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Citation for published version:

Laforest, S, Pelletier, M, Denver, N, Poirier, B, Nguyen, S, Walker, BR, Durocher, F, Homer, NZM, Diorio, C, Andrew, R & Tchernof, A 2019, 'Estrogens and glucocorticoids in mammary adipose tissue: Relationships with body mass index and breast cancer features', *The Journal of Clinical Endocrinology & Metabolism* (JCEM), vol. 105, no. 4, pp. e1504-e1516. https://doi.org/10.1210/clinem/dgz268

Digital Object Identifier (DOI):

10.1210/clinem/dgz268

Link:

Link to publication record in Edinburgh Research Explorer

Document Version: Peer reviewed version

Published In: The Journal of Clinical Endocrinology & Metabolism (JCEM)

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Title: Estrogens and glucocorticoids in mammary adipose tissue: Relationships with body
 mass index and breast cancer features

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- 24 Short Title: Steroid hormones in breast adipose tissue
- 25 *Keywords*: estradiol, estrone, cortisol, cortisone, breast cancer, adiposity

26 *Word count*: 5275

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30 Disclosure summary: AT is the recipient of research grant support from Johnson &

31 Johnson Medical Companies, Medtronic and Pzifer for studies unrelated to this publication.

32 The authors have nothing to disclose.

33 This work was supported by an inter-center grant from the CMDO to AT, CD and FD. SL

34 is the recipient of PhD scholarships from Fonds de recherche du Québec-santé (FRQS)

35 and the Canadian Institutes of Health Research (CIHR). SL's internship in RA's lab was

36 made possible thanks to the support of the Institute of Nutrition and Functional Foods and

37 the scholarship from the International internship program of the *Fonds de recherche Nature*

38 et Technologies du Québec (FRQNT) as well as a Michael Smith Foreign Study

39 Supplement travel scholarship. CD holds an Investigator Awards (Senior) from the FRQS.

40 RA is funded by the Wellcome Trust. BRW is a Wellcome Trust Senior Investigator.

42 ABSTRACT

43 **Context:** Adipose tissue is an important site for extragonadal steroid hormone biosynthesis 44 through the expression and activity of P450 aromatase, 11β -HSD1 and 17β -HSDs. The 45 contribution of steroid hormones produced by adjacent adipose tissue for the progression 46 and survival of breast tumors is unknown.

47 Objective: To quantify estrogens (estradiol, estrone) and glucocorticoids (cortisol,
48 cortisone) in breast adipose tissue from both healthy and diseased women and their
49 relationships with adiposity indices and breast cancer prognostic markers.

50 **Design and setting:** Breast adipose tissue was collected at time of surgery.

51 Patients: Pre- and post-menopausal women undergoing partial mastectomy for treatment
52 of breast cancer (n=17) or reduction mammoplasty (n=6) were studied.

53 Interventions: Estrogen and glucocorticoid relative amounts were determined by liquid
 54 chromatography-tandem mass spectrometry.

55 **Results:** The targeted steroids were reliably detected and quantified in mammary adipose 56 tissues. Women with ER+/PR+ tumor had higher estradiol levels than women with ER-57 /PR- tumor (P < 0.05). Ratio of estradiol-to-estrone was higher in lean women compared 58 to women with a BMI ≥ 25 kg/m² (P < 0.05). Mixed-model analyses showed that estradiol, 59 cortisone and cortisol were negatively associated with tumor size (P < 0.05). Relationships 60 between glucocorticoids and tumor size remained significant after adjustment for BMI. The cortisol-to-cortisone ratio was negatively associated with tumor stage (P < 0.05) 61 62 independently of BMI.

63 Conclusions: We reliably quantified estrogens and glucocorticoids in breast adipose tissue
64 from healthy women and women suffering from breast cancer. Our findings suggest that
65 smaller breast tumors are associated with higher levels of estradiol and cortisol in adipose
66 tissue.

67 **PRÉCIS**

- 68 Estrogens and glucocorticoids were measured by LC-MS/MS in breast fat tissue from
- 69 women with or without breast cancer. Tumor size was negatively associated with estradiol
- 70 and glucocorticoid levels.

72 INTRODUCTION

73 Being overweight or obese is a well-known risk factor for postmenopausal breast cancer 74 ^{1,2}. Obesity is also linked to a poorer prognosis in women with breast cancer regardless of their menopausal status ^{1,3,4}. Women with obesity have more aggressive tumors, higher 75 76 mortality rates, incidence of metastases and increased risk of recurrence ^{1,5-8}. Central 77 obesity, as measured by waist circumference (WC), is an emerging risk factor for both preand postmenopausal breast cancer ⁹. Furthermore, the efficacy of chemotherapy, 78 79 radiotherapy, surgery and endocrine therapy is reduced in women with obesity and possibly more so with increased visceral fat accumulation ¹⁰⁻¹⁴. The mechanisms underlying higher 80 81 risk and reduced treatment efficacy are not fully understood. Altered secretion of 82 adipokines, growth factors and steroids by dysfunctional mammary adipose tissue may contribute to a pro-inflammatory, growth-promoting microenvironment for cancer cells ¹⁵. 83 84 Recent evidence from human studies has shown that local breast adipose tissue does 85 present an altered biological profile, as described above, concomitant with body mass index (BMI) increases ¹⁶⁻²¹. Reports from Iyengar and collaborators have showed that mean 86 87 adipocyte cell size from breast adipose tissue was positively associated with BMI, 88 increased aromatase expression and inflammatory markers such as crown-like structures and was also related to menopausal status ^{19,20}. 89

Expression and activity of several steroidogenic enzymes present in adipose tissue have
been linked to increased adiposity ²². For example, higher rates of androgen-to-estrogen
conversion through aromatization in adipose tissue ²² have been proposed as a mechanism
for the obesity-related increase in breast cancer risk. Findings from our group suggest that
known estrogenic 17β-HSD (type 1, 7 and 12)-mediated conversion of estrone (E1) to

95 estradiol (E2) is five times higher in differentiated adipocytes than in preadipocytes 23 . 96 Increased mean adipocyte size is associated with higher expression level and activity of 97 11 β -HSD type 1 which locally converts cortisone to active cortisol through oxoreductase 98 activity 24,25 .

99 Despite their well-known anti-inflammatory effects, glucocorticoids (GC) could contribute 100 to breast cancer initiation, progression and survival via the activation of the glucocorticoid 101 receptor (GR) or by increasing aromatase expression via the GC response element (GRE) on exon I.4 of the CYP19A1 gene ²⁶. Moreover, 11β-HSD1 expression increases with 102 estrogen receptor β (ER β) activation ²⁷. In a rodent model of breast cancer, increased GC 103 104 levels nurtured the transition from DCIS (ductal carcinoma in situ) to IDC (invasive ductal 105 carcinoma) while administration of RU-486 was able to partially block this effect, i.e. prevent breast cancer progression to IDC ²⁸. Considering the slow turnover of GC in 106 adipose tissue ²⁹, there is biological plausibility for autocrine and paracrine effects of active 107 GC, such as cortisol, in the tumoral microenvironment. 108

Hence, steroid dynamics in breast adipose tissue and cancer appear to be involve more than overexpression and increased activity of aromatase. We currently have very little information about the relative importance of adipose tissue dysfunction markers such as altered steroid conversion to tumor progression and aggressiveness or patient prognosis. Moreover, considering the complexity of potential enzymatic hormone conversion in adipose tissues, direct measurement of active hormone levels and their precursors has become highly relevant.

Liquid-chromatography followed by tandem mass spectrometry (LC-MS/MS) is
recognized as the gold standard to quantify endogenous steroids in plasma, however, tissue

118 steroids are harder to measure than those in plasma due to the complexity of the matrix and 119 the need to homogenize uniformly. Measurements are confounded by the low abundance 120 and the poor ionisation profile of steroids as well as the higher concentrations of lipids in 121 adipose tissue than in plasma, which result in higher susceptibility of so-called matrix 122 effects. We aimed to characterize the relationship of locally-produced cortisol and E2 as 123 well as inactive steroid cortisone and main E2 precursor E1, with adiposity and prognostic 124 markers in a sample of women with or without breast cancer. We hypothesize that 125 independently of menopausal status: i) E2 and cortisol breast adipose tissue levels as well 126 as the ratio E2/E1 and cortisol on cortisone are increased with adiposity; ii) worst clinical 127 breast cancer features, such as tumor stage, size and grade are associated with lower 128 adipose tissue steroid levels; and iii) estrogen relative amounts are increased in women 129 with ER+/PR+ tumors. We also investigated if these relationships were independent of 130 total adiposity i.e. reflecting the micro-environment or a characteristic of the adiposity 131 state.

133 METHODS

134 Study sample and data collection

135 The study protocol was approved by the Research Ethics Committees of Laval University 136 Medical Center (DR-002-136). All patients signed a written, informed consent prior to 137 surgery. Breast adipose tissue was obtained from women undergoing partial or total 138 mastectomy for treatment of breast cancer (n = 17) or reduction mammoplasty (n = 6). 139 Fresh tissue specimens were acquired from residual resected breast tissues that were not 140 required for clinical diagnosis at least 1 cm away from the tumor margins. Information on 141 clinicopathologic and anthropometric factors was collected from in-person interviews, 142 phone-call interviews and medical records.

143 *Laboratory methods*

144 *Cell sizing*

Breast adipocyte size was measured as previously described ^{30,31} in formalin-fixed adipose
tissue. Briefly, 250 breast adipocytes from 10 randomly chosen areas at × 40 magnification

147 using Calopix software (Tribvn) were measured for each subject, in a blinded fashion.

148 Liquid chromatography-tandem mass spectrometry (LC-MS/MS)

149 Standards and solvents

150 E1 and E2 were obtained from Steraloids, Inc (Newport, USA). Cortisone, cortisol, formic

151 acid (FA; \geq 98%), iodomethane (\geq 99%), were from Sigma-Aldrich, Inc. (Dorset, UK). 1-

152 (2,4-dinitro-5-fluorophenyl)-4-methylpiperazine (PPZ) was obtained from TCI chemicals

153 (Chuo-ku, Tokyo, Japan). HPLC grade glass distilled solvents (acetone; ethyl acetate,

154 EtOAc; water) were from Fisher Scientific UK Limited (Leicestershire, UK). Analytical

155	reagent (AR) grade ethanol (EtOH) and HPLC grade glass distilled solvents (acetonitrile;
156	methanol, MeOH) and LCMS grade (acetonitrile; formic acid, FA; water) solvents were
157	from VWR (England, UK).

158 Instrumentation

159 Cortisone, cortisol, E1 and E2 were measured by LC-MS/MS, using a UHPLC Shimadzu 160 Nexera X2 system (UK) coupled to a Sciex QTRAP[®] 6500+ (SCIEX, Warrington, UK) 161 equipped with an electrospray ionization interface (ESI). Mass spectrometry conditions 162 were previously described in conjunction with ion spray voltage (5500 V) and source 163 temperature (500°C) ³².

164 Sai

Sample preparation

Following enrichment of frozen adipose tissue samples (~200 mg) with three internal standards (IS), 9,11,12,12-[${}^{2}H_{4}$]-cortisol (D₄-F 5 ng; Cambridge Isotopes Laboratory, England, UK), 2,3,4-[${}^{13}C_{3}$]-17 β -estradiol and 2,3,4-[${}^{13}C_{3}$]-estrone (${}^{13}C_{3}E2$, ${}^{13}C_{3}E1$ respectively; 5 ng; Sigma-Aldrich, Inc (St. Louis, USA)), analytes were extracted as described below.

Briefly, frozen adipose tissue samples were homogenized (Model Pro 200, ProScientific, Inc, Monroe, CT, USA) in EtOH:EtOAc (1 mL; 1:1) and immediately frozen on dry ice and stored at -80°C overnight. The following morning, samples were thawed on wet ice and sonicated (8 x 15 second bursts with 1-minute gaps; Ultrasonic cleaner, Branson Ultrasonic Inc, Danbury, CT, USA). Samples were subjected to centrifugation (3200 *g*, 45 minutes, 4°C; Heraeus Megafuge 16R, ThermoFisher Scientific, Germany). The supernatant was transferred into a new glass tube and dried down under oxygen-free 177 nitrogen (OFN; 60°C). Samples were resuspended in aqueous MeOH (30% v/v, 5 mL). 178 Solid-phase extraction was carried out after conditioning C18 Sep-Pak columns (12cc, 2g; 179 Waters, Wilmslow, UK; MeOH (2 x 10 mL), followed by H₂O (2 x 10 mL)). The adipose 180 extract was loaded, and the column was washed with H₂O (10 mL) followed by aqueous 181 MeOH (5%, 10 mL). Steroids were eluted with MeOH:CH₃CN (1:1, 10 mL). The eluent 182 was dried down under OFN at 60°C prior to derivatization of estrogens. Generation of 1-183 (2,4-dinitrophenyl)-4,4-dimethylpiperazinium (MPPZ) derivatives of E1 and E2 has already been described ³²⁻³⁴. Derivatization was carried out by incubating (1h, 60°C) with 184 185 (CH₃)₂CO (70 µL), NaHCO₃ (10 µL, 1M, Sigma-Aldrich, Inc (St-Louis, USA)) and PPZ 186 $(10 \,\mu\text{L}, 1 \,\text{mg/mL}, \text{dissolved in (CH_3)}_2\text{CO})$ followed by addition of CH_3I; $(100 \,\mu\text{L})$ and further incubation (2h, 40°C) as previously described ^{33,34}. Samples were dissolved in 187 H₂O:CH₃CN (70 µL; 70:30) and transferred to LC vials. 188

189 Liquid chromatography parameters

Following injection (30 μ L), analytes were separated on an ACE 2 Excel C18-PFP (150 × 2.1 mm, 2 μ m; HiChrom, Reading, England, 40°C) column. The elution process started with mobile phase compositions of 90:10 H₂O with 0.1% FA (solution A) and CH₃CN with 0.1% FA (solution B) which was maintained for 1 minute. This was followed by an 11-minute linear gradient to 50% solution B, which was maintained for 2 minutes, before returning to 10% solution B at 15 minutes and maintained for 3 minutes, all at a constant flow rate of 0.5 mL/min.

197

Linearity and lower limit of quantitation (LLOQ)

Blank samples and aliquots containing estrogens (5, 7.5, 10, 15, 25, 50, 100, 200, 500,
1000 pg/sample), GC (50, 75, 100, 150, 250, 500, 1000, 2000, 5000, 10000 pg/sample) and

200 IS (500 pg) were analysed by LC–MS/MS. Calibration curves were plotted as the peak area 201 ratio (standard/IS) versus amount of analytes (GC or estrogens). Calibration lines of best 202 fit were acceptable if the regression coefficient, r, was > 0.98. Weightings of 1/x were used 203 for all four steroids. LLOQs were 50 pg, 15 pg, 100 pg and 75 pg for E2, E1, cortisol and 204 cortisone, respectively. Values below the confirmed LLOQ were calculated as LLOQ 205 divided by 3 i.e. the lowest acceptable signal-to-noise ratio. The values were then converted 206 to pmol/kg according to the weight of the corresponding adipose tissue sample. This 207 transformation was performed to avoid null values to calculate steroid ratios as described 208 below.

209 Calculated ratios

All steroid amounts were converted into pmol/kg. Those values were then used to calculate
product and substrate ratios, as described below. The ratio of cortisol:cortisone was used
as a marker of 11β-HSD1 enzyme activity. The ratio of E2:E1 was used as a marker of
estrogenic 17β-HSD enzyme activity.

214 Gene expression

215 Tissues were homogenized in Qiazol buffer (Qiagen, Germantown, MD, USA) and total 216 RNA was extracted using the RNeasy mini kit on-column DNase (Qiagen, Hilden, DE) 217 treatment following the manufacturer's instructions. First-strand cDNA synthesis was 218 accomplished using 1 ug of RNA in a reaction containing 200 U of Superscript IV Rnase 219 H-RT (Invitrogen Life Technologies, Burlington, ON, CA). cDNA corresponding to 20 ng 220 of total RNA was used to perform fluorescent-based Realtime PCR quantification using 221 the LightCycler 480 (Roche Diagnostics, Mannheim, DE). Reagent LightCycler 480 222 SYBRGreen I Master (Roche Diagnostics, Indianapolis, IN, USA) was used as described 223 by the manufacturer. The conditions for PCR reactions were: 45 cycles, denaturation at 224 95°C for 10 sec, annealing at 58-60°C for 10 sec, elongation at 72°C for 14 sec and then 225 72°C for 5 sec (reading). Oligoprimer pairs were designed by GeneTool 2.0 software 226 (Biotools Inc, Edmonton, AB, CA) and their specificity was verified by blast in the 227 GenBank database. The synthesis was performed by IDT (Integrated DNA Technology, 228 Coralville, IA, USA) (Table 1). Normalization was performed using the following 229 reference genes ATP synthase O subunit (ATP5O), hypoxanthine guanine phosphoribosyl 230 transferase 1 (HPRT1) and glyceraldehyde-3-phosphate dehydrogenase (GAPDH). 231 Quantitative Real-Time PCR measurements were performed by the CHU de Québec 232 Research Center (CHU) Gene Expression Platform, Quebec, Canada and were compliant 233 with MIQE guidelines.

234 Statistical analyses

235 Differences in breast adipocyte diameter between women with breast cancer (cases) and 236 women without breast cancer (controls) or according to menopausal status were assessed 237 by Student's t-test. Cell size frequency distribution differences between case and control 238 subjects or according to menopausal status were assessed by the Kolmogorov-Smirnov (KS) test. Women were subdivided in categories of BMI [lean ($< 25 \text{ kg/m}^2$) or overweight 239 240 and obese ($\geq 25 \text{ kg/m}^2$)] or according to their estrogen and progesterone receptor (ER/PR) 241 status. Hormone levels or ratios between those subgroups were assessed by Student's t-242 test. Satterthwaite approximation was used when variances were deemed unequal 243 according to a conservative Folded F statistic (P < 0.10). Exact p-values computed using 244 non-parametric Wilcoxon tests showed similar results. Women were subdivided according 245 to their respective tumor size, according to tertiles of the distribution. A mixed-model was 246 performed to evaluate the relationship between hormone levels and tumor size (in tertiles 247 or continuous; as determined by the best AIC fit for the model), grade (categorical) and 248 stage (categorical). A repeated statement was incorporated into the model to account for 249 the non-constant variance among the residuals i.e. specifying a variance component 250 covariance structure in the model. Non-normally distributed variables were log-251 transformed to achieve normality and linearity. Models were further adjusted for BMI (as 252 a continuous variable) to account for total adiposity. Adjustments for menopausal status 253 and current use of hormonal derivatives (as combined indicator variables) were also 254 performed as they are identified as confounders in the breast cancer literature. Spearman 255 correlation coefficients were computed to assess the relationship between steroid relative 256 amount, prognostic factors and relative expression of genes. P-values < 0.05 were 257 considered significant. All statistical analyses were performed with SAS software (SAS 258 Institute, Cary, NC, USA).

260 **RESULTS**

Table 2 shows the characteristics of the study sample. Women were overweight with a median BMI of 25.6 kg/m² and a median age of 55 years. Postmenopausal status was equally balanced across case and control women. Table 3 presents the clinicopathological features of pre-menopausal and post-menopausal breast cancer patients. Most women presented with a unilateral breast lesion. Breast tumors were mainly of ductal histology (82%). Only one woman presented a HER2+ tumor and 13 women had an ER+/PR+ tumor. Half of the women had stage 2 breast cancer as classified by TNM score ³⁵.

Women with or without breast cancer were comparable for primary clinical characteristics, except for mean fat cell size which was 13.3 μ m higher in the control women (*t-test*, P = 0.0072). This difference was also shown by the adipocyte size distribution curve comparison (*KS*, P = 0.0041) (**Figure 1A**). Postmenopausal women had a 9.7 μ m higher mean adipocyte size compared to premenopausal women as well as a right-shifted adipocyte size distribution, however this difference was not statistically significant (**Figure 1C**). **1B**). The distribution remained similar when considering only cancer cases (**Figure 1C**).

275

Quantification of steroids

Table 4 shows adipose tissue steroid relative amounts and calculated ratios. E1 and cortisol
relative amounts were quantified in each of the 23 samples of mammary adipose tissue. E2
level was below the limit of quantification for 2 samples. Cortisone level was below the
limit of quantification for 5 samples, 4 of which were obtained from control women.

280 11β-HSD1 mRNA abundance was positively correlated with the cortisol:cortisone ratio 281 (r = 0.4929; P = 0.0198) and negatively associated with the cortisone relative amount

282	($r = -0.4850$; P = 0.0221). In women with ER+/PR+ tumor, 17 β -HSD12 transcript amount
283	($r = 0.5804$; P = 0.0479) was correlated with E1 relative amount in breast adipose tissue.
284	17 β -HSD12, 17 β -HSD7 or CYP19A1 were not correlated with E2 relative amount.
285	Women with cancer had higher expression level of 17β -HSD12 mRNA compared to
286	controls (P = 0.0231). E2 relative amount was positively associated with ER β mRNA
287	expression ($r = 0.6083$, P = 0.0358), but not ER α mRNA expression in women with
288	ER+/PR+ tumor. In women with breast cancer, abundance of CYP19A1 transcript was
289	positively correlated with 11 β -HSD1 mRNA expression ($r = 0.5353$, P = 0.0326).

290 Adiposity

291 The ratio of E2 to E1 was higher in lean women compared to women with a BMI ≥ 25 kg/m^2 (P = 0.0335) (Figure 2A) (postmenopausal, n=15, P=0.0072; premenopausal, n=8, 292 293 P=NS) in the entire cohort, even if CYP19A1 expression was higher in women with a 294 BMI ≥ 25 kg/m² (p<0.05). This difference remained significant when considering only 295 cancer cases (P = 0.0393, n = 17), or only ER+/PR+ patients (P = 0.0436, n = 13). The 296 difference, although the same magnitude as above, was no longer statistically significant 297 when considering ER+/PR+ patients with invasive carcinoma (stage \geq 1) (P = 0.0900, 298 n = 11). E2 levels were higher in lean women than in overweight and obese women in 299 cancer cases (P = 0.0494) (Figure 2B) (postmenopausal, n=11, P=0.0325; premenopausal, 300 n=6, P=NS). cancer cases (P = 0.0494) (Figure 2B). There was no difference in levels of 301 cortisol and cortisone between lean and overweight/obese women (data not shown).

302 Breast cancer clinical features

Adipose tissue E2 level (log-transformed) was inversely associated with tumor size (categorical variable) (**Figure 3A**) (P = 0.0281, n = 17) (postmenopausal, n=11, P=0.0283;

305 premenopausal, no convergence), but this relationship did not reach statistical significance when tumor size was treated as a continuous variable ($\beta = -0.0199$, P = 0.1089, n = 17). 306 307 Further adjustment for menopausal status and use of hormonal derivatives did not alter the 308 results (P = 0.0488, n = 17) whereas the relationship was no longer significant after 309 adjusting for BMI (P = 0.1569, n = 17) or when considering only patients with invasive 310 carcinoma (stage ≥ 1) (P = 0.1157, n = 14). Including only women with ductal carcinoma 311 generated similar effect size. Although the adjusted models were no longer significant, the non-adjusted model remained significant (data not shown). CYP19A1 mRNA expression 312 313 level was positively correlated with tumor size (r = 0.4875, P < 0.05), whereas expression 314 of ER α mRNA was negatively associated with this variable (r = -0.5435, P = 0.0296). 315 There was no relationship between adipose tissue estrogen levels and tumor stage or tumor 316 grade in our sample.

The ratio of cortisol to cortisone was lower in women with cancer compared to control women (P < 0.0001). This difference was driven by higher cortisone levels and not lower cortisol levels. In fact, cortisone levels were higher in cancer patients than in control patients (P = 0.0055). These differences remained when combining controls with women with ER-/PR- breast cancer, who shared similar characteristics (**Table 5**), and comparing them to women with ER+/PR+ tumor (P = 0.0097 and P = 0.0171, respectively) (data not shown).

No difference was detected between control women and cancer-positive women with respect to cortisol (data not shown). Both log-transformed cortisol and cortisone levels were inversely associated with tumor size [($\beta = -0.01873$, P = 0.0007) and ($\beta = -0.05048$, P < 0.0001)] independent of BMI, menopausal status and current use of hormonal 328 derivatives $[(\beta = -0.02135, P = 0.0027)$ and $(\beta = -0.03636, P < 0.0001)]$ (Figure 3BC) 329 when tumor size was treated as a continuous variable. When we stratified according to 330 menopausal status, the mixed-models remained significant for postmenopausal women 331 (n=11) (Figure 3BC; P=0.0055, P<0.0001) but did not converge for premenopausal 332 women (n=6). Those relationships were still significant when including only patients with 333 invasive tumors [Cortisol: ($\beta = -0.02376$, P = 0.0078); ($\beta_{adj} = -0.02977$, P = 0.0079) and 334 Cortisone ($\beta = -0.03328$, P < 0.0001); ($\beta_{adj} = -0.03650$, P < 0.0001)]. Contrary to 335 estradiol, cortisone and cortisol were not associated with tumor size (all TNM stages 336 included or with only invasive tumors) (P = 0.0904 and P = 0.1894). Tumor stage was 337 negatively associated with cortisol/cortisone ratio independent of BMI ($\beta 2 = -1.8720$, **3D**) 338 $\beta 3 = -1.8316;$ P = 0.0410) (Figure (postmenopausal, n=11, P=0.0660: 339 premenopausal, n=6, P=0.3096). In sensitivity analyses, including only women presenting 340 a ductal carcinoma histology phenotype, the models generate similar results (data not 341 shown). There was no relationship between adipose tissue glucocorticoid levels and tumor 342 grade in our sample.

Higher levels of E2 and E1 were detected in the adipose tissue of women with ER+/PR+ tumor compared to women with ER-/PR- tumor (P = 0.0163 and P = 0.0134, respectively) (**Figure 3E**) (postmenopausal, n=11, P=0.1077 and P=0.4324, respectively; premenopausal, no convergence).

348 **DISCUSSION**

349 To our knowledge, this is the first study to report a LC-MS/MS quantification method 350 combining analysis of cortisone, cortisol, E1 and E2 extracted from breast adipose tissue 351 from both healthy women and women with breast cancer. Several lines of evidence showed 352 that adipose tissue might play an active role in tumor initiation and progression ^{18,36,37}. As 353 such, the notion of an active cross-talk between adipose and tumor cells has been put forward in the literature ^{15,38,39}. Yet, the contribution of adipose tissue to the steroid 354 355 hormone relative amounts and their possible actions remain to be fully elucidated. Most of 356 the steroidogenic pathways have been studied directly in breast tumors or *in vitro*. Reports 357 of endogenous steroid hormones in breast adipose tissue are mostly comprised of analyses 358 of estrogens (E1 and E2) and their corresponding fatty acyl-esters as well as androgen precursors, namely androstenedione and testosterone ⁴⁰⁻⁴³. Of note, a recent study 359 360 characterised more than 20 steroids, including androgens, progestogens and estrogens in breast adipose tissue by gas chromatography-tandem mass spectrometry (GC-MS/MS)⁴⁴. 361 362 A limitation of that study is the lack of data regarding normal breast adipose tissue from 363 healthy controls. In vitro uptake of E1, E2 and cortisol from culture media by female 364 abdominal adipose tissue explants was reported as more than two-fold lower compared to 365 progesterone and testosterone, highlighting the possible contribution of steroid conversion in adipose tissue as a source of estrogens and GC for the tumor ⁴⁵. 366

367

Quantification of steroids

We have been able to quantify these four steroids in most of our breast adipose samples. Interestingly, we found similar ranges in relative amounts for estrogens as those reported by Honma and collaborators in breast cancer tissue, using LC-MS/MS ⁴⁶. These findings 371 suggest that breast adipose tissue is a potent source of sex hormones for the tumor. One of 372 the strengths of our study is the use of stable isotope dilution LC-MS/MS instead of the 373 historically used ELISAs (enzyme-linked immunosorbent assay). ELISAs for steroid 374 measurements have several drawbacks such as nonspecific antibody interactions, 375 inconsistent reproducibility and inadequate sensitivity ⁴⁷. Moreover, they usually require 376 separate assays for each compound of interest, demanding a large quantity of tissue. Using 377 three stable isotope labelled standards in our protocol allowed us to normalize for loss of 378 analytes during the extraction process.

379 Contrary to our hypothesis, our data, reported as pmol/kg of whole adipose tissue, showed 380 a decrease in the ratio of E2/E1 and a decrease of E2 with increasing adiposity, as assessed 381 with the BMI, suggesting little impact of aromatase conversion per mass unit in adipose 382 tissue. This can also be explained by the higher affinity of androstenedione as a substrate for aromatase compared to testosterone, as previously reported ⁴⁸. Marchand et al. reported 383 that higher circulating E2 level is directly associated with increased fat mass ⁴⁹. Simpson 384 385 et al. reported that increased aromatization in obesity was due to a higher number of cells and not to higher conversion activity per adipose tissue mass unit ⁵⁰. Our results are also 386 387 consistent with another study where the authors found a positive correlation between E1/E2ratio from visceral fat and BMI in postmenopausal women ⁵¹. Contrary to Savolainen-388 389 Peltonen, we found that 17β-HSD type 12 mRNA expression is higher in adipose tissue from cancer patients compared to controls ⁴³. We cannot exclude that the lower levels of 390 391 E2 observed in adipose tissue of overweight and obese women are due to increased uptake by the tumor cells as previously proposed by Savolainen-Peltonen⁴³. However, we 392 393 observed this difference in our entire sample and with all women with cancer, including those with ER- status. Effect size were not modified when stratification by ER+/PR+ was
performed.

396 We were unable to find a significant association between BMI and cortisol levels in breast 397 adipose tissue. We acknowledged that this may be due to our limited sample size. However, 398 we found lower and non-quantifiable cortisone levels in breast adipose tissue of our control 399 women (4 out of six). Our group of control women had higher mean adipocyte size than 400 our women with cancer. This is not surprising, as our control women tended to have higher 401 BMI than the women with cancer, although not statistically significant probably because 402 of our small sample. The adjpocyte size difference reported between pre- and postmenopausal women is similar to the findings of Iyengar and collaborators¹⁹. The lower 403 404 levels of cortisone could be partially explained by a higher activation rate of cortisol or a 405 lower inactivation of cortisol to cortisone by 11β-HSD1, because our gene expression 406 results suggest that 11β-HSD1 expression is positively associated with the 407 cortisol:cortisone ratio and negatively associated with the cortisone relative amount These 408 results are consistent with previous findings from our team which showed that 11β-HSD1 409 activity and expression is positively associated with adipocyte size, at least in the abdominal subcutaneous and omental depots ^{24,25}. A previous study with obese subjects 410 411 undergoing bariatric surgery has indeed found lower levels of cortisone in adipose tissue 412 of obese subjects before weight loss and when compared to control with no difference in adipose tissue cortisol levels among those groups ⁵². No difference in tissue relative amount 413 414 of cortisol could be explained by concomitant higher clearance by 5α -reductase, also 415 increased in obesity ⁵³. However, our data showed no decrease of cortisone with higher

BMI which suggests a different catabolism of cortisol and cortisone in women with breastcancer.

418 Some steroids were previously proposed to be increased in adipose tissue during obesity 419 such as cortisol, which could also act as an immune suppressor in breast tissue. As reported 420 by Cirillo et al., many different tumor types produce active cortisol which inhibits tumorspecific CD8+ T proliferation *in vitro*⁵⁴. Infiltration of CD8+ cells was linked to improved 421 cancer-specific survival by Mohammed and collaborators ⁵⁵. However, breast cancer was 422 423 one of the types of cancer not showing any difference between 11β -HSD1/2 expression 424 between cancer and matched normal epithelial tissues, which suggest a paracrine, possibly by adjacent adjose tissue instead of an autocrine effect by cortisol ⁵⁴. In the same order of 425 ideas, immunohistochemistry of 11β-HSD1 showed a presence of the enzyme in 64% of 426 breast tumors and 97% of matched adjacent tissue ⁵⁶ and GR protein level was higher in 427 breast tumor vs normal epithelial tissue ⁵⁷. Increases in GC, particularly cortisol, can induce 428 aromatase expression via the GRE on exon I.4²⁶. On the other hand, cortisol via binding 429 430 to GR is an activator of the estrogen sulfotransferase, which inactivates estrogens by adding a sulfate group and limiting its binding to ER 58 . 431

We did observe higher cortisone levels in adipose tissue from women with cancer with no change in cortisol. As per our initial hypothesis, we did observe a decrease of estradiol, cortisol and cortisone adipose tissue levels according to tumor size and a decrease of cortisol/cortisone ratio with increasing tumor stage. As stated previously, the decrease of estradiol, but not estrone with tumor size could represent an increased uptake by the tumor cells⁴². The higher cortisol/cortisone ratio between cancer patients in comparison to control women and the further decrease of this ratio with tumor stage point to a dual effect of 439 glucocorticoids in breast cancer related to stage of the disease. Increases in 440 cortisol/cortisone ratio at the lower stages could increase estrogen production via activation 441 of aromatase, but at a later stage, decreases in cortisol/cortisone could be explained by a 442 negative feedback loop through increased estrogen production by the tumor and a 443 concomitant lower expression of 11β -HSD1. However, we were unable to demonstrate a 444 relationship between cortisol/cortisone ratio with E2/E1 ratio or E2 and E1 in our sample, 445 contrary to previous results in visceral adipose tissue, but similar to results in subcutaneous tissue ⁵¹. 446

447 Despite our relatively low number of patients in both groups, we found that median E2 448 adipose tissue level was higher in women ER+/PR+ than ER-/PR-. Hennig et al. reported 449 higher adipose tissue E2, androstenedione and androsterone levels in women with ER+ 450 breast cancer compared to ER-, but no difference in E1 or any of the other androgens and progestagens ⁴⁴. Falk et al. measured sex steroids by radioimmunoassay and found a 451 significant difference only for testosterone and a trend for higher E2, E1 and 452 androstenedione adipose tissue levels in ER+/PR+ compared to ER-/PR- samples ⁴⁰. We 453 454 found that E2 relative amount was associated with ER β mRNA, but not that of ER α in 455 women with ER+/PR+ tumor. ER β expression is known to be increased with E2 456 production, not $ER\alpha$, either suggesting a negative feedback or no effect depending on depot 457 origin ⁵⁹. ER β is only expressed in mature adipocytes in adipose tissue whereas ER α is present in both the stroma-vascular fraction and in mature adipocytes 60 . Lower ER α 458 expression has been linked with adipose tissue dysfunction ⁶¹. However, a relationship with 459 460 adipocyte cell size and ER α expression level was not observed in our study.

Limitations of our study comprised the relatively low number of participants which is counterbalanced by the wide range of BMI and prognostic marker values of our cohort. Of note, our total sample number is in the same order of magnitude compared to previous literature on breast adipose tissue steroid measurements ⁴⁰⁻⁴³, representing the difficulty in obtaining those samples for research purposes. Moreover, we have included in our analysis, control samples i.e. samples from healthy women.

467 Data on steroid concentration in breast adipose tissue from healthy control and women with cancer is scarce in the literature. Most publications investigated this relationship in 468 postmenopausal women ^{40,41,43} with the exception of Hennig et al. ⁴⁴, who did include 469 470 premenopausal women (n=6 out of 51) in their analyses. They did not, however, separate 471 according to menopausal status due their low number of premenopausal women. We 472 acknowledge the potential difference in hormone metabolism between pre- and 473 postmenopausal women. Because the main source of estrogens shifts from gonads to 474 peripheral tissues in menopause, we adjusted for menopausal status and age in our analysis. 475 We also performed stratification to further alleviate concerns about the influence of 476 menopausal status, but these analyses should be interpreted with caution. It should be noted 477 that inference in our study is limited to postmenopausal women, as our number of 478 premenopausal women did not allow us to investigate fully their steroid concentration as a 479 separate group. Most of our findings were still significant when including only 480 postmenopausal women. Of note, tissue steroid concentration in our study did not vary as 481 a function of menopause. Our cohort represents normal demographics in breast cancer, i.e. increased prevalence in older, postmenopausal women. Hence, we suggest that these 482 483 findings are relevant in this context.

484 The use of a cross-sectional design does not allow for causal inferences and we 485 acknowledge that there might be reverse causality as higher tumor stage and size may cause 486 changes in steroid metabolism rather than the opposite. Another constraint is that we cannot 487 quantify the distance from the tumor at which the adipose tissue samples were taken. 488 However, as the defined margins were all included in paraffin blocks for 489 clinicopathological assessment directly at the hospital, we can attest that our adipose 490 samples were taken at least 1 cm from the tumor extremities. Hennig et al reported no 491 difference between steroid levels between two sample locations (less than 0.5 cm and more than 5 cm), except for E2⁴⁴. Due to standard of care and acceptable limit margins in 492 493 Quebec for clinicopathological assessment, it was not possible to include adipose tissue 494 less than 0.5 cm away from the tumor in a research project. Therefore, the variation of E2 495 due to the relative distance from the tumor was likely limited.

497 **CONCLUSION**

498 We were able to quantify estrogens and glucocorticoids in breast adipose tissue from both 499 healthy women and women suffering from breast cancer. There is clear indication that 500 steroid hormone metabolism is different among those two subgroups. Moreover, relative 501 amounts of sex steroids in adipose tissue appear to be related to BMI, especially for E2, 502 whereas differences in glucocorticoids levels appear to be more closely related to cancer 503 progression. As such, estradiol levels were lower in women with larger tumors 504 independently of age and menopausal status and glucocorticoid breast adipose tissue levels 505 were negatively associated with tumor size, independently of age, menopausal status and 506 BMI.

508 ACKNOWLEDGMENTS

We would like to acknowledge the collaboration of the study participants. The authors thank Kaoutar Ennour-Idrissi and Lucie Tellier for data collection from the medical records. We are also grateful to Scott Denham from the Mass Spectrometry Core, Edinburgh Clinical Research Facility and to Serge Simard, MSc for technical and scientific expertise.

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739		

741 FIGURE HEADINGS

Figure 1: Size distribution of breast adipocytes. (A) Comparison between control and
case women (*t-test*, P=0.0072; KS, P=0.0041; n=22). (B) Comparison between
premenopausal and postmenopausal women (*t-test*, P=0.0510; *KS*, P=0.0864; n=22). (C)
Comparison between premenopausal and postmenopausal women with cancer (*t-test*,
P=0.0552; *KS*, P=0.0480; n=16).

Figure 2: Adiposity and estrogens in breast adipose tissue. (A) Difference of the estradiol/estrone ratio according to BMI status using 25 kg/m² as a cut-off (P=0.0335, n=23). (B) Difference of estradiol levels in women with cancer according to BMI status using 25 kg/m² as a cut-off (P=0.0494, n=17). Data on graphs are mean. Open circles represent premenopausal women data point.

752 Figure 3: Breast cancer clinical features and breast adipose tissue steroid levels. (A) 753 Mixed-model regression between estradiol relative amount (log-transformed) and tertiles 754 of tumor size adjusted for menopausal status and current intake of hormonal derivatives 755 $(\beta_2 = -0.9785, \beta_3 = -1.1197; P=0.0488)$. (B) Mixed-model regression between cortisol 756 relative amount (log-transformed) and tumor size (continuous) adjusted for BMI, 757 menopausal status and current intake of hormonal derivatives ($\beta = -0.02135$, P=0.0027). 758 (C) Mixed-model regression between cortisone relative amount (log-transformed) and 759 tumor size (continuous) adjusted for BMI, menopausal status and current intake of 760 hormonal derivatives ($\beta = -0.03636$, P < 0.0001). (D) Mixed-model between 761 cortisol/cortisone ratio (log-transformed) and tumor stage adjusted for BMI ($\beta_2 = -1.8720$, 762 β_3 = - 1.8316; P=0.0410). (E) Difference in estradiol and estrone levels in ER-/PR- vs ER+/PR+ women (P=0.0134 and P=0.0454). Data on graphs are mean. n=17. Open circles 763 764 represent premenopausal women data point.

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767 LIST OF ABBREVIATIONS

- 768 11β-HSD1/2: 11β-hydroxysteroid dehydrogenase type 1 and 2
- 769 ${}^{13}C_3E1: 2,3,4-[{}^{13}C_3]$ -estrone
- 770 ${}^{13}C_3E2: 2,3,4-[{}^{13}C_3]-17\beta$ -estradiol
- 771 17β-HSD: 17β-hydroxysteroid dehydrogenase
- 772 AR: Analytical reagent
- 773 BMI: Body mass index
- 774 CD8+ T: Cytotoxic T cell
- 775 CI: Confidence interval
- 776 D₄F: 9,11,12,12 [²H₄]-cortisol
- 777 DCIS: Ductal carcinoma in situ
- E1: Estrone
- E2: Estradiol
- 780 ELISA: Enzyme-linked immunosorbent assay
- 781 ER: Estrogen receptor
- 782 ER β : Estrogen receptor beta
- 783 ESI: Electrospray ionization
- 784 EtOAc: Ethyl acetate
- 785 EtOH: Ethanol
- 786 FA: Formic acid
- 787 GC: Glucocorticoids
- 788 GC-MS/MS: Gas chromatography-tandem mass spectrometry
- 789 GR: Glucocorticoid receptor

- 790 GRE: Glucocorticoid response element
- 791 HER2: Human epidermal growth factor receptor 2
- 792 HPLC: High-performance liquid chromatography
- 793 HRT: Hormonal replacement therapy
- 794 IDC: Invasive ductal carcinoma
- 795 IQR: Interquartile range
- 796 IS: Internal standards
- 797 KS: Kolmogorov-Smirnov
- 798 LC-MS/MS: Liquid chromatography-tandem mass spectrometry
- 799 LLOQ: Lower limit of quantitation
- 800 MeOH: Methanol
- 801 MPPZ: 1-(2,4-dinitrophenyl)-4,4-dimethylpiperazinium
- 802 MRM: Multiple reaction monitoring
- 803 OFN: Oxygen-free nitrogen
- 804 PPZ: 1-(2,4-dinitro-5-fluorophenyl)-4-methylpiperazine
- 805 PR: Progesterone receptor
- 806 UHPLC: Ultra-high-performance liquid chromatography
- 807 WC: Waist circumference
- 808

TABLES

Gene Symbol	Description	GenBank	size (pb)	Primer sequence 5'→3' S/AS
HSD11B1	Homo sapiens hydroxysteroid 11-beta dehydrogenase 1 (HSD11B1), 3 transcripts	NM_005525	85	TGTGCCCTGGAGATCATCAA A/TGATCAGAAGAGTGGTCC AGAGTG
CYP19A1	Homo sapiens cytochrome P450, family 19, subfamily A, polypeptide 1 (CYP19A1), 11 transcripts	NM_000103	123	AAGAGGCAATAATAAAAGGAA ATCCAGAC/CGACAGGCTGG TACCGCATGCTC
HSD17B12	Homo sapien hydroxysteroid (17-beta) dehydrogenase 12 (HSD17B12)	NM_016142	145	CCCACTCTTGACCATCTATT CTG/CTTCCGGATTTTAGCCA GTTTTGTA
HSD17B7	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 7 (HSD17B7)	NM_016371	293	TCCACCAAAAGCCTGAATCT CTC/GGGCTCACTATGTTTCT CAGGC
Er alpha	Homo sapiens estrogen receptor 1 (ESR1), 6 transcripts	NM_000125	293	TGCAAAATCTAACCCCTAAG GAAGTG/CTCCCAGTACCCA CAGTCCATCTC
Er beta	Homo sapiens estrogen receptor 2 (ESR2), 5 transcripts	NM_001437	114	ACGCCGTGACCGATGCTTTG G/TCGCATGCCTGACGTGGG ACA
Atp5o	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (ATP5O)	NM_001697	267	ATTGAAGGTCGCTATGCCAC AG/AACGACTCCTTGGGTATT GCTTAA
Hprt1	Homo sapiens hypoxanthine phosphoribosyltransferase 1 (HPRT1)	NM_000194	157	AGTTCTGTGGCCATCTGCTT AGTAG/AAACAACAATCCGC CCAAAGG
GAPDH	Homo sapiens glyceraldehyde-3- phosphate dehydrogenase (GAPDH)	NM_002046	194	GGCTCTCCAGAACATCATCC CT/ACGCCTGCTTCACCACC TTCTT

Table 1: Sequence primers and gene description

813	Table 2: Clinical characteristics of the women with breast cancer and
814	control women

		Women	
Variables Median (Q1-Q3)	All (n=23)	Controls (n=6)	Cases (n=17)
Age (years)	55.0 (50.1-62.9)	53.7 (44.9-57.5)	55.9 (53.2-63.2)
Menopausal status			
Premenopausal n (%)	8 (35)	2 (33)	6 (35)
Postmenopausal n (%)	15 (65)	4 (67)	11 (65)
Anthropometrics			
BMI (kg/m ²)	25.6 (24.3-28.2)	27.1 (24.3-29.4)	25.4 (24.5-26.8)
WC (cm)	_	_	94 (86-99)
Breast adipocyte mean diameter (µm)	75.5 (67.3-87.5) ^a	87.9 (86.2-89.5)	74.1 (66.1-80.3) ^b
Hormonal derivatives			
Current oral contraceptive use (yes) n (% of premenopausal)	6 (75)	1 (50)	5 (83)
Current HRT use (yes) n (% of postmenopausal)	5 (33)	2 (50)	3 (27)

815 ^an=22, ^bn=16. **Key:** BMI, body mass index; HRT, hormone replacement therapy; WC, waist circumference.

	Premenopausal (n=6)	Postmenopausal (n=11)	All (n=17)
Characteristics n (%)			
Lesion			
unilateral	5 (83.33)	10 (90.91)	15 (88.24)
bilateral	1 (16.67)	1 (9.09)	2 (11.76)
Histology			
Ductal	6 (100.00)	8 (72.73)	14 (82.35)
Others ^a	0 (0.00)	3 (27.27)	3 (17.65)
Receptor status			
ER+/PR+	5 (83.33)	8 (72.73)	13 (76.47)
TNM status			
0	0 (0.00)	3 (27.27)	3 (17.65)
1	2 (33.33)	4 (36.36)	6 (35.29)
2	4 (66.67)	4 (36.36)	8 (47.06)

Table 3: Characteristics of the tumor 816

817 818 ^aincludes lobular, mucinous and metaplastic carcinomas. Key: ER, estrogen receptor; PR, progesterone

receptor.

819 **Table 4: Relative amounts of steroid and associated ratios in adipose**

820 **tissue**

	Women				
Steroids Median (Q1-Q3)	All (n=23)	Controls (n=6)	Cases (n=17)	*p-value	
Cortisone (pmol/kg)	5501 (1829-16748)	326 (298-1829)	7817 (2811-16748)	0.0055	
Cortisol (pmol/kg)	22671 (16040- 33719)	22142 (16040-46171)	22671 (18473- 32688)	0.5697#	
Estrone (pmol/kg)	3744 (2847-9899)	7855 (3062-18998)	3744 (2514-7415)	0.1611	
Estradiol (pmol/kg)	2320 (1537-4160)	2747 (1796-2937)	2306 (1537-4160)	0.4211	
Ratio Cortisol/Cortisone	4.13 (2.47-15.09)	42.28 (15.09-56.09)	3.35 (2.26-5.54)	<0.0001	
Ratio Estradiol/Estrone	0.46 (0.32-0.66)	0.54 (0.15-0.65)	0.44 (0.33-0.63)	0.6059	

821 *Student t-test p-values calculated with log-transformed variables. *Satterthwaite adjusted p-value

823 Table 5: Clinical characteristics of the women with ER-PR- breast cancer 824 and control women

Variables	Control (n=6)	ER-/PR- (n=4)
Median (Q1-Q3)		
Age (years)	53.7 (44.9-57.5)	54.9 (53.6-60.29)
BMI (kg/m ²)	27.1 (24.3-29.4)	26.0 (25.5-26.6)
Premenopausal n (%)	2 (33)	1 (25)
Postmenopausal n (%)	4 (67)	3 (75)

825 **Key:** BMI, body mass index; ER, estrogen receptor; PR, progesterone receptor.







