

# Specific adaptations of estrogen receptor $\alpha$ and $\beta$ transcripts in liver and heart after endurance training in rats

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**Abstract** Estrogens exert their biological roles mainly through estrogen receptors (ER) that function as ligand-activated transcription factors. ER content in a cell is regulated by many factors and is decisive for estrogen action. The purpose of the present study was to investigate the influence of an 8-wk endurance training program on ER expression in the liver, right atrium (RA), and left ventricle (LV) of intact and ovariectomized (Ovx) rats. We measured ER $\alpha$  and ER $\beta$  mRNA content by reverse transcription-polymerase chain reaction (RT-PCR). We found an important increase in ER $\alpha$  mRNA levels in the liver (300%;  $P < 0.01$ ) and in ER $\beta$  mRNA levels in the RA (200%;  $P < 0.05$ ), and a marked decrease in ER $\alpha$  (80%;  $P < 0.01$ ) and ER $\beta$  (40%;  $P < 0.05$ ) transcripts content in the LV of intact rats after endurance training. On the other hand, ER $\alpha$  mRNA levels were depressed by 50% ( $P < 0.01$ ) in the liver, and increased by 60% ( $P < 0.01$ ) in LV of Ovx rats after exercise training. These results first indicate that endurance training is associated with modifications of ER transcripts levels in the liver, LV, and RA of

female rats. More specifically, these effects are tissue and isoform-specific and the direction of the response (increase or decrease) is different in intact and Ovx rats. It is suggested that some of the adaptations to endurance training in liver and heart may be mediated by an ER-dependent mechanism.

**Keywords** Gene expression · Ovariectomy · Right atrium · Left ventricle · Ovarian steroids

## Introduction

It is now well recognized that the effects of estrogens are not limited to the female reproductive system [1, 2]. Almost all organs of the body in both male and female are under estrogenic influence, and especially metabolic tissues such as liver and heart [1]. The biological actions of estrogens occur mainly when they bind to one of the two known estrogen receptor (ER) isoforms—ER $\alpha$  and ER $\beta$ —specifically distributed among tissues [1, 2]. These nuclear receptors are ligand-activated transcription factors and can stimulate or repress the expression of genes encoding proteins with important physiological functions [1, 2]. In addition, ER can be stimulated independently of estrogen-binding, by phosphorylation of the N-terminal ligand-independent activation function (AF-1 domain), present on the ER gene [3]. Activation of the AF-1 domain allows ER to stimulate the transcription of estrogen-regulated genes [3]. Therefore, some estrogenic effects involving an ER-mechanism are mediated in the absence of estrogens [3].

In the liver, the synthesis of angiotensinogen, blood coagulation factors, LDL receptor, and lipoproteins are under the control of an ER $\alpha$ -dependent mechanism [4–7],

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the ER $\beta$  isoform being absent in liver [8]. Estrogens also act in heart through ER $\alpha$ , where estradiol treatment reduces myocardial injury associated with ischemia-reperfusion [9, 10], and through ER $\beta$ , where estrogens exert a protection against left ventricular hypertrophy [11]. Since the capacity of a cell to mediate an estrogenic response is correlated to the intracellular ER content, it is of interest to study the stimuli that influence ER expression [7, 12, 13].

Hepatic ER $\alpha$  content is principally influenced by growth hormone action, but other pituitary and glucocorticoid hormones are involved [14–16]. It is known from the ovariectomized (Ovx) rat model that endogenous estradiol negatively regulates hepatic ER $\alpha$  content [17–19], whereas hyperphysiological doses of 17 $\beta$ -estradiol after exogenous administration or during pregnancy stimulates its synthesis [20–22]. Nevertheless, the changes in circulating estradiol levels associated with the estrous cycle do not affect ER $\alpha$  content in liver, unlike the fluctuations in ER $\alpha$  levels reported in uterus [16, 20, 21]. On the other hand, a positive regulation of ER levels by endogenous estrogens was reported in female rat atria using an Ovx model [23]. However, few studies looked at the regulation of ER content in heart, making it less defined and maybe not only endogenous estrogens are implicated. Taken together, it can be concluded that the regulation of ER expression is tissue-specific and can be influenced by several factors.

In addition to the above-mentioned stimuli, an endurance-trained state has also been reported to influence ER expression [24–26]. However, the information in this regard is scarce and limited to skeletal muscle. Two studies conducted by Lemoine et al. [24, 25] indicated changes in ER $\alpha$  transcripts levels after a 7-wk endurance training program in skeletal muscle. Furthermore, Wiik et al. [26] found higher ER $\alpha$  and ER $\beta$  transcripts levels in vastus lateralis of highly endurance-trained than in moderately active men. Lemoine et al. [24, 25] put forward the concept that an ER-mediated mechanism could contribute to the training adaptations of skeletal muscle. We hypothesized that ER expression is also modulated following endurance training in liver and heart. The rationale for this hypothesis is based on the evidence that both endurance training and estrogen signaling pathway share physiological targets in liver and heart. Examples of this are repression of the expression and the activity of hepatic lipase in liver [27, 28] and improvement of cardiac recovery after ischemia/reperfusion [29, 30]. In a first step, to increase our understanding of the link between ER and exercise training in these tissues, we determined the effects of an 8-wk endurance training program on ER $\alpha$  and ER $\beta$  transcripts expression in liver and 2 heart chambers, left ventricle (LV) and right atrium (RA) of intact female rats.

It is well recognized that Ovx in rodents results in an increased adiposity [31, 32], and induces several metabolic

disturbances including hepatic steatosis, hypercholesterolemia, hyperinsulinemia [31, 33], and decreased insulin sensitivity [34]. Several of these disturbances are prevented by exercise training [34, 35]. Furthermore, exercise training and estrogen therapy augment cardiac and endothelial function in Ovx rats [36–39]. It is not known if ER transcripts are affected after exercise training in Ovx rats. This question is of interest especially that ER mRNA levels cannot rely on endogenous ovarian hormones anymore. Therefore, a second aim of the present study was to determine the changes in ER transcripts expression following endurance training in liver and heart of ovariectomized rats. We have shown that endurance training is associated with tissue- and isoform-specific ER changes in intact female rats and that the direction of the response is modified by the removal of ovaries.

## Materials and methods

### Animal care

Female Sprague–Dawley rats ( $n = 32$ ) weighing 180–200 g (8-wk old) were obtained from Charles River (St-Constant, PQ) and housed individually. The 12:12-h light–dark cycle started at 6:00 AM, and room temperature was maintained at 20–23°C. All animals were given free access to the usual pellet rat chow (12.5% fat; 63.2% carbohydrate; 24.3% protein; kJ, Agribrands Purina Canada, Woodstock, ON) and tap water. They were treated similarly in terms of daily manipulations. The experiments described in this report were conducted according to the guidelines of the Canadian Council on Animal Care after institutional approval.

### Surgery

Two days after their arrival in our laboratory, the rats were randomly assigned to ovariectomy (Ovx) or sham-operation (Sham) groups. Ovx was performed according to the technique described by Robertson et al. [40]. The animals were injected with antibiotics (Tribrissen 24%; 0.125 ml/kg sc) for 3 days, beginning on the day before surgery. For surgery, the rats were anesthetized with a mixture of ketamine–xylazine (61.5–7.6 mg/kg, i.p.).

### Groups

Two days after surgery, Ovx and Sham rats were submitted to training (Tr), or remained sedentary (Sed), which resulted in a total of 4 groups ( $n = 8$ /group): sham-operated (Sham), sham-operated + endurance training (ShamTr), ovariectomized

(Ovx), and ovariectomized + endurance training (OvxTr). Body weight and food intake in g were monitored every other day. All rats were killed 8 wk after surgical manipulation in the Tr or Sed state.

#### Endurance training protocol

Exercise training consisted of continuous running on a motor-driven rodent treadmill (Quinton Instruments, Seattle, WA), 5 times/wk for 8 wk. The rats progressively ran from 15 min/day at 15 m/min, 0% slope, up to 60 min/day at 26 m/min, 4% slope, for the last 4 wk. At the end of this 8-wk period, animals were sacrificed 48 h after the last exercise bout.

#### Tissue sampling

Rats were killed between 08:00 and 11:00 AM. Food was removed from the cage 2–3 h before sacrifice. Immediately after complete anesthesia (pentobarbital sodium; 50 mg/kg i.p.), the abdominal cavity was opened along the median line of the abdomen. Several organs and tissues were removed in the following order: liver, heart, uterus, femur, mesenteric, urogenital, and retroperitoneal fat depots, along with the right triceps surae (soleus, plantaris, and gastrocnemius). The liver median lobe was freeze-clamped and the heart was divided in its 4 chambers. The liver median lobe, the right atrium (RA), and the left ventricle (LV) were processed for ER $\alpha$  and ER $\beta$  mRNA determination and quantification. All tissue samples were weighed (Mettler AE 100), immediately frozen in liquid nitrogen, and stored at  $-78^{\circ}\text{C}$  until further analysis. Finally, the right femur wet weight was obtained following a short-boiling period in a 10% KOH solution in order to remove the surrounding tissue.

#### Isolation of RNA and RT-PCR for hepatic ER $\alpha$

ER $\alpha$  mRNA assessment in liver was conducted using RT-PCR while real time RT-PCR was used for determination

of ER $\alpha$  and ER $\beta$  mRNA in heart because of technical problems. Total RNA was isolated from rat frozen liver with Trizol reagent (Life Technologies, Rockville, MD) according to manufacturer's specifications. Total RNA was then treated with RNase-free DNase I under a standard protocol. The integrity and quality of the purified RNA were controlled by formaldehyde denaturing agarose gel electrophoresis and measurement of the  $A_{260}/A_{280}$  ratio. First-strand cDNA was synthesized in a final volume of 40  $\mu\text{l}$  containing first-strand buffer, 2  $\mu\text{g}$  RNA, 2  $\mu\text{g}$  hexanucleotide primer (Life Technologies), and avian myeloblastosis virus reverse transcriptase (12 units/ $\mu\text{g}$  RNA, Life Technologies). About 5  $\mu\text{l}$  of first-strand cDNA were added to a PCR mixture and amplified for 27 cycles by incubation at  $95^{\circ}\text{C}$  for 1 min, at  $55^{\circ}\text{C}$  for 50 s and at  $72^{\circ}\text{C}$  for 1 min 10 s, with final incubation at  $72^{\circ}\text{C}$  for 3 min, all in a Robocycler gradient 40 thermocycler (Stratagene, La Jolla, CA). The ER $\alpha$  primer sequences are enumerated in Table 1. Amplification of 18S RNA by oligonucleotides was achieved according to the protocol provided by the manufacturer (Ambion, Austin, TX). Control RT-PCR omitted reverse transcriptase or RNA from the reaction mixture. After amplification, the samples were loaded and electrophoresed on 1.5% agarose gel. Bands stained by ethidium bromide were counted and analyzed with the Storm 840 imaging system and ImageQuant software (Version 4.2, Molecular Dynamics, Sunnyvale, CA). To validate RT-PCR as a tool for the semi-quantitative measurement of mRNA, dose–response curves were charted for different amounts of total RNA extracted from the rat liver, and the samples were quantified in the linear phase of PCR amplification. These data were normalized to the corresponding values of 18S RNA. All samples were run in duplicate. The intra- and inter-assay coefficients of variation were 6 and 8%, respectively.

#### Isolation of RNA and semi-quantitative real-time PCR for ER $\alpha$ and ER $\beta$ in heart

Total RNA was extracted from frozen hearts with Trizol (Invitrogen Life Technologies, Inc., Burlington, ON)

**Table 1** Oligonucleotide primers used for reverse transcriptional polymerase chain reaction (RT-PCR), and real-time quantitative PCR

Gene	Sense primer (5'–3')	Antisense primer (5'–3')	Accession No.
Erz	GCGGCTGCCACTGACCATG	CCTCGGGGTAGTTGAACACGG	NM_012689
ER $\alpha$ (real time)	CCAAAGCCTCGGGAATGG	AGCTGCGGGCGATTGAG	NM_012689
ER $\beta$ (real time)	TTGGTGTGAAGCAAGATCACTAGAG	AACAGGGCAGGCACAACCTG	NM_012754
GADPH (real time)	TTCAATGGCACAGTCAAGGC	TCACCCATTTGATGTTAGCG	NM_017008
Beta Actin	ACCCACACTGTGGCCCATCTA	GCCACAGGATTCCATACCCA	NM_031144

Estrogen receptor-alpha (ER $\alpha$ ); estrogen receptor-beta (ER $\beta$ ); glyceraldehyde-3-phosphate dehydrogenase (GADPH). 18S primer pair universal 18S (Ambion, Lot N: 115R057113A) was bought directly from Ambion

according to manufacturer's protocol. To remove genomic DNA, RNA samples were incubated with 2 U deoxyribonuclease I (DNase I; Invitrogen Life Technologies, Inc., Burlington, ON)/ $\mu\text{g}$  RNA for 30 min at 37°C. The cDNA synthesis was performed as previously described. PCR was carried out in the iCycler IQ Real time PCR detection system (Bio-Rad Laboratories, Inc., Hercules, CA), using SYBR<sup>®</sup> green chemistry. Samples were analyzed in duplicate or triplicate. For amplification, 2  $\mu\text{l}$  of diluted cDNA was added to a 20  $\mu\text{l}$  reaction mixture containing 1 $\times$  iQ SYBR Green Supermix (Bio-Rad Laboratories, Inc.) and 200 nM forward and reverse primers. The thermal cycling program was 95°C for 2 min, followed by 40 cycles of 95°C for 30 s, 60°C for 30 s, and 72°C for 30 s. Primers were purchased from Invitrogen Life Technologies, Inc. Primer pairs were chosen to minimize primer dimerization and to generate an amplicon between 150 and 350 bp. The primers used are enumerated in Table 1. Optical data were collected during the annealing step of each cycle. Following PCR, reaction products were melted for 1 min at 95°C, and then the temperature was lowered to 55°C, and then gradually increased to 95°C in 1.0°C increments, 10 s per increment. Optical data were collected over the duration of the temperature increase with a dramatic drop in fluorescence occurring. This was done to ensure that only one PCR product was amplified per reaction.

The relative expression of the RT-PCR products was determined using the  $\Delta\Delta\text{Ct}$  method. This method calculates relative expression using the equation: Fold induction =  $2^{-[\Delta\Delta\text{Ct}]}$ , where Ct = the threshold cycle, i.e. the cycle number at which the sample's relative fluorescence rises above the background fluorescence and  $\Delta\Delta\text{Ct}$  = [Ct gene of interest (unknown sample) – Ct GAPDH (glyceraldehyde-3-phosphate dehydrogenase) (unknown sample)] – [Ct gene of interest (calibrator sample) – Ct GAPDH

(calibrator sample)]. One of the control samples was chosen as the calibrator sample and used in each PCR. Each sample was run in duplicate and the mean Ct was used in the  $\Delta\Delta\text{Ct}$  equation. GAPDH was chosen for normalization instead of 18S RNA because the latter is rapidly amplified and taking into consideration the high sensitivity of real time RT-PCR, it is difficult to determine Ct for 18S RNA. GAPDH showed consistent expression relative to other housekeeping genes among the treatment groups in our array experiments.

#### Statistical analysis

Values are expressed as means  $\pm$  SE. Statistical analyses were performed by a two-way ANOVA for non-repeated measures. Fisher's PLSD post hoc test was used in the event of a significant ( $P < 0.05$ ) ratio.

#### Results

Comparison of body composition parameters in Sham and Ovx rats confirms that ovariectomies were well performed (Table 2) [31]. Ovx rats showed higher ( $P < 0.01$ ) body weight, energy intake, intra-abdominal fat depot, and right triceps surae weights than Sham rats. Moreover, uterus weight was significantly lower in Ovx rats ( $P < 0.01$ ), indicating total ovariectomy.

Endurance training led to increased ( $P < 0.05$ ) body weight and energy intake in Sham rats, suggesting that intact animals did not experience negative energetic balance during the training period (Table 2). Exercise training resulted in a significant ( $P < 0.01$ ) decrease in intra-abdominal fat depot weight when expressed per unit of body weight in both Sham and Ovx rats reflecting the

**Table 2** Body weight, daily energy intake, intra-abdominal fat depots, left ventricle weight, femur weight, right triceps surae weight, and uterus weight, in sham-operated (Sham), sham-

operated + endurance training (ShamTr), ovariectomized (Ovx), and ovariectomized + endurance training (OvxTr) rats

	Sham	ShamTr	Ovx	OvxTr
Body weight (g)	287 $\pm$ 7	330 $\pm$ 14 <sup>+</sup>	394 $\pm$ 9**	377 $\pm$ 9*
Energy intake (kJ/day)	259 $\pm$ 6	288 $\pm$ 10 <sup>+</sup>	321 $\pm$ 8**	310 $\pm$ 8
Intra-abdominal fat depot weight (g)	15 $\pm$ 2	14 $\pm$ 1	33 $\pm$ 2**	23 $\pm$ 3* <sup>+</sup>
Intra-abdominal fat depot weight/body weight (g/g)	0.053 $\pm$ 0.005	0.044 $\pm$ 0.003 <sup>++</sup>	0.084 $\pm$ 0.005**	0.061 $\pm$ 0.007*** <sup>++</sup>
Left ventricle weight (mg)	617 $\pm$ 28	660 $\pm$ 35	732 $\pm$ 27**	791 $\pm$ 30**
Left ventricle weight/body weight (mg/g)	2.11 $\pm$ 0.06	1.99 $\pm$ 0.04	1.84 $\pm$ 0.03**	2.09 $\pm$ 0.09 <sup>+</sup>
Femur weight (mg)	728 $\pm$ 18	838 $\pm$ 39 <sup>++</sup>	772 $\pm$ 18	851 $\pm$ 13 <sup>++</sup>
Femur weight/body weight (mg/g)	2.54 $\pm$ 0.07	2.60 $\pm$ 0.05	1.96 $\pm$ 0.04**	2.25 $\pm$ 0.05*** <sup>++</sup>
Right triceps surae weight (g)	2.03 $\pm$ 0.05	2.21 $\pm$ 0.10	2.46 $\pm$ 0.05**	2.50 $\pm$ 0.04**
Uterus weight (g)	0.61 $\pm$ 0.08	0.81 $\pm$ 0.07	0.11 $\pm$ 0.01**	0.10 $\pm$ 0.01**

Values are means  $\pm$  SE,  $n = 8$  rats/group. \* $P < 0.05$  and \*\* $P < 0.01$ : significantly different from corresponding Sham group. <sup>+</sup> $P < 0.05$  and <sup>++</sup> $P < 0.01$ : significantly different from corresponding sedentary rats

**Table 3** Summary of the results on ER transcripts

	Effect of Tr (Sham)		Effect of Ovx		Effect of Tr (Ovx)	
	ER $\alpha$	ER $\beta$	ER $\alpha$	ER $\beta$	ER $\alpha$	ER $\beta$
Liver	↑300%	⊗	↑200%	⊗	↓50%	⊗
LV	↓80%	↓40%	↓80%	↓100%	↑60%	N.S.
RA	N.S. (↑150%)	↑200%	↑200%	N.S.	N.S.	N.S.

Estrogen receptor-alpha (ER $\alpha$ ); estrogen receptor-beta (ER $\beta$ ); ovariectomy (Ovx); training (Tr); left ventricle (LV); right atrium (RA)

Comparisons are made with respective control groups. ⊗—The isoform is not expressed in that tissue. N.S.—No significant effect, otherwise all other indicated changes are significant,  $P < 0.05$

effectiveness of the running program. In addition, femur weight was significantly increased ( $P < 0.01$ ) with training in both Sham and Ovx groups.

The RT-PCR technique was used to demonstrate ER transcripts changes in response to endurance training in the liver and heart chambers (LV and RA) of intact and Ovx female rats. Our results confirmed that ovariectomy induced tissue-specific alterations of ER mRNA levels [41], and showed for the first time that endurance training affects ER expression in liver and heart.

A summary of the results on ER transcripts is presented in Table 3. ER $\alpha$  mRNA levels were increased by 200% in the liver and RA and decreased by 80% in LV of sedentary Ovx compared to Sham rats ( $P < 0.05$ ; Fig. 1). Different effects of endurance training on ER $\alpha$  expression were observed in the liver and heart chambers, and between intact and Ovx rats. Hepatic ER $\alpha$  transcripts content was significantly higher (300%,  $P < 0.01$ ) in Sham, but 50% lower ( $P < 0.01$ ) in Ovx rats after endurance training (Fig. 1A). On the other hand, ER $\alpha$  mRNA levels in LV were decreased by 80% ( $P < 0.05$ ) in Sham, and increased by 60% ( $P < 0.01$ ) in Ovx rats after endurance training (Fig. 1B). There were no significant effects of training on ER $\alpha$  transcripts content in RA (Fig. 1C).

ER $\beta$  transcripts levels were completely depressed by Ovx in LV (100%;  $P < 0.01$ ; Fig. 2A), but remained unchanged in RA of sedentary rats (Fig. 2B). ER $\beta$  mRNA levels were 40% lower ( $P < 0.05$ ; Fig. 2A) in LV, but 200% higher ( $P < 0.05$ ; Fig. 2B) in RA of Sham rats after endurance training. However, endurance training did not change ER $\beta$  mRNA content in either LV or RA of Ovx rats (Fig. 2A and B).

## Discussion

The purpose of the present study was to investigate the effects of an 8-wk endurance training regimen on ER transcripts levels in liver and heart of female rats. We found significant alterations of ER expression in the liver, left ventricle, and right atrium of endurance-trained animals.

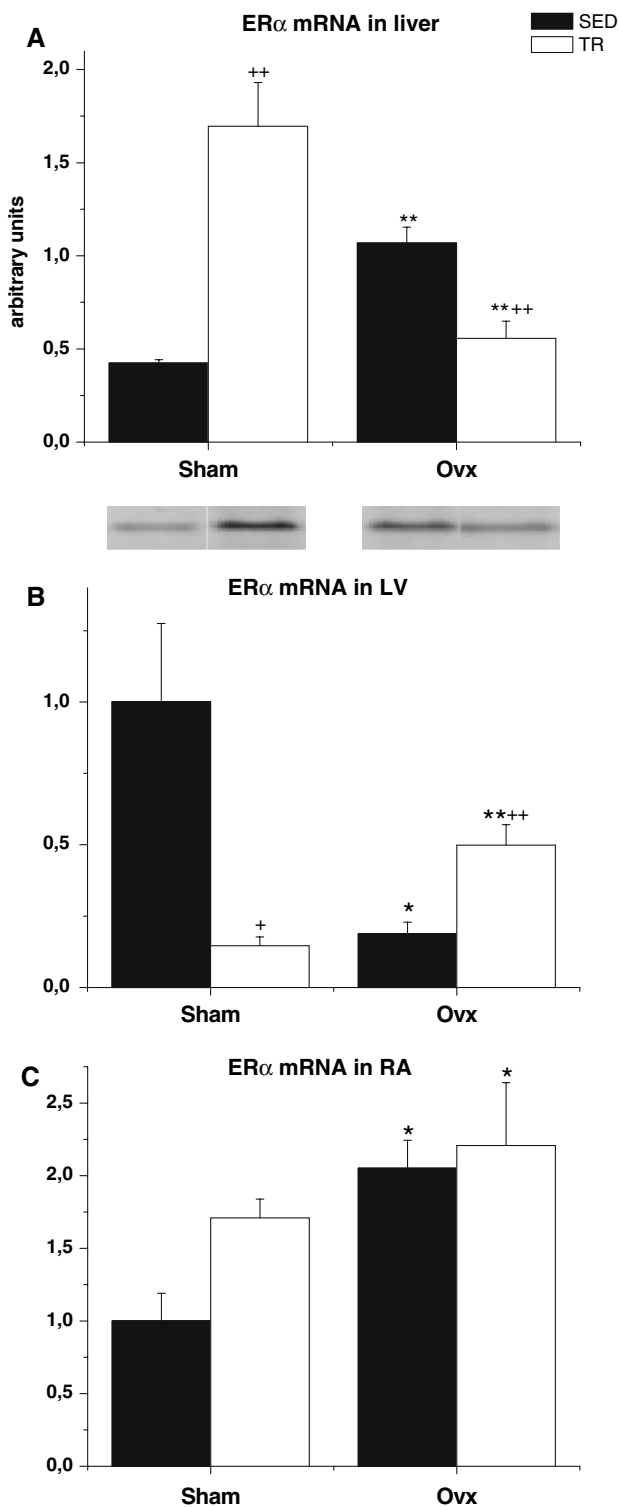
More specifically, we showed that these effects are tissue and isoform-specific and that the direction of the changes in transcripts levels is different in Ovx than in intact rats.

### Training in Sham rats

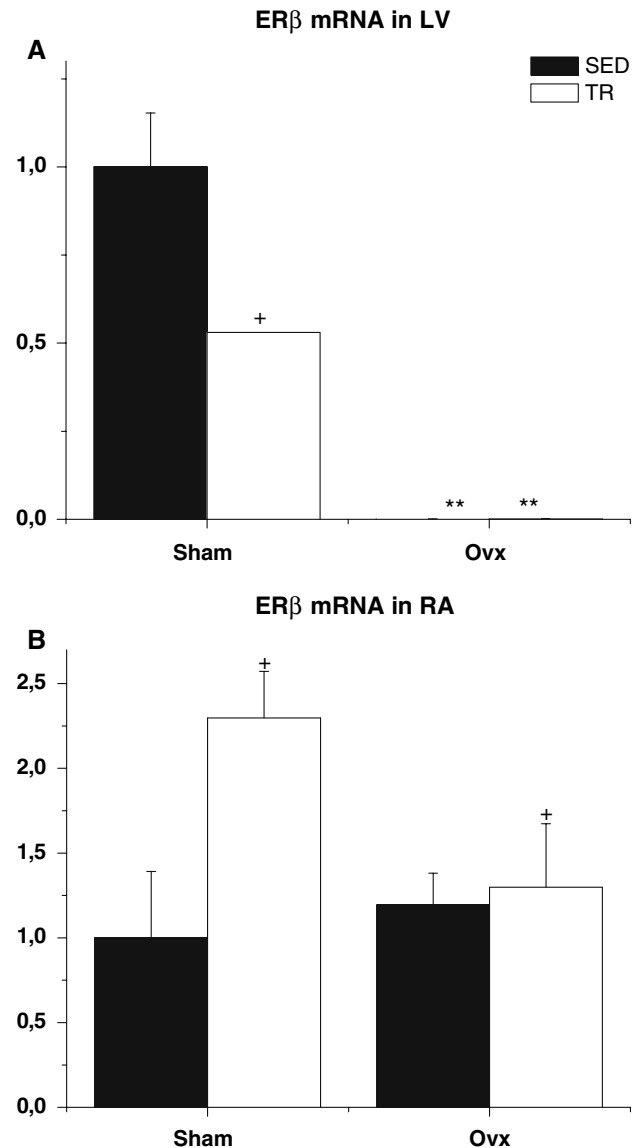
The most novel finding of the present study is the clear effect of endurance training on ER transcripts content in liver and heart. We found an important increase (300%) in hepatic ER $\alpha$  mRNA levels, accompanied by a decreased expression of both ER $\alpha$  and ER $\beta$  genes in LV, and an increase (200%) in ER $\beta$  transcripts content in RA following endurance training in intact rats. Our results are in line with the tissue-specific regulation of ER, which has been shown to be under the control of several factors including pituitary and gonadal hormones [14, 15, 23]. The present results extend these findings by indicating that an endurance-trained state also influence ER gene expression in liver and heart.

Our results raise the question as to why ER expression changes following endurance training in liver and heart. Endurance training is known to stimulate the expression of transcription factors, leading to the synthesis of proteins involved in the adaptations of skeletal muscle to training. For instance, the increased expression of the transcription factor NRF-1 takes part in the mitochondrial biogenesis processes in muscle in response to endurance training [42]. ERs are themselves transcription factors and variations in their transcripts levels may lead to stimulation or repression of target genes [1, 2]. Lemoine et al. [24, 25] were first to report an effect of endurance training on ER $\alpha$  gene expression in skeletal muscle. They put forward the concept that an ER-mediated mechanism could contribute to the training adaptations of skeletal muscle [24, 25]. The stimulation of mitochondrial biogenesis is an adaptation to training that may also be mediated by an ER-dependent mechanism in skeletal muscle, as a positive correlation was reported between ERs levels and citrate synthase activity [26]. Our data suggest that this concept could be extended to liver and heart. It seems likely that endurance training





**Fig. 1** ER $\alpha$  mRNA levels in the liver, left ventricle (LV), and right atrium (RA) of sham-operated (Sham), sham-operated + endurance training (ShamTr), ovariectomized (Ovx), and ovariectomized + endurance training (OvxTr) rats. Values are means  $\pm$  SE,  $n = 8$  rats/group. \* $P < 0.05$  and \*\* $P < 0.01$ : significantly different from corresponding Sham group. \* $P < 0.05$  and \*\* $P < 0.01$ : significantly different from corresponding sedentary rats. Values were normalized to values of Sham-sed groups in (B) and (C)



**Fig. 2** ER $\beta$  mRNA levels in left ventricle (LV) and right atrium (RA) of sham-operated (Sham), sham-operated + endurance training (ShamTr), ovariectomized (Ovx), and ovariectomized + endurance training (OvxTr) rats. Values are means  $\pm$  SE,  $n = 8$  rats/group. \* $P < 0.05$  and \*\* $P < 0.01$ : significantly different from corresponding Sham group. \* $P < 0.05$ : significantly different from corresponding sedentary rats. Values were normalized to values of Sham-sed groups. Values for Ovx and OvxTr rats were  $0.001 \pm 0.0003$  and  $0.001 \pm 0.001$ , respectively

and estrogens share target genes in the liver. For instance, both endurance training and estrogens, through ER $\alpha$ , repressed the expression and activity of hepatic lipase, leading to increased levels of circulating HDL [22, 27, 28, 43]. We found important increases in hepatic ER $\alpha$  mRNA levels in endurance-trained rats. It is thus possible that the stimulation of the estrogen signaling pathway through endurance training participates in the repression of hepatic lipase by training.

Aside from the liver, the physiological role of estrogens on the myocardium is poorly understood, making the interpretation of our training effects on ER transcript levels more difficult. However, several associations exist for the ER content and myocardial functions and dysfunctions [44, 45]. For instance, ER are involved in the stability of cardiac intercalated discs and an up-regulation of ER content in myocardium is related to myocardial pressure load as a compensatory mechanism [44, 45]. The decreased levels of ER $\alpha$  mRNA content found in LV of our endurance-trained rats may, therefore, be interpreted as an indicator of an improvement in heart function. Nevertheless, it is difficult to reconcile the important decrease in both ER $\alpha$  and ER $\beta$  genes in LV following endurance training with the known cardioprotective effect associated with training and with the administration of estrogens [46, 47]. It is possible that both of these actions occur through a different mechanism. Recent evidence suggests that ER $\beta$  attenuation of cardiac dysfunction following trauma-hemorrhage is mediated via a nuclear as well as a mitochondrial action [48]. This reemphasizes our limited understanding of the effects and the mechanism of action of ER $\alpha$  and ER $\beta$  on the cardiovascular system [49]. The 80% decrease in LV ER $\alpha$  in Sham rats subjected to exercise can be related to observed increase of LV mass in these animals. It is well known that physiological cardiac hypertrophy, caused by chronic exercise training and defined as athlete's heart is part of beneficial adaptive responses of the cardiovascular system to exercise training. Recently it was reported that estrogen treatment attenuates the increase in relative heart weight in the ovariectomized rats in association with ER $\alpha$  [50]. Therefore, training-related lowering of ER $\alpha$  in Sham can serve to promote development of physiological LV hypertrophy.

#### Effects of Ovx

It has been proposed that endogenous estrogens may be repressive on hepatic ER $\alpha$  gene expression of intact rats, as an elevation of ER $\alpha$  mRNA levels is observed after Ovx [17, 18]. Accordingly, we found an increase in hepatic ER $\alpha$  mRNA content after Ovx. On the other hand, the consequences of ovaries removal on ER transcripts levels in heart are poorly defined. A positive regulation of ER $\alpha$  levels by endogenous estrogens was reported in female rat atria following Ovx [23], while other reports showed no significant changes [41]. Our study is the first, to our knowledge, to report effects of Ovx on both ER isoforms transcripts levels in two different heart chambers of the female rat. We observed decreased ER $\alpha$  and ER $\beta$  mRNA levels in LV, and increased content of ER $\alpha$  transcripts in RA by Ovx. These results indicate the positive and

negative regulation of ER mRNA levels in LV and RA, respectively, by sex steroids. Therefore, our findings in liver and heart chambers of Ovx rats further highlight the reported tissue-specific regulation of ERs by endogenous ovarian hormones.

#### Training in Ovx rats

Taking into account the fact that ER transcripts were largely modified by the Ovx, the second major finding of the present study is the different effect of endurance training found on ER transcripts levels according to the presence or the absence of ovaries. The training adaptations of ER $\alpha$  mRNA content in the liver and LV of Ovx animals were opposite to the training response observed in Sham rats. In addition, while exercise training altered ER $\beta$  mRNA content in LV and RA of Sham rats, it had no effect in Ovx rats. These findings suggest that the presence or absence of ovaries influence the response of ER gene expression to the stimulus of exercise. This is in line with the action of estradiol in liver that has been reported to depend on the presence of the pituitary hormones [12, 14, 15]. Thus, it seems that the responsiveness of the ER gene in liver and heart to the stimulus of exercise training is different according to the presence of ovarian steroids. The response of ER transcripts to exercise training in the present Ovx rats also suggest that exercise training compensated for the ovarian steroid deprivation in liver and heart. It is important to recall that the transcriptional activity of ER can be modified by factors other than estrogens [3, 51]. Exercise-induced activation of MAPK can lead to phosphorylation of the AF-1 domain of the ER gene, thus resulting in the stimulation or the repression of ER transcriptional activity [52, 53].

Although ER $\alpha$  is the main receptor known to mediate the estrogenic effects, both isoforms mediate specific actions, and the relative importance of the two isoforms is not well understood [2]. Our results also suggest that the ER gene expression in RA is less sensitive to different stimuli, e.g. ovariectomy (ovarian steroids deprivation) and exercise training, than the ER gene expression in the liver and LV.

There are limitations to the present study that need to be addressed. It could be argued that our training program disrupted the estrus cycle of intact rats and altered ER expression, since circulating estrogens can regulate the levels of their own receptors [23, 54, 55]. However, training-induced disruption of the estrus cycle is often associated with body weight loss and low energy intake [56, 57]. This was hardly the case in our study since intact female rats gained body weight and increased their energy intake with training. Moreover, changes in ERmRNA

levels following endurance training were also observed in Ovx rats. Although plasma estradiol levels were not measured in the present study, it seems likely that endurance training was indeed responsible for the changes that we observed on ER expression in both intact and Ovx rats. On the other hand, the physiological interpretation of the present data is limited by the sole measurement of transcripts levels. However, the demonstration that these ER gene transcripts are modified with endurance training in liver and heart constitutes an interesting opening in a new avenue of research linking ER as a possible contributor to the training adaptations in these tissues.

In conclusion, we showed, using the RT-PCR technique that endurance training is associated with modifications of ER gene expression in liver, right atrium, and left ventricle of the heart. We demonstrated that these training effects are tissue and isoform specific and that the responsiveness of the ER gene to the stimulus of endurance training is different in intact and Ovx animals. It is suggested that ER gene expression may mediate some of the adaptive effects of endurance training in liver and heart.

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