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1 **Estrogen Biosynthesis in Breast Adipose Tissue during Menstrual Cycle in Women with**
2 **and without Breast Cancer**

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Abbreviations: BMI, body mass index; 17 β -HSD, 17 β -hydroxysteroid dehydrogenase; E₁, estrone; E₂, estradiol; E₂-FAE, E₂ fatty acyl ester; FEI, free estradiol index; IQR, interquartile range; IS, internal standard; LC-MS/MS, liquid chromatography-tandem mass spectrometry; STS, steroid sulfatase; WHR, waist to hip ratio.

21 **ABSTRACT**

22 Circulating estrogens fluctuate during the menstrual cycle but it is not known whether this
23 fluctuation is related to local hormone levels in adipose tissue. We analyzed estrogen
24 concentrations and gene expression of estrogen-regulating enzymes in breast subcutaneous
25 adipose tissue in premenopausal women with (n=11) and without (n=17) estrogen receptor-
26 positive breast cancer. Estrone (E₁) was the predominant estrogen in premenopausal breast
27 adipose tissue, and E₁ and mRNA expression of *CYP19A1* in adipose tissue correlated
28 positively with BMI. Adipose tissue estradiol (E₂) concentrations fluctuated during the
29 menstrual cycle, similarly to the serum concentrations. In women with breast cancer median
30 adipose tissue E₁ (1519 vs. 3244, $P<0.05$) and E₂ (404 vs. 889 pmol/kg, $P<0.05$) levels were
31 lower in the follicular than in the luteal phase whereas in control women no significant
32 differences were observed. In the follicular phase, mRNA expressions of *HSD17B1* (median
33 0.06; interquartile range 0.05-0.07 vs. 0.17; 0.03-0.2, $P=0.010$) and *CYP19A1* (0.08; 0.07-
34 0.14 vs. 0.22; 0.09-0.54, $P=0.025$) were lower in women with breast cancer than in controls.
35 In conclusion, the changes in adipose tissue E₁ and E₂ concentrations and the estrogen-
36 regulating *CYP19A1* and *HSD17B1* during the menstrual cycle may be related to
37 dysfunctional local estrogen metabolism in women with breast cancer.

38 Words: 200

39 **Key words:** adipose tissue, aromatase, breast cancer, estrogen, menstrual cycle

40

41 Main text: 1964

42 **Introduction**

43 In premenopausal women, estrogens are produced both in the ovaries and through peripheral
44 conversion of adrenal precursor androgens [1, 2]. Increased circulating estrogen and androgen
45 concentrations have been linked to premenopausal breast cancer [3], but the breast cancer
46 risk, unlike in postmenopausal women, may be inversely associated with adiposity [4, 5].

47 In the breast, the mammary gland and the ductal system are abundantly surrounded by
48 adipose tissue (AT). Therefore, AT might provide a local source of estrogen. In AT,
49 aromatase converts androstenedione and testosterone to estrone (E₁) and estradiol (E₂),
50 respectively [6, 7]. Estrone may also be formed from E₁ sulfate by steroid sulfatase (STS) and
51 be further converted to E₂ by 17 β -hydroxysteroid dehydrogenases (17 β -HSDs) [8, 9].

52 Estradiol fatty acyl esters (E₂-FAE) may be released by hormone-sensitive lipase (LIPE) [10].

53 Although estrogen metabolism has been studied to some extent in postmenopausal AT [11-
54 13], little is known about the menstrual cycle -related estrogen concentrations in the breast
55 AT of premenopausal women [14]. Yet, there is evidence that the circulating estrogen
56 concentrations during the follicular phase of the menstrual cycle may be associated with an
57 increased breast cancer risk [15, 16]. On the other hand, breast tumor proliferation is
58 increased in the luteal phase [17]. Thus, exploring the regulation of estrogen synthesis in AT
59 during the menstrual cycle becomes of interest.

60 We assessed E₁, E₂, and E₂-FAE concentrations in the breast subcutaneous AT according to
61 the phase of the menstrual cycle. We also studied adipose tissue mRNA expressions of
62 aromatase (*CYP19A1*), *STS*, 17 β -HSDs (*HSD17B1*, *HSD17B7* and *HSD17B12*), and *LIPE*,
63 which all drive toward the formation of biologically active E₂. Since the fluctuation of
64 estrogens during the menstrual cycle may play a role in premenopausal breast cancer, we
65 compared AT estrogen levels and the estrogen-metabolizing gene expressions in women with

66 and without breast cancer.

67 **Methods**

68 *Subjects and study design*

69 We collected AT samples from premenopausal women operated for estrogen receptor-positive
70 breast cancer (mastectomy; n=11) and women undergoing reduction mammoplasty (control;
71 n=17). Detailed clinical information (Table 1) was gathered at the preoperative visit and from
72 medical records. Women using oral contraceptives were excluded.

73 Blood samples were obtained preoperatively and processed as described [13]. Two
74 subcutaneous AT samples (1 g each) from the breast specimens were immediately snap frozen
75 in liquid nitrogen [13]. The phase of the menstrual cycle was based on serum progesterone
76 and E₂ (luteal phase concentrations >7 nmol/L and 0.37-0.77 nmol/L, respectively).

77 Women with and without breast cancer were comparable in the primary clinical
78 characteristics only waist to hip ratio (WHR) was slightly higher in control women (Table 1).
79 Five women with cancer and six controls (45% vs. 35%, $P=0.34$) had a levonorgestrel-
80 releasing intrauterine device (Mirena®). Due to the very low systemic absorption of
81 levonorgestrel into systemic circulation, these women were included in the study. Serum
82 follicle stimulating hormone (FSH) and sex-hormone binding globulin (SHBG)
83 concentrations were similar in women with or without Mirena (data not shown). Women in
84 the follicular (n=12) or the luteal phase (n=11) of the menstrual cycle did not differ in their
85 primary clinical characteristics.

86 *Ethical approval*

87 The study complies with the Declaration of Helsinki and was approved by the Ethics
88 Committee of Helsinki University Hospital. Informed consent was obtained from all subjects.

89 *Quantification of hormones and SHBG*

90 For determination of E₁, 30 µL of ¹³C₃-E₁ (2.5 nM; IsoSciences) was added as an internal
91 standard (IS) to a 250 mL aliquot of serum or 200 mg of AT homogenized in 1 mL of distilled
92 water. The AT samples were then processed as described in [18]. E₁ in serum and purified
93 tissue samples was quantified by liquid chromatography-tandem mass spectrometry (LC-
94 MS/MS) as previously described [18]. The limit of quantification (LOQ) of E₁ was 10
95 pmol/L (signal to noise-ratio S/N=10). For tissue E₁, interassay variation of the control
96 samples [18] was 1.8% and 4.9% in three consequent assays.

97 For E₂ and E₂-FAE, serum and AT samples were processed as described [19]. Pooled
98 subcutaneous breast AT [18], and serum with added E₂-17-stearate [20] were used as control.
99 E₂ and hydrolyzed E₂-FAE were analyzed by E₂ LC-MS/MS as described [2] with a LOQ of
100 15 pmol/L (S/N=10). Interassay imprecision of endogenous E₂-FAE in pooled AT was 15%
101 in nine consequent assays. Interassay variation of the serum control samples was 11% and
102 7.1% for E₂, and 6.2% and 18% for E₂-FAE, respectively, in nine or ten consequent assays.

103 For serum progesterone, ¹³C₃-progesterone was added as an IS (IsoSciences) and the samples
104 were extracted with diethylether. Extracts and calibrators (Riedel-deHaën, Buchs,
105 Switzerland) were analyzed on a LC-MS/MS system equipped with an API 3000 triple
106 quadrupole mass spectrometer (PE Sciex, Foster City, CA), an Agilent series 1200 HPLC
107 system with a binary pump (Waldbronn, Germany) and a SunFire C18 column (2.1 x 50 mm;
108 Waters, Milford, MA). The mobile phase was a linear gradient consisting of methanol and 50
109 mM ammonium acetate in water, at a flow rate of 250 µL/min and the gradient was: 0 min,
110 50% methanol; 1.5 min 95% methanol; 5 min 95% methanol; and 5.5-10 min 50% methanol.
111 Progesterone was detected as protonated ion in the positive mode with the following
112 transitions: m/z 315 to m/z 109 and IS, m/z 318 to m/z 112. The data was processed with the

113 Analyst Software (Ver. 1.6, Sciex).

114 Serum SHBG was measured by immunoassay [13].

115 ***Preparation and quantification of mRNA***

116 Total RNA was isolated and purified [19], and 1.0 µg of RNA was reverse transcribed into

117 cDNA. Real-time PCR was performed as described [19]. Data were normalized to the

118 geometric mean of two reference genes, importin 8 (*IPO8*) and lysine-specific demethylase

119 2B (*KDM2B*).

120 ***Statistical analysis***

121 Data are expressed as median (range) or median (IQR) unless otherwise stated. The statistical

122 tests were done with SPSS Statistics software, version 22.0. Normality was assessed with the

123 Shapiro-Wilk test. Between-group differences were evaluated with the Student's t test, or for

124 nonparametric variables, with Mann-Whitney U test. For multiple comparisons, analysis of

125 variance with the LSD post hoc test (parametric variables) or the Kruskal-Wallis test (non-

126 parametric variables) were used. For comparisons between the cancer and control groups we

127 used analysis of with adjustment for WHR. For pairwise comparisons, paired samples t test

128 was used (parametric) and Wilcoxon signed ranks test (nonparametric). Correlation was

129 assessed using Spearman's correlation. The level of significance was $P < 0.05$.

130 **Results**

131 Estrone was the predominant estrogen in premenopausal AT, the concentration being

132 approximately three times higher than that of E₂ (Figure 1A, Table 2). Adipose tissue

133 concentrations of E₁ ($P < 0.0001$), E₂ ($P < 0.0001$), and E₂-FAE ($P < 0.0001$) were higher than

134 the corresponding serum levels (Figure 1), and AT E₁ and E₂ correlated positively with their

135 corresponding serum concentrations ($r=0.77$, $P=0.001$ and $r=0.73$, $P<0.0001$, respectively).

136 Adipose tissue E_1 , but not E_2 or E_2 -FAE, correlated positively with mRNA expressions of
137 *HSD17B12* ($r=0.42$, $P<0.05$), *CYP19A1* ($r=0.57$, $P<0.01$), *STS* ($r=0.47$, $P<0.05$) and *LIPE*
138 ($r=0.44$, $P<0.05$). Furthermore, AT E_1 ($r=0.52$, $P=0.008$) and expression of *CYP19A1*
139 ($r=0.60$, $P<0.001$) correlated positively with BMI.

140 During the menstrual cycle, the E_2 levels in AT followed the changes detected in serum, while
141 E_1 and E_2 -FAE levels did not change significantly (Table 2). Serum progesterone did not
142 correlate with AT hormone levels (data not shown). Stratification of the AT hormone
143 concentrations according to the cancer status demonstrated that E_1 and E_2 concentrations were
144 lower in the follicular than the luteal phase of the menstrual cycle in women with breast
145 cancer but not in the control women (Figure 1B-C). In the luteal phase, the E_2 level tended to
146 be higher in women with cancer than in control women, but the difference did not reach
147 statistical significance ($P=0.06$ when adjusted for WHR).

148 In the follicular phase, the expression of *HSD17B1* (median 0.06; interquartile range 0.05-
149 0.07 vs. 0.17; 0.03-0.2, cancer vs. control, $P=0.010$) and *CYP19A1* mRNA (0.08; 0.07-0.14
150 vs. 0.22; 0.09-0.54, cancer vs. control, $P=0.025$) were lower in women with breast cancer
151 than in control women. There were no statistically significant differences in the expressions of
152 *STS*, *LIPE*, *HSD17B7* or *HSD17B12*. Serum FSH correlated negatively with the expression of
153 *CYP19A1* ($r=-0.65$, $P=0.03$), and with AT E_1 ($r=-0.80$, $P=0.01$), and E_2 ($r=-0.81$, $P=0.005$)
154 in women with cancer.

155 Discussion

156 Although ovarian follicles are the principal site of estrogen biosynthesis in premenopausal
157 women, our data indicate that AT synthesizes a considerable amount of estrogens, and AT

158 estrogen levels change significantly during the menstrual cycle. We found that E₁ levels in
159 premenopausal breast subcutaneous AT were about eight times higher than in serum. Adipose
160 tissue E₁ concentration correlated positively with BMI and mRNA expressions of aromatase
161 (*CYP19A1*), *HSD17B12* and *STS* genes, all related to E₁ bioavailability. This supports active
162 local synthesis of E₁, an important precursor for E₂, in premenopausal AT.

163 In theory, there are two possible explanations for the novel finding of fluctuating E₂
164 concentrations in AT reflecting the menstrual cycle serum E₂ levels. First, we cannot rule out
165 the possibility that circulating estrogens would directly influence their AT concentrations.
166 However, no active transport mechanism transferring estrogen from serum to AT against a
167 concentration gradient, has been reported so far. Second, it is theoretically possible that AT
168 estrogen is regulated by mechanisms resembling those in the ovaries. The synthesis of E₂ in
169 the ovaries is controlled by pituitary gonadotropins, as well as cytokines and growth factors
170 [21]. FSH induces estrogen biosynthesis by regulating the transcription of *CYP19A1* and
171 *HSD17B1* in ovarian granulosa cells [22, 23], and the mRNA expression of these genes is
172 decreased during the luteinization process [24, 25]. Furthermore, FSH receptors have been
173 found in the human AT [26]. Data on the regulation of estrogen concentrations in AT during
174 menstrual cycle, however, is scarce: in one previous study, AT E₂ level as determined with
175 immunological methods correlated with the time since last menses [14]. While our current
176 findings indicate that estrogen levels may be regulated in AT during the menstrual cycle, our
177 data do not give direct evidence that these fluctuations could be gonadotropin-controlled.

178 In women with breast cancer, follicular phase AT E₁ and E₂ levels were lower than luteal
179 phase levels, whereas there was no significant difference in control women. Also, in cancer
180 patients, the follicular phase gene expressions of the two estrogen regulating enzymes,
181 *CYP19A1* and *HSD17B1* were lower than in controls. These findings may imply that estrogen
182 synthesis in the tumor-bearing breast AT is dysregulated. There is evidence that adipocytes

183 near the tumor go through a phenotypic change into cancer-associated adipocytes [27], which
184 could possibly affect local estrogen production, at least in part explaining our findings.
185 Contrary to our data on postmenopausal women [13] the overall E₂ or E₂-FAE levels in
186 women with and without breast cancer did not differ. This emphasizes the differences
187 between premenopausal and postmenopausal breast cancer, and the need to take into account
188 hormonal fluctuations when studying premenopausal women.

189 Our study has limitations. We acknowledge the relatively small number of women in our
190 study, which may affect the power in the comparisons between the subgroups. Also, the
191 limited amount of samples did not allow us to analyze protein levels or activities of the
192 steroidogenic enzymes, or the concentrations of other possible regulators of hormone
193 synthesis, such as luteinizing hormone or AT progesterone levels. Although our findings are
194 preliminary in nature, to our knowledge this is the first study to use LC-MS/MS to show
195 menstrual cycle-dependent fluctuation in AT estrogen levels, and furthermore, comparisons
196 between premenopausal women with and without breast cancer.

197 In conclusion, E₁, E₂, and the E₂ fatty acyl ester concentrations were higher in the breast
198 subcutaneous AT than in serum. E₂ concentrations fluctuated during the menstrual cycle, not
199 only in serum, but also in AT. The changes in E₁ and E₂ concentrations and the relative
200 mRNA expression levels of estrogen-regulating *CYP19A1* and *HSD17B1* in AT during the
201 menstrual cycle may be related to dysfunctional local estrogen metabolism in women with
202 breast cancer.

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214

215 **Figure legends**

216

217 **Fig. 1. A.** Serum and adipose tissue estrone (E_1), estradiol (E_2), and E_2 -fatty acyl ester (FAE)
218 concentrations (pmol/l in serum and pmol/kg in adipose tissue) in all premenopausal women
219 ($n=28$). The hormone concentrations in women with ($n=11$) and without ($n=17$) breast cancer
220 were comparable, thus, thus data from all women were combined. The data are expressed as
221 median and interquartile range. $*P < 0.0001$, adipose tissue compared with the respective
222 serum concentrations (paired samples t test and the Wilcoxon signed ranks test). **B.** Adipose
223 tissue estrone (E_1) and **C.** estradiol (E_2), concentrations (pmol/kg) in premenopausal women
224 with and without breast cancer stratified according to the phase of menstrual cycle. Six
225 women in both groups were in the follicular phase of the cycle and six control women and
226 five women with cancer in the luteal phase of the cycle. The data are expressed as median and
227 interquartile range. $*P < 0.05$, student's t test for parametric and Mann-Whitney U test for
228 non-parametric variables

229

230

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305

306

307 Table 1. Clinical characteristics.

	Breast cancer	Control	
	N=11	N=17	P-
			value
Age¹, years	45 (27-49)	40 (21-50)	0.18
Body mass index¹, kg/m²	24 (22-34)	28 (19-33)	0.15
Waist to hip ratio¹	0.82 (0.69- 0.91)	0.88 (0.76- 1.00)	0.006
Serum follicle stimulating hormone¹, IU/L	3.8 (0.9-12.7)	4.3 (1.7-10.5)	0.69
Sex hormone binding globulin¹, nmol/L	60.0 (14.8-92)	59.9 (18.9-122)	0.92
Age at menarche¹, years	13 (11-15)	13 (11-16)	0.23
Age at first labour¹, years	29 (20-34)	29 (17-36)	0.75
Number of labours¹	2 (0-2)	2 (0-3)	0.41
History of hormonal contraception¹, years	0 (0-14)	0 (0-15)	0.17
Ductal breast cancer histology²	10 (91)		
Estrogen receptor positive breast cancer²	11 (100)		
Progesterone receptor positive breast cancer²	10 (91)		

308 The data are expressed as median (range)¹ or n (%)².

309

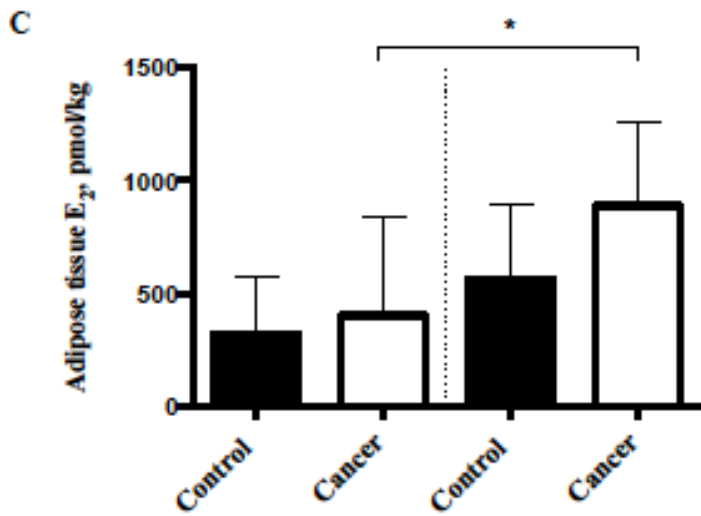
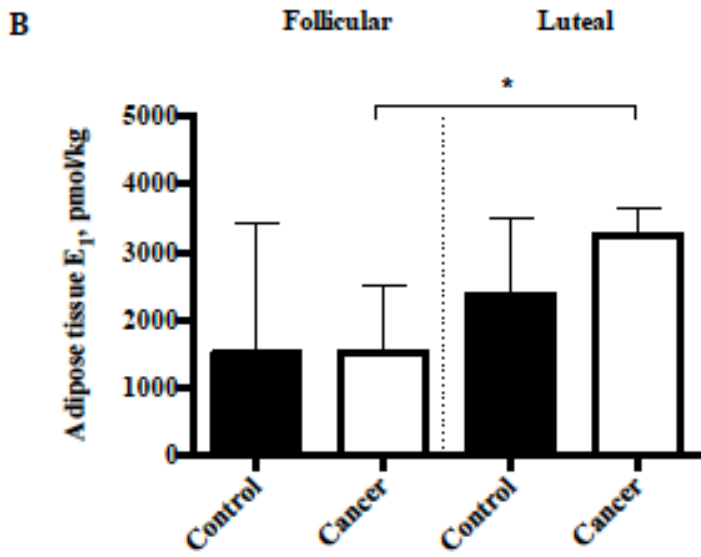
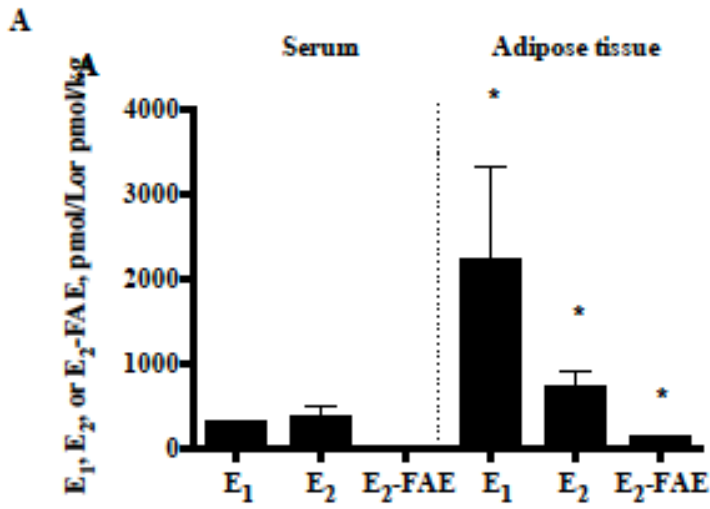
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Table 2. Serum and adipose tissue E₁, E₂, and E₂-FAE levels in premenopausal women according to the phase of menstrual cycle.

	Phase of Menstrual cycle			<i>P</i> -value
	Follicular N=12	Periovulatory N=5	Luteal N=11	
Adipose tissue				
E ₁	1519 (1119-3055)	2213 (1904-3574)	2616 (1992-3592)	0.14
E ₂	326 (204-700) ^{**,*1}	975 (880-1539) ^{**,*2}	767 (495-916) ^{*1,*2}	0.003
E ₂ -FAE	91 (60-128)	133 (100-250)	115 (101-179)	0.088
Serum				
E ₁	200 (104-290)	315 (304-417)	271 (161-336)	0.10
E ₂	182 (104-336) ^{#,*}	575 (506-820) ^{#,**}	399 (282-457) ^{*,**}	<0.0001
E ₂ -FAE	8 (5-11) ^{**}	23 (13-177) ^{*,*}	9 (5-17) [*]	0.024

The data are expressed as median (interquartile range), pmol/L in serum and pmol/kg in adipose tissue. **P*<0.05, ***P*<0.01, ****P*<0.001, #*P*<0.0001

312 Figure
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