

Estrogen receptor α and β in uterine fibroids: a basis for altered estrogen responsiveness

Panagiotis Bakas, M.D.,^a Angelos Liapis, M.D.,^a Spiros Vlahopoulos, Ph.D.,^b Maria Giner, M.D.,^a Stella Logotheti, B.Sc.,^b Georgios Creatsas, M.D.,^a Aggeliki K. Meligova, B.Sc.,^c Michael N. Alexis, Ph.D.,^{b,c} and Vassilis Zoumpourlis, Ph.D.^b

^a2nd Department of Obstetrics and Gynaecology, Aretaieio Hospital, University of Athens; ^bBiomedical Application Unit; and ^cMolecular Endocrinology Programme, Institute of Biological Research and Biotechnology, National Hellenic Research Foundation, Athens, Greece

Objective: To investigate the relative expression and the DNA-binding status of estrogen receptors α and β in fibroids and normal myometrial tissue to explore the molecular basis of altered estrogen responsiveness of leiomyomas.

Design: Biopsy samples from uterine fibroids and adjacent normal myometrial tissue at the follicular phase of the menstrual cycle.

Setting: Aretaieio University Hospital and the National Hellenic Research Foundation, Athens, Greece.

Patient(s): Thirty-five patients who underwent hysterectomy or myomectomy because of myoma symptoms.

Intervention(s): None.

Main Outcome Measure(s): Deoxyribonucleic acid-binding status of estrogen receptors α and β .

Result(s): The level of messenger RNA expression of estrogen receptor α and β and the level of estrogen receptor as a whole are increased on average to a similar extent in leiomyomas compared with normal myometrium. Occasionally, however, estrogen receptor α is disproportionately increased in leiomyomas, and this appears to increase the amount of estrogen receptor α that binds to the estrogen-responsive element of estrogen target genes as homodimer rather than as heterodimer with estrogen receptor β .

Conclusion(s): The estrogen receptor α -to-estrogen receptor β expression ratio rather than the individual expression levels determines the fraction of DNA-binding homodimers of estrogen receptor α and possibly the growth potential of myomas. (Fertil Steril® 2008; ■: ■-■. ©2008 by American Society for Reproductive Medicine.)

Key Words: Uterine myomas, estrogen receptor alpha, estrogen receptor beta, myometrium, fibroids

Uterine leiomyoma (also known as myoma or fibroid) is the most common benign gynecologic tumor and is present in at least 20% of women at reproductive age and in 40% to 50% of women older than 40 years of age (1). It is estimated that <50% of uterine myomas produce symptoms, and they are usually discovered by clinical examination. Although the etiology of uterine fibroids is unknown, their development is considered to be estrogen dependent, because they have the ability to enlarge during pregnancy and to shrink during menopause, ovariectomy, and other hypoestrogenic conditions (2–4). Aromatase P450 is often overexpressed in myomas, causing in situ synthesis of estrogen to increase, and this is believed to contribute to the growth of leiomyomas (5). Recent studies therefore continue to focus on the role of estrogen receptor (ER) in the pathogenesis of leiomyomas (6).

Estrogens act mainly through two ER subtypes, ER α and ER β , which function as ligand-dependent regulators of transcription in a manner that depends on the selection of cognate cofactors provided by the cell and the structure of the ER target gene promoter, as well as that of the ligand (7–9). The two forms of ER bind to DNA enhancer elements (estrogen-responsive elements [EREs]) or to other transcription factors (e.g., Activator Protein 1 [AP1]) in the promoter of ER target genes as a heterodimer, as well as homodimers (10, 11). Estrogen receptor α and ER β exhibit significant homology in the DNA-binding domain and the ligand-binding domain, where receptor dimerization functions are known to reside. However, the ligand-binding specificity and the transcriptional activity of the two forms of ER are known to differ substantially (8, 9). In fact, it has been reported that ER α and ER β have opposite effects on AP1 sites and that they probably have different roles as regards regulation of AP1-dependent genes, including genes involved in the control of cell growth and viability (12). In the mammary gland, ER α and ER β are pivotal for tissue development and terminal differentiation, respectively (13, 14). In line with

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Reprint requests: Vassilis Zoumpourlis, Ph.D., National Hellenic Research Foundation, 48 Vas. Constantinou Ave, 116 35 Athens, Greece (FAX: 30-210-7273677; E-mail: vzub@eie.gr).

this, 17β -E₂ (hereafter referred to as E₂) has been reported to promote the growth of immortalized mammary cells through ER α and inhibit it through ER β (15). In mammary hyperplasia of the usual type, it is the ER α -to-ER β ratio rather than the individual receptor amounts that reportedly is associated with the risk that this lesion will develop into invasive carcinoma (16). Work with ER α -positive breast cancer cells engineered to express increasing amounts of ER β has shown that, although ER β could regulate genes that are not targeted by ER α alone, it increasingly affected genes regulated by ER α as well, although nearly half of ER α target genes were not affected even when the level of ER β was set to be much higher than that of ER α (17). Notably, however, key genes involved in the hormonal control of cell proliferation and apoptosis were among those of ER α target genes that were affected by ER β , implying that the ER α -to-ER β expression ratio could influence tumor cell growth and consequently breast cancer prognosis and treatment as well (17).

Although endothelial and connective tissue cells of leiomyomas and normal myometrium express only ER β , smooth muscle cells express both ER subtypes (18). Whether and how the ER α -to-ER β expression ratio could influence the estrogen-dependent development of leiomyomas is not known presently. Several studies have demonstrated that the level of expression of ER β messenger RNA (mRNA) in both leiomyomas and normal myometrium is lower than that of ER α (19–22). However, there is conflicting evidence as regards the level of ER α mRNA in the leiomyomas compared with the adjacent myometrium, with some studies reporting an increase in the diseased tissue compared with normal (19) and others reporting no change (20, 22). There is also conflicting evidence as regards the relative abundance of the two forms of ER in the fibroids and how this could impact fibroid response to hormonal therapy, because some studies report that the ER α -to-ER β ratio is higher in the diseased tissue compared with normal (22), whereas others report an increase in the ER α -to-ER β ratio after treatment with a GnRH analogue (21). In addition, it has been reported that the levels of ER α and ER β in myometrial cells change during the menstrual cycle and that the patterns of changes are similar (4). The present study aims at estimating mRNA and protein levels as compared with the DNA-binding activities of two ER subtypes in leiomyomas and normal myometrium during the follicular phase of the menstrual cycle.

MATERIALS AND METHODS

Patient Selection and Tissue Collection

Biopsy samples of uterine leiomyomas and adjacent myometrium were taken from 35 women at the follicular phase of the menstrual cycle (between day 5 and day 9). Dating of endometrium was performed with use of Noyes criteria (23). The criterion for inclusion in the study was pending operation for hysterectomy or myomectomy because of subfertility, menorrhagia, or other bothering myoma symptoms. Criteria for exclusion from the study were adenomyo-

sis, malignancy, or hormonal therapy given within 3 months before operation. Selected patients were subjected to a complete preoperative workup to exclude any other known possible cause of their problem. All specimens originated from 2nd Department of Obstetrics and Gynaecology, Aretaieio Hospital, School of Medicine, University of Athens, and were examined by the same histopathologists. The study took place with the permission of the local ethics committee. All patients gave informed consent for participating in the study.

Ribonucleic Acid Quantification

Ribonucleic acid extraction Total RNA was isolated from normal and pathologic (leiomyoma) myometrial samples with use of TRIzol (GIBCO BRL, Grand Island, NY) according to the manufacturer's instructions, and RNA concentration of the samples was measured with use of a U-2000 spectrophotometer (Hitachi, Tokyo, Japan).

Complementary DNA synthesis Messenger RNA transcription into complementary DNA (cDNA) was performed with use of 2 μ g of total RNA and SuperScript ribonuclease (RNase) H–reverse transcriptase (Invitrogen, Carlsbad, CA), according to manufacturer's instructions.

Polymerase chain reaction Estrogen receptor α and ER β mRNA levels were assessed with use of semiquantitative multiplex reverse transcriptase–polymerase chain reaction (RT-PCR) as previously described (24). Estrogen receptor α and ER β cDNA fragments were coamplified with a larger reference cDNA fragment of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (25). The ratio of the relative amounts of the amplified products in myoma samples is taken to reflect the relative amounts of ER α and ER β mRNAs and was compared with the relative amounts in the corresponding myometrium. A 441–base pair (bp) fragment of ER α cDNA was amplified with use of primers (forward: 5'-TGC CAA GGA GAC TCG CTA CTG-3', reverse: 5'-GGG GGC TCA GCA TCC AAC AAG-3') corresponding to bases 896–916 and 1316–1336 of the published sequence (GenBank accession number NM 000125). A 268-bp fragment of ER β cDNA was amplified with use of primers (forward: 5'-CGA TGC TTT GGT TTG GGT GAT-3', reverse: 5'-GCC CTC TTT GCT TTT ACT GTC-3') corresponding to bases 1400–1420 and 1648–1667 of the published sequence (GenBank accession number AB006590). Oligonucleotide primers and Taq polymerase were purchased from Invitrogen. Polymerase chain reaction conditions for ER α were 94°C for 5 minutes, followed by 25 cycles of 94°C for 30 seconds, 60°C for 30 seconds, and 74°C for 30 seconds, and a final step of 74°C for 10 minutes. Polymerase chain reaction conditions for ER β were 95°C for 5 minutes, followed by 30 cycles of 95°C for 30 seconds, 60°C for 30 seconds, and 72°C for 30 seconds, and a final step of 72°C for 10 minutes. The RT-PCR reaction products were analyzed by polyacrylamide gel electrophoresis (PAGE), stained with ethidium bromide, and quantified with use of a Crossfield 5400/Rip/

Laser/CMYK Densitometer (Crossfield/PS Electronic Services, Shefford Beds, UK). Normalized levels of ER α and ER β mRNA were obtained by dividing the corresponding RT-PCR reaction product densities by the respective GAPDH product densities.

Protein Preparation and Analysis

Preparation of cytosol and nuclear extracts Frozen leiomyoma and myometrium samples were homogenized in ice-cold TEM (10 mmol/L tris[hydroxymethyl]aminomethane [Tris] pH 7.4, 1 mmol/L ethylenediaminetetraacetic acid [EDTA], 2 mmol/L Na₂Mo₄, and 10% glycerol) (Sigma, St. Louis, Mo) in a tissue weight-to-buffer volume ratio of 1:4 (grams per milliliter) with use of an Ultra-Turrax T25 homogenizer (Fischer Scientific, Schwerte, Germany). The homogenate was centrifuged initially at 1,000 \times g for 10 minutes, and the resulting supernatant was centrifuged at 10,000 \times g for 60 minutes. The resulting cytosol (supernatant) was used to assay ER by immunoprecipitation-PAGE-immunoblotting and ligand binding. The protein concentration of the cytosol was measured by the Bradford method (26).

For the preparation of the nuclear extracts, fine slices of the leiomyoma and myometrium samples were homogenized in ice-cold TSM (25 mmol/L Tris pH 7.5, 5 mmol/L KCl, 0.5 mmol/L MgCl₂, 0.5 mmol/L dithiothreitol [DTT], 0.5 mmol/L phenylmethylsulfonyl fluoride [PMSF]) with use of a Teflon-glass homogenizer (Thomas, Philadelphia, PA). The nuclear fraction was pelleted, washed thoroughly with isotonic buffer (25 mmol/L Tris pH 7.5, 5 mmol/L KCl, 0.5 mmol/L MgCl₂, 0.5 mmol/L DTT, 1 mmol/L PMSF, and 0.2 mmol/L sucrose), and lysed with TET (25 mmol/L Tris pH 7.5, 1 mmol/L EDTA, 0.1% Triton, 0.5 mmol/L DTT, 0.5 mmol/L PMSF). Nuclear debris was removed by centrifugation at 55,000 \times g for 1 hour at 4°C. The protein concentration of the resulting nuclear extract (supernatant) was measured by the method of Bradford (26).

Immunoprecipitation, PAGE, and immunoblotting Immunoprecipitation was carried out with use of 40 μ g of protein A-Sepharose 6MB (Sigma, St. Louis, MO), mixed with 0.2 μ g of the C-311 antibody against ER α (Santa Cruz Biotechnology, Santa Cruz, CA) in nondenaturing immunoprecipitation buffer (0.05 mol/L Tris pH 7.9, 1 mmol/L DTT, 0.1 mol/L KCl, 0.1% Nonidet P-40, 20% glycerol) (Sigma), and incubated for 2 hours at 4°C under rotary shaking. The resin with the bound antibody was collected by centrifugation (1,500 \times g, 1 minute) and washed three times for 5 minutes with 1 mL of ice-cold immunoprecipitation buffer; 0.5 mg of nuclear extract protein was added, and the mixture was further incubated for 4 hours at 4°C under rotary shaking. The resin was then washed and collected by centrifugation as above. The pellet was resuspended in 2 \times sodium dodecyl sulfate (SDS)-PAGE sample buffer and subjected to SDS-PAGE and immunoblotting analysis, as already described (27). Immunostaining was developed by using ECLplus (Amersham Biosciences, Piscataway, NJ) and quantified by using

a Storm 860 phosphorimager (Molecular Dynamics, Sunnyvale, CA).

Estrogen receptor assessment by ligand binding The assessment of ER in the cytosol of leiomyomas and adjacent myometrium was carried out with use of standard Scatchard plot analysis as previously described (28). In brief, aliquots (150 μ L; 400 μ g protein) of cytosol were incubated overnight at 4°C with increasing concentrations (0.1–8 nmol/L in 50 μ L TEM) of tritiated 17-[2, 4, 6, 7-3H] 17 β -E₂ (95 Ci/mmol; NEN) and with (50 μ L of) a 1,000-fold molar excess of diethylstilbestrol in TEM (Sigma), or with (50 μ L of) a 200-fold molar excess of the ER α -selective ligand propyl pyrazole triol (PPT) (29) in TEM, or with (50 μ L of) TEM alone. After incubation, the samples were mixed with a dextran-coated charcoal pellet (prepared from 50 μ L of 10% Norit A preincubated with 0.1% dextran T70; USB Corp., Cleveland, OH) for 20 minutes at 4°C to remove free tritiated E₂. Norit A-dextran T70 was pelleted by centrifugation at 1,500 \times g for 10 minutes, and 200 μ L of clear supernatant was removed and counted with use of 4.2 mL of Ultima-gold XR scintillation fluid (PerkinElmer, Waltham, MA) and a Wallac 1409 DSA scintillation counter (Wallac/PerkinElmer, Waltham, MA). The levels of total ER and ER α were obtained from the difference in tritiated E₂ binding in the absence and presence of diethylstilbestrol and PPT, respectively, and were expressed as femtomoles (of specifically bound tritiated E₂) per milligram of protein. The protein concentration was measured by the Bradford method (26).

Electrophoretic mobility shift assay Electrophoretic mobility shift assay (EMSA) was performed with use of the ERE of the *Xenopus* vitellogenin A2 gene (vitERE), as reconstituted by annealing of the following oligonucleotides: 5'-TCAAAGTCAGGTCACAGTGACCTGATCAAAGA-3' and 5'-TCTTTGATCAGGTCAGTGTGACCTGACTTTG A-3'. The reconstituted vitERE was end-labeled with γ -³²P-adenosine triphosphate (ATP) with use of T4 polynucleotide kinase, and the labeling reaction products were purified by PAGE with use of an 8% polyacrylamide gel. Protein DNA binding reactions were carried out by mixing 2,000 cpm of γ -³²P-labeled oligonucleotide with 20 μ g of nuclear extract protein in binding buffer (50 mmol/L HEPES pH 8.0, 500 mmol/L NaCl, 0.5 mol/L PMSF, 0.5 mg/mL bovine serum albumin, 1 mmol/L EDTA, 20% glycerol) plus 1 mmol/L DTT and 150 μ g/mL poly(2'-deoxyinosic-2'-deoxycytidilic acid) sodium salt (poly(dI-dC)) (Sigma). The reaction mixture was left at room temperature for 30 minutes, and the samples were subsequently subjected to electrophoresis on a 6% polyacrylamide gel at 150 V for 90 minutes, dried, and visualized by autoradiography. For the supershift assay, the reaction mixture was incubated with antibodies to ER α or ER β (Santa Cruz Biotechnology) for 30 minutes at 4°C.

Statistical Analysis

Statistical analysis was performed with use of MedCalc software version 7.6.0.0 (MedCalc Software, Mariakerke, Belgium). The levels for ER α and ER β mRNA relative to

GAPDH mRNA in leiomyomas and the corresponding normal myometrium were compared with use of the paired samples *t*-test (and confirmed by the Wilcoxon paired samples test, i.e., the nonparametric equivalent of the paired samples *t*-test). For comparing the ER status of leiomyomas and normal myometrium, the nonparametric Wilcoxon paired samples test was used, because the data were not normally distributed, and results were plotted with use of a box-and-whisker graph.

RESULTS

Patient Characteristics

Biopsy samples of uterine leiomyomas and adjacent myometrium during the follicular phase of their cycle from 35 women who underwent myomectomy or total abdominal hysterectomy for uterine fibroids were included in this study. The mean age of the patients was 45.7 ± 2.9 years (95% confidence interval 44.7–46.7 years), the mean body mass index was 28.1 ± 1.3 kg/m² (95% confidence interval 27.7–28.6 kg/m²), and mean parity was 1.9 ± 0.7 (95% confidence interval 1.7–2.2).

Estrogen Receptor Status of Leiomyomas and Normal Myometrium

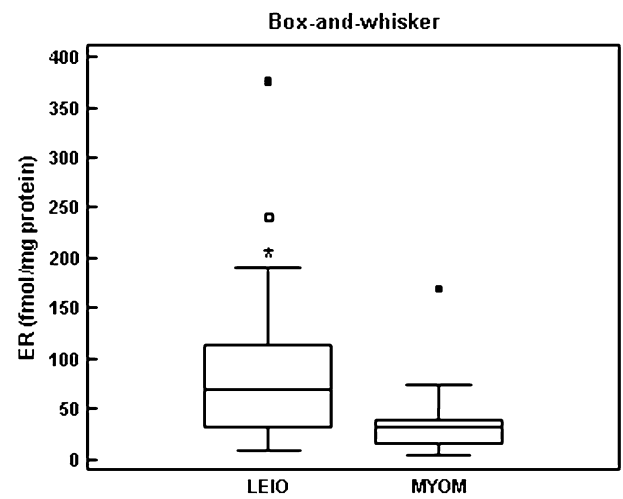
Initially we studied tritiated E₂ binding of the ER in the cytosol of leiomyomas and the adjacent myometrium with use of Scatchard plot analysis. We found that the level of ER in the cytosol of leiomyomas (median level, 67 fmol/mg protein) was significantly higher (Wilcoxon paired samples test; $P < .002$) compared with that of myometrium (median level, 30 fmol/mg protein) (Fig. 1). Then we tried to use PPT, which reportedly exhibits a relative (to E₂) binding affinity for ER α (RBA α) that is much higher than that (RBA β) for ER β (29), to estimate the fraction of E₂ binding of ER that could be attributed to ER α alone, with inconsistent results. The reason for this could be that the ER α -to-ER β binding selectivity of PPT, as determined with use of purified ER α and ER β and a fluorescence polarization assay previously described (27, 30), was found not higher than 100 (RBA α = 50.2 ± 2.4 , RBA β = 1.4 ± 0.3 , RBA α /RBA β = 36; RBA α and RBA β of E₂ are set equal to 100). This caused substantial inhibition of E₂ binding of purified ER β by PPT at concentrations of the ligand capable of inhibiting E₂ binding of purified ER α only partially (data not shown), which rendered the estimation of the fraction of ER α in mixtures with ER β with use of PPT fairly inaccurate.

Estrogen Receptor α and ER β Gene Expression of Leiomyomas and Normal Myometrium

Next we examined whether the higher level of ER in the myoma cytosol was due to an increase in the level of gene expression of ER α , ER β , or both. We therefore determined the levels of mRNA of ER α and ER β in the myomas and the adjacent myometrium using semiquantitative multiplex RT-PCR. To account for differences during the reverse

FIGURE 1

Estrogen receptor levels (median values) in the cytosol of leiomyomas (LEIO) compared with normal myometrium (MYOM) from 35 patients operated on for symptomatic fibroids. Median values of ER levels in leiomyoma and the corresponding myometrium samples, as assessed by Scatchard plot analysis with use of tritiated E₂, were calculated by using the nonparametric Wilcoxon paired samples test, expressed as femtomoles of ER per milligram of cytosol protein and plotted with use of a box-and-whisker graph. The *boxes* contain the values between the upper (75th percentile) and lower (25th percentile) quartiles, the *lines* across the boxes correspond to the medians, and the *whiskers* extend to the highest and lowest values, excluding outliers (*open square*) and extreme values (*filled squares*). *Significantly different compared with myometrium ($P < .05$).

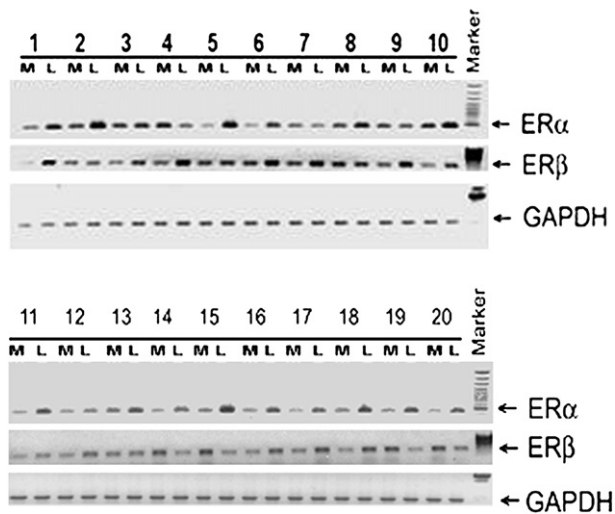


Bakas. Estrogen receptors α and β in fibroids. *Fertil Steril* 2008.

transcription (RT) step, GAPDH cDNA was also amplified and used as a reference standard in all the samples (Fig. 2). Unlike the levels of cytosolic ER we determined by tritiated E₂ binding, the levels of mRNA of ER α and ER β relative to those of GAPDH, deduced as described in Materials and Methods, were distributed quite normally. The mean ratio of mRNA levels of ER α relative to GAPDH was 1.7 ± 1.7 (95% confidence interval 1.1–2.3) in the myometrium and 3.2 ± 2.4 (95% confidence interval 2.4–4.1) in the myomas. The mean level of ER α mRNA in the myomas was found 1.9 times higher (paired samples *t*-test; $P < .0001$) than in the adjacent myometrium. Using a similar approach we found that the mean ratio of mRNA levels of ER β relative to GAPDH was 1.0 ± 0.5 (95% confidence interval 0.8–1.3) in the myometrium and 1.8 ± 1.2 (95% confidence interval 1.4–2.3) in the myomas. The mean level of ER β mRNA in the myomas was found 1.8 times higher (paired samples *t*-test; $P < .0025$) than in the myometrium. Thus, it appears that

FIGURE 2

Estrogen receptor α and ER β mRNA levels in leiomyomas and the corresponding normal myometrium, as compared with the respective GAPDH mRNA levels. Estrogen receptor α , ER β , and GAPDH mRNA levels in the myometrium (M) and the corresponding leiomyoma samples (L) from each individual patient were obtained by semiquantitative multiplex RT-PCR. Typical ethidium bromide-stained gels of the RT-PCR reaction products from the samples of 20 patients are shown.



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the above reported increase in the level of ER in the cytosol of myomas compared with myometrium is due to similar increases in the level of mRNA of ER α and ER β .

Deoxyribonucleic Acid-binding Activity of Leiomyoma and Myometrial ER

To find whether and how up-regulation of myoma ER affects the DNA-binding activity of the receptor we carried out comparative EMSA analysis of the ER of myomas and the respective myometrial samples using the vitERE as bait. We invariably observed that vitERE binding of protein in nuclear extracts from leiomyoma samples was higher as compared with normal myometrium. Then we focused on the samples in which up-regulation of ER α mRNA relative to ER β mRNA was most noticeable (e.g., samples 5 and 10 of Fig. 2). Initially we examined whether the higher level of ER α mRNA expression of these leiomyomas was accompanied by a higher level of the receptor in the nuclear extract. Immunoprecipitation and immunoblotting analysis using ER α -specific antibodies and quantification of ER α by phosphorimaging revealed that leiomyoma levels of ER α from, for example, samples 5 and 10 were 9.3-fold and 6.2-fold higher, respectively, as compared with the respective myome-

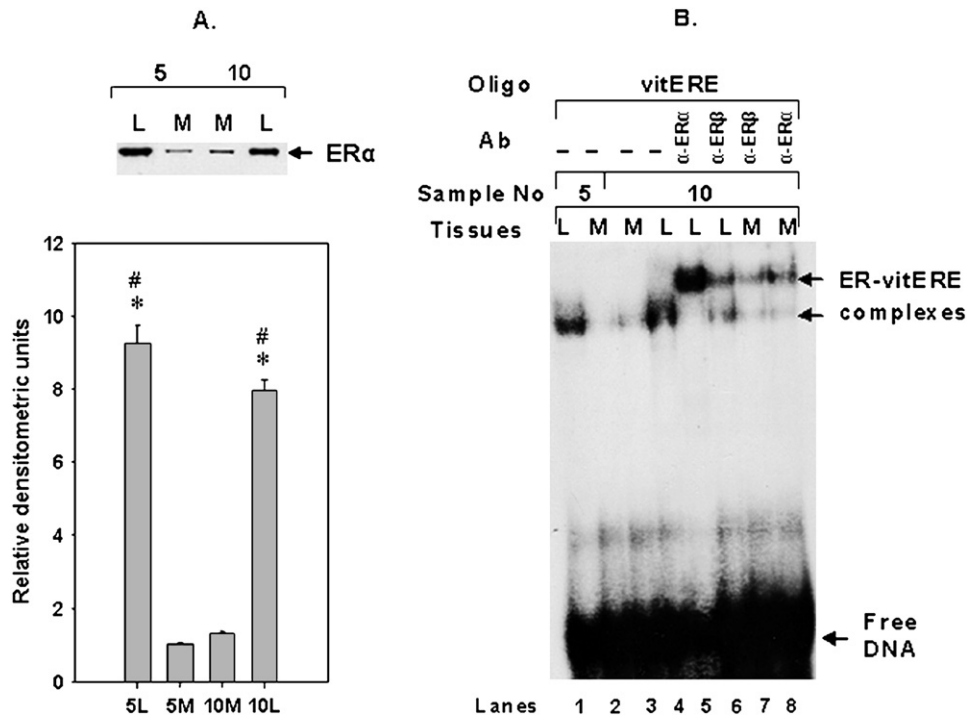
trial levels (Fig. 3A). Next we compared vitERE binding of proteins in the nuclear extracts of leiomyoma and the respective adjacent myometrium. Quantification of the vitERE-protein complexes using phosphorimaging detected 11.2-fold and 7.8-fold higher vitERE binding activity in leiomyoma samples 5 and 10, respectively, compared with the adjacent myometrium (Fig. 3B; lanes 1–4). Supershift analysis of the ERE-binding activity of leiomyoma sample 10 and the respective myometrium using antibodies specific for ER α and ER β followed by phosphorimaging revealed that the participation of ER α in the ER-vitERE complexes exceeded that of ER β by 9.4-fold (Fig. 3B; compare the supershifted bands in lanes 5 and 6), as compared with 2.2-fold in the adjacent myometrium (Fig. 3B; compare the supershifted bands in lanes 7 and 8). Thus, the 6.2-fold increase in the level of ER α in myoma sample 10 compared with the adjacent myometrium (Fig. 3A) is faithfully reflected by a comparable increase in the participation of ER α in ER-vitERE complexes (Fig. 3B; compare the supershifted bands in lanes 5 and 8). Similarly, the increased participation of ER β in the ER-vitERE complexes in myoma sample 10 compared with the adjacent myometrium (Fig. 3B; compare the supershifted bands in lanes 6 and 7) faithfully reflects the mean 1.8-fold increase in the level of ER β mRNA in the myomas compared with the myometrium reported above. Interestingly, Fig. 3B shows, in addition, that ER-vitERE complexes containing ER α alone occurred to a much higher extent with myoma than with myometrial ER α (Fig. 3B; compare the nonsupershifted bands in lanes 6 and 7).

DISCUSSION

Estrogens regulate a variety of physiologic processes in many different tissues and organs. They act predominantly through the two hormone-binding forms of ER, ER α and ER β , both of which function as ligand-dependent regulators of transcription by binding to enhancer elements in the promoter of estrogen target genes in heterodimeric or homodimeric form (9–11). Uterine leiomyomas express ER α and ER β , and their development is known to be estrogen dependent (4–6), but the molecular basis of this dependency is not clearly understood. In the present study we used myomas and adjacent myometrium from 35 women at the follicular phase of their menstrual cycle to examine whether ER α mRNA and protein levels are up-regulated in the myomas compared with the adjacent myometrium and how this up-regulation might impact the ERE-binding activities of ER α and ER β . We found that the level of expression of both ER α and ER β mRNA is by and large higher in leiomyomas compared with normal myometrium (Fig. 2), causing the level of ER to increase accordingly (Fig. 1), and that the abundance of ER α relative to ER β can increase in leiomyomas compared with the myometrium to a level that is high enough to allow leiomyoma ER α to bind to the vitERE as a homodimer, as well as a heterodimer with ER β (Fig. 3). Specifically, the data of Figure 3B suggest that up-regulation of ER α in myoma sample 10 (Fig. 3A) caused the receptor to bind to

FIGURE 3

Protein levels and DNA-binding status of ER α in leiomyomas and the corresponding normal myometrium. (A) Representative immunoprecipitation, PAGE, and immunoblotting analysis of ER α in nuclear extracts from normal myometrium (M) and leiomyoma (L) samples no. 5 (5M and 5L) and no. 10 (10M and 10L), selected for their high ER α -to-ER β mRNA expression ratio (see samples numbered in Fig. 2). (B) vitERE binding of protein in the nuclear extracts from normal myometrium and leiomyoma samples no. 5 and no. 10, as assessed by EMSA (lanes 1 vs. 2 and 3 vs. 4, respectively). Supershifting of vitERE-binding of protein in the nuclear extracts from normal myometrium and leiomyoma sample no. 10, as assessed by EMSA in the presence of antibodies to ER α (a-ER α , lanes 5 and 8) and ER β (a-ER β , lanes 6 and 7). *Significantly different compared with 5M ($P < .05$; t -test). #Significantly different compared with 10M ($P < .05$; t -test). Ab = antibodies.



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the ERE as a homodimer (not supershifted with antibodies to ER β), as well as a heterodimer with ER β , and that this is not the case with ER α in the corresponding normal myometrium. Notably, formation of an ERE-binding homodimer of ER α occurred in spite of the participation of leiomyoma ER β in ER-vitERE complexes being nearly twice as high as that of myometrial ER β , suggesting that the formation of ERE-binding homodimers of ER α depends on the ER α -to-ER β ratio rather than the individual amounts of the two ER subtypes. That ER β is usually up-regulated in the myomas compared with the myometrium is further substantiated by the data of Figure 2. Thus, it appears that the formation of ERE-binding homodimers of ER α is most likely the result of higher up-regulation of ER α compared with ER β rather than the result of down-regulation of ER β during the myometrium-to-myoma transition.

Estrogen receptor α is considered to be a key player in the development of uterine fibroids, because ER α mRNA and protein levels are often up-regulated in fibroids compared

with myometrium (19, 22). In line with this notion, adenovirus-mediated expression of a dominant negative ER α was recently shown to inhibit tumor growth in nude mice (6). In addition, ER α is considered to be a key player during progression of benign mammary proliferative disorders to invasive carcinomas. Recent evidence suggests, however, that a high ER α -to-ER β ratio rather than the level of ER α alone is what characterizes those cases of mammary hyperplasia of the usual type that are likely to progress to breast cancer (16). In fact, there is ample evidence that, although ER α expression is maintained in breast cancer compared with benign tumors or normal mammary tissue, ER β expression is decreased (31). Interestingly, work with HC11 immortalized mammary epithelial cells has revealed that, although E $_2$ promotes cell growth through ER α , at the same time it causes cells to undergo apoptosis through ER β (15). Thus, it appears that the establishment of a high ER α -to-ER β ratio, as the result of down-regulation of the expression of ER β during progression of benign mammary tumors to breast cancer, is somehow associated with enhancement of tumor

growth, and this could reflect stimulation of ER α -dependent cell proliferation, suppression of ER β -dependent cell death, or both. The physiologic significance of the disproportionate up-regulation of the ER α -to-ER β mRNA ratio during uterine fibroid development (e.g., Fig. 2, sample 5), and the relative abundance of ER α over ER β that is likely to result, as regards proliferation and survival of myoma smooth muscle cells, is, however, unexplored.

It has been reported that ER α and ER β can form heterodimers that bind DNA with an affinity similar to that of ER α homodimers and greater than that of ER β homodimers (10); that ER β can regulate gene expression in a ligand-independent manner (17, 32); and that the ERE-dependent transcriptional activity of ER α in the presence of E₂ is negatively modulated by ER β , which thus appears to act as a dominant suppressor of ER α (30, 32). The transcriptional activity of ER α through other transcription factors in the promoter of estrogen target genes also is modulated negatively by ER β . Most important, E₂ activation of cyclin D1 gene expression through ER α is completely inhibited by ER β (33). The opposing action and dominance of ER β over ER α in E₂ activation of both ERE-dependent and AP1-dependent gene expression supports a role for ER β as a dominant negative inhibitor of the proliferative and antiapoptotic effects of E₂ through ER α . In line with this notion, it has been proposed that the modulation of ER α -dependent cyclin D1 expression by ER β could be defective in leiomyomas (34). Our data provide a mechanistic basis as to how this might come about, because they show that in leiomyomas with a high ER α -to-ER β ratio, formation of ER α homodimers is increased substantially, and this may cause the opposing action of ER β on, for example, ER α -mediated activation of cyclin D1 gene expression to decrease accordingly, with potentially pronounced implications as regards tumor growth. In addition, our data indicate that a substantial increase in the fraction of ER α homodimers, as detected with use of vitERE as bait, could be the molecular determinant that differentiates fast-growing myomas from stationary ones. A study is currently in progress to test this hypothesis.

No effective medical treatment is available presently to women who have symptomatic fibroids and want to avoid surgery (35). Because leiomyomas may depend on an ER α -to-ER β ratio higher than normal for growth in the presence of estrogen, agents causing down-regulation of ER α and/or up-regulation of ER β could be used against fibroids. Interestingly, indole-3-carbinol, a constituent of *Brassica* vegetables reportedly endowed with potent anticancer activity in rodent models of carcinogenesis (36), was recently shown to down-regulate ER α expression without altering ER β expression of MCF-7 breast cancer cells and to cause the levels of ERE-bound ER α and ER β to decrease and increase, respectively, and the stimulation of the proliferation of the cells by E₂ to drop as a consequence (37). Thus, agents that can decrease the inherently high ER α -to-ER β ratio of MCF-7 cells may also inhibit the estrogen-dependent proliferation of these cells, apparently by improving promoter occupancy of ER

target genes by ER β relative to ER α . Whether such agents could have an impact on the growth of fibroid smooth muscle cells, which are known to express ER β as well as ER α (18), is presently a matter of conjecture.

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