-0.031	0.093	0.116	0.165	0.130	-0.100
Androstenedione	-0.139	0.016	-0.180	-0.315*	0.073	0.072	-0.189	0.111	-0.108
Testosterone	0.017	0.074	-0.120	-0.132	0.018	0.088	-0.200	0.157	-0.178
Free Testosterone	-0.052	-0.062	-0.052	-0.126	-0.092	-0.052	-0.288	0.068	-0.085
SHBG	0.040	0.169	-0.233	0.006	0.100	0.174	0.074	0.218	-0.254
DHEAS	0.058	0.112	-0.077	-0.130	0.146	0.211	0.039	0.120	-0.055
<u>Feces:</u>									
E1	0.207	0.005	0.324*	0.151	-0.061	-0.008	-0.052	-0.139	0.228
E2	0.173	-0.021	0.321*	0.079	-0.089	-0.016	0.015	-0.193	0.226
E3	0.035	-0.075	0.157	0.007	-0.134	-0.068	-0.008	-0.205	0.087
Total estrogens	0.123	-0.070	0.293	0.072	-0.139	-0.071	-0.047	-0.207	0.204
<u>Urine:</u>									
E1	0.237	0.256	0.050	0.052	0.233	0.259	-0.008	0.236	-0.086
E2	0.294	0.314*	-0.063	0.261	0.341*	0.324*	-0.084	0.346*	-0.230
E3	0.142	0.266	-0.294	0.267	0.336*	0.273	-0.055	0.364*	-0.470**
E3-3G	0.167	0.382*	-0.407**	0.196	0.511**	0.431**	-0.007	0.445*	-0.548**
2-OHE1	0.336	0.401*	0.028	0.159	0.312	0.412*	0.145	0.161	-0.031
4-OHE1	0.057	0.033	0.117	0.021	-0.079	-0.034	0.024	-0.120	0.231
2-OHE2	0.561**	0.591**	0.039	0.420*	0.588**	0.597**	0.066	0.389*	-0.161
16 α -OHE1	-0.038	0.237	-0.374*	-0.146	0.366	0.113	0.007	0.267	-0.397*
2-OHE1/4-OHE1	0.406*	0.478**	0.015	0.146	0.424*	0.498**	0.164	0.336	-0.171
2-OHE1/16 α -OHE1	0.279	0.094	0.332	0.215	-0.054	0.039	0.027	-0.047	0.262
2-OHE1/E1	0.185	0.197	0.037	0.110	0.084	0.153	0.093	0.052	0.038
4-OHE1/E1	-0.251	-0.321	0.027	-0.038	-0.393*	-0.396*	-0.080	-0.329	0.245
16 α -OHE1/E1	-0.247	0.017	-0.430*	-0.215	0.137	0.060	0.032	0.103	-0.333

Variables	Total Kcal	Total Fat	Total CHO	Total Proteins	Total SFA	Total MUFA	Total PUFA	Fat/Fiber Ratio	Total Fibers
E3/16 α -OHE1	0.339	0.058	0.296	0.554**	0.011	0.031	-0.031	-0.039	0.128
2-OHE	0.399*	0.460*	0.032	0.222	0.385*	0.470*	0.135	0.217	-0.060
16-OHE	0.224	0.370*	-0.294	0.308	0.514**	0.439*	0.005	0.301	-0.414*
2-OHE/16 α -OHE	-0.062	-0.134	0.191	-0.143	-0.226	-0.160	-0.131	-0.194	0.355
2-MeO-E1	0.253	0.243	0.134	0.093	0.181	0.247	0.070	0.043	0.067
15 α -OHE1	-0.017	0.062	-0.169	-0.028	0.068	0.072	0.125	0.070	-0.168
16 β -OHE1	0.088	0.395*	-0.432*	0.027	0.514**	0.460*	0.025	0.358	-0.438*
16-ketoE2	0.102	0.302	-0.266	-0.027	0.389*	0.372*	0.081	0.306	-0.344
16-epiE3	0.185	0.385*	-0.329	0.151	0.507**	0.503**	0.116	0.324	-0.356
17-epiE3	0.250	0.374*	-0.151	0.184	0.325	0.373*	0.194	0.132	-0.153
Total urinary estrogens	0.469*	0.580**	-0.088	0.323	0.599**	0.634**	0.101	0.351	-0.254

* p \leq 0.05** p \leq 0.01.

CHO = carbohydrates; SFA = saturated fatty acids; MUFA = mono-unsaturated fatty acids; PUFA = poly-unsaturated fatty acids.