



Disturbed estrogen and progesterone action in ovarian endometriosis

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ABSTRACT

Endometriosis is a very common disease in pre-menopausal women, where defective metabolism of steroid hormones plays an important role in its development and promotion. In the present study, we have examined the expression of 11 estrogen and progesterone metabolizing enzymes and their corresponding receptors in samples of ovarian endometriomas and control endometrium. Expression analysis revealed significant up-regulation of enzymes involved in estradiol formation (aromatase, sulfatase and all reductive 17 β -hydroxysteroid dehydrogenases) and in progesterone inactivation (AKR1C1 and AKR1C3). Among the estrogen and progesterone receptors, ER α was down-regulated, ER β was up-regulated, and there was no significant difference in expression of progesterone receptors A and B (PRAB). Our data indicate that several enzymes of estrogen and progesterone metabolism are aberrantly expressed in endometriosis, which can lead to increased local levels of mitogenic estradiol and decreased levels of protective progesterone. Changes in estrogen receptor expression suggest that estradiol may also act via non-estrogen receptor-mediated pathways, while expression of progesterone receptors still needs further investigation.

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1. Introduction

Endometriosis is a complex disease that is defined as the presence of endometrial glands and stroma outside the uterine cavity and musculature (Olive and Schwartz, 1993). It is most commonly diagnosed in women of reproductive age (Berkley et al., 2005). There are three different types of endometriosis, namely ovarian endometriosis, peritoneal endometriosis and deep endometriotic nodules, which represent three different entities with different pathogenesis (Nap et al., 2004). Pathophysiological mechanisms have been extensively studied on animal models, but due to species-specific metabolism and reproductive physiology they still cannot replace studies on human samples (Grümmer, 2006).

In general, endometrial implants behave like normal endometrium in their response to hormones, where estradiol (E2) stimulates proliferation and progesterone (P) promotes differentiation (Olive and Schwartz, 1993). In pre-menopausal women the ovaries are the primary sources of E2, but E2 can also be produced in peripheral tissue from inactive precursors (Labrie et

al., 2000). Locally, in endometriotic tissue, E2 can be synthesised in two ways: by the aromatase pathway, which includes conversion of ovarian or adrenal androstenedione to the weakly estrogenic estron (E1); this is further converted to active E2 (Bulun et al., 1999, 2000, 2001) by 17 β -hydroxysteroid dehydrogenases (17 β -HSDs) types 1, 7 and 12. The reverse reaction is catalyzed by the oxidative 17 β -HSDs types 2, 4 and 8 (Luu-The, 2001; Penning, 2003; Midnich et al., 2004; Vikho et al., 2004; Luu-The et al., 2006).

The use of aromatase inhibitors for the treatment of endometriosis has been successful, as this is the rate-limiting step in local E2 production (Bulun et al., 2001; Patwardhan et al., 2008). The other pathway, the so-called sulfatase pathway, includes sulfatase and sulfotransferase. Sulfatase converts estrogen sulfate to E1, which should be activated by the reductive 17 β -HSDs, while sulfotransferase inactivates E1 and E2, forming their sulfates. The production of estrogens from circulating levels of estrogen sulfate is another important alternative or complementary pathway to ovarian steroidogenesis.

P action can also be regulated at the pre-receptor level. Locally P can be converted to the less active 20 α -hydroxyprogesterone (20 α -OHP) and/or to 5 α -reduced metabolites (Pollow et al., 1975). The major human reductive 20 α -HSDs are aldo-keto reductases AKR1C1 and AKR1C3, with the latter also acting as a reductive 17 β -

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Table 1
Patients details: endometriosis group

Sample	Age	Diagnosis	Phase of menstrual cycle
13	33	Endometriosis ovarii	Proliferative
14	24	E. ovarii	Proliferative
15	33	E. ovarii	Late proliferative/early secretory
16	30	E. ovarii	Late proliferative/early secretory
18	44	E. ovarii	Proliferative
19	27	E. ovarii	Early secretory
20	30	E. ovarii	Late proliferative/early secretory
21	34	E. ovarii	Proliferative
22	32	E. ovarii	Secretory
23	27	E. ovarii	Secretory
24	36	E. ovarii	Proliferative
25	33	E. ovarii	Proliferative
26	28	E. ovarii	Late proliferative/early secretory
27	40	E. ovarii	Proliferative
28	29	E. ovarii	Secretory
29	39	E. ovarii	Late proliferative/early secretory
30	31	E. ovarii	^a
31	28	E. ovarii	Secretory
32	37	E. ovarii	^a
36	47	E. ovarii	Proliferative
37	28	E. ovarii	Proliferative
41	37	E. ovarii	Secretory
43	26	E. ovarii	^a
45	25	E. ovarii	Proliferative

^a Not determined.

HSD and potentially also involved in E2 synthesis (Penning et al., 2000).

The abundance and distribution of estrogen receptors (ERs) and progesterone receptors (PRs) will determine whether E2 or P will have a particular effect (Pearce and Jordan, 2004). There are two distinct ERs: ER α and ER β , where ER α has a proliferative effect and ER β acts as a repressor of ER α . Also PRs exist in two isoforms, as products of the same gene; where PRA lacks 164 amino acids from the N-terminal end of PRB (Oehler et al., 2000). PRB acts as transcriptional activator of progesterone target genes, while PRA has dual role which is cell- and promoter-specific. It can act as transcriptional factor or as repressor of PRB function (Vegeto et al., 1993).

In the present study, we have examined the mRNA levels of 11 estrogen- and progesterone-metabolizing enzymes (17 β -HSD types 1, 2, 4, 7, 8 and 12, sulfatase, sulfotransferase, aromatase, AKR1C3 and AKR1C1) and steroid receptors (ER α , ER β and PRAB) in specimens of ovarian endometriosis and control endometrium.

2. Materials and methods

2.1. Tissue samples

A total of 34 samples were collected: 24 of ovarian endometriomas and 10 of control endometrium, from women with *Uterus myomatosus* or *Myoma uteri* (Tables 1 and 2). The study was approved by the National Medical Ethics Committee of the Republic of Slovenia. RNA extraction, cDNA synthesis, real-time PCR and

Table 2
Patients details: control group

Sample	Age	Diagnosis	Phase of menstrual cycle
1	42	U. myomatosus	Non-proliferative
2	47	M. uteri	Late secretory
3	43	U. myomatosus	Early secretory
5	45	U. myomatosus	Proliferative
6	50	U. myomatosus	Early secretory
8	42	M. uteri	Late secretory
9	47	U. myomatosus	Proliferative
10	48	M. uteri	Proliferative
11	43	U. myomatosus	Early secretory
25	60	M. uteri	Atrophic endometrium

statistical analyses were performed as previously reported (Šmuc et al., 2007a,b). Expression analysis of 17 β -HSD types 4, 8 and 12, AKR1C1 and AKR1C3 was performed on the 24 samples, and for another 9 genes on 16 samples of ovarian endometriomas.

2.2. Western blotting

Aliquots from the protein fractions of ovarian endometriomas (isolated from the same samples as for RNA) were separated by SDS-PAGE. The proteins were transferred from the gels to membranes and incubated with 5% non-fat milk to avoid non-specific binding. The membranes were then incubated overnight with primary antibodies: mouse anti-AKR1C3 (1:1000, Sigma-Aldrich, USA), mouse anti-PR-B (1:500, Santa Cruz, USA). The next day, the membranes were incubated with the secondary antibodies (anti-mouse peroxidase-conjugated, dilutions 1:3000 and 1:10,000, Jackson ImmunoResearch Laboratories Inc., USA) for 1 h. Supersignal West Pico Chemiluminescence Substrate (Pierce Biotechnology) was used for the detection of the bound antibodies, according to the manufacturer instructions.

2.3. Immunohistochemistry

Formalin-fixed, paraffin-embedded samples containing ovarian endometriomas were dewaxed in xylene and rehydrated. The sections were incubated in H₂O₂ to block endogenous peroxidase. After antigen retrieval in sodium citrate buffer, sections were incubated with anti-AKR1C3 (1:500, Sigma-Aldrich, USA), anti-ER α (1:20, DakoCytomation, Denmark), or anti-PR-B (1:50, Santa Cruz, USA) antibodies. The peroxidase-antiperoxidase complex with DAB as substrate was used to detect bound antibodies.

3. Results and discussion

Aromatase is a key enzyme in local estrogen production (Noble et al., 1996) and aromatase inhibitors have proven to be beneficial for the treatment of endometriosis in some patients (Amsterdam et al., 2005). We saw that aromatase transcript levels were significantly higher ($p < 0.001$) in the endometriosis group, compared to normal (Fig. 1I) (Šmuc et al., 2007b), confirming previously published data (Noble et al., 1996; Bulun et al., 2005). The aromatase pathway also includes 17 β -HSDs, which convert E1 to the highly active E2, and vice versa. In endometriotic tissue, all of the reductive 17 β -HSDs were up-regulated. Although Zeitoun et al. (1998) have previously reported no differences in steady-state 17 β -HSD type 1 mRNA between eutopic endometrial and endometriotic samples in different phases of the cycle, our published data (Šmuc et al., 2007a,b) show higher levels of type 1 transcript in ovarian endometriomas (Fig. 1A) ($p < 0.001$). Also, Dassen et al. (2007) recently reported up-regulation of 17 β -HSD type 1 in deep infiltrating endometriosis. We have also shown higher levels of 17 β -HSD type 7 in endometriotic tissue compared to normal (Šmuc et al., 2007a,b) (Fig. 1B) ($p < 0.001$). The expression of 17 β -HSD type 12 has not yet been examined either in normal or diseased endometrium; here, we detected 17 β -HSD type 12 mRNA in ovarian endometriosis and in normal endometrium tissue samples, and for the first time, we show that also 17 β -HSD type 12 is up-regulated in the diseased endometrium (Fig. 1C) ($p = 0.0016$).

For oxidative 17 β -HSDs, Zeitoun et al. (1998) have reported an absence of 17 β -HSD type 2 mRNA transcripts in extra-ovarian endometriotic tissue with Northern blotting; however, our more sensitive study saw this transcript in both the control and the endometriosis group, although we did not detect any significant changes in the expression levels between these groups (Šmuc et al., 2007a,b) ($p > 0.05$) (Fig. 1D). In contrast, Matsuzaki et al. (2006) and Dassen et al. (2007) have shown lower expression of 17 β -HSD type 2 in deep infiltrating endometriotic tissue. There have been no reports of decreased levels of 17 β -HSD type 2 in ovarian endometriosis. For 17 β -HSD types 4 and 8, we did not find significant differences in their expression levels (Fig. 1E and F), although Dassen et al. (2007) have reported a significant down-regulation of 17 β -HSD type 4 in deep infiltrating endometriosis. However, at least to our knowledge, we are the

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