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Gene expression profile of estrogen receptors alpha and beta in rat brain during aging and following high fat diet

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ABSTRACT

The "sex-hormone" estrogen-17 β promotes several cognitive functions and is a master regulator of brain bioenergetics via the estrogen receptors α and β (*ER* α and *ER* β). In this work, by using Real-Time PCR analysis, we evaluated the effect of aging and high fat diet (HFD) on *ER* α and *ER* β expression in rat hippocampus and cortex. In young rats, *ER* β is abundant in cortex and *ER* α and *ER* β expression; in hippocampus and cortex. In young rats, *ER* β is abundant in cortex and *ER* α and *ER* β expression; in hippocampus *ER* α increases and *ER* β decreases. *ER* expression patterns in rat brain are also affected by the administration of an HFD. In cortex, after 4 weeks of HFD, *ER* β transcripts are down-regulated, whereas *ER* α levels remain unchanged; after 12 weeks, both ER α and *ER* β expression is up-regulated. In the hippocampus, the level of *ER* β transcripts does not change following HDF, whereas *ER* α expression is affected by HDF, in a time-dependent manner: it increases after the 4-week treatment and decreases after 12 weeks. Possible involvements of these receptors in the control of cortex and hippocampus functions during aging and in the modulation of energetic metabolism and feeding behaviour are discussed.

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1. Introduction

Estrogens are sex hormones involved in the development and maintenance of normal reproductive functions [1–3]. These hormones exert their biological effects through the interaction of two primary estrogen receptors, members of the nuclear steroid receptors superfamily: estrogen receptor-alpha (ER α), and estrogen receptor-beta (ER β).

 17β -Estradiol (E2) is the most potent and predominant form of estrogen; besides its recognized role in reproduction, it shows a number of effects on cognition and brain function [4–7]. In different brain regions, aromatase activity catalyzes E2 synthesis from androgen precursors and/or cholesterol. E2 is neurotrophic and neuroprotective and some recent results, although controversial, suggest

* Corresponding author. E-mail address: ilaria.verderame@unina.it (M. Verderame). that E2 synthesized within hippocampus and prefrontal cortex may contribute to the normal processes of memory consolidation [8–11]. The effects of E2 on several brain processes decline with age, including the ability to preserve the blood–brain barrier and to protect from stroke and neuroinflammation [12,13].

Accordingly with E2 functions, both $ER\alpha$ and $ER\beta$ subtypes are widely distributed in the brain regions, in particular in the hippocampus, the frontal cortex, and the amygdala [4,14]. The alteration in the normal distribution of ERs has been observed in hippocampal neurons of Alzheimer's disease (AD) patients [15].

It has been demonstrated that the two ER subtypes play different roles in E2-regulated gene expression and neuronal physiology [16]; the differential activation of $ER\alpha$ or $ER\beta$ is able to influence the expression of neuroprotective genes [17,18].

Data on the differential expression of *ER* genes in the central nervous system (CNS), mostly achieved in females

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and collected in the rat, the mouse or the human, are often contradictory [19–21]. In the rat hippocampus, $ER\alpha$ is transcriptionally more active than $ER\beta$ [7], and a significant decrease in the number of neurons positive to estrogen receptors during aging is observed [22]. In addition, *ERs* seem to have a gender specificity; indeed, *ERβ* expression in the female hippocampus is greater than in the male one [23]. In the cerebral cortex of the same animals, the limited data available show *ERβ* as the predominant receptor subtype immunolocalized [24].

In mouse, $ER\alpha$ is the predominant subtype in the hippocampus and in most of hypothalamus, whereas in the cerebral cortex $ER\beta$ is the main receptor subtype observed [25]. Conversely, in human and primates $ER\beta$ is the primary receptor reported in hippocampus [26,27]. In humans, it has been found that following the neurological diseases influencing memory, such as AD, the expression of $ER\alpha$ decreases while $ER\beta$ increases [28–31].

The relationship between E2 and ERs in the brain involves the regulation of food intake. In women, the daily food intake changes in relation to the E2 secretion during the menstrual cycle: during the follicular phase, the daily food intake is lower in respect to the luteal phase [32,33]. Also in rat, the increase in plasma E2 concentration during the preovulatory phase is associated with a transient decrease in food intake [34]. In the CNS, both $ER\alpha$ and $ER\beta$ are expressed in areas associated with satiety and feeding, such as hypothalamus and pituitary [35]; it has been also demonstrated that ER activation is involved in the metabolic effect of insulin signaling in the brain [36]. Finally, E2 has been proposed to act directly and indirectly to decrease orexigenic or anorexigenic peptides and food intake: for instance, $ER\alpha$ null mice are obese, insulin resistant, and show decreased energy expenditure [35,37].

In this study, by using Real-Time PCR analysis, we investigated the expression pattern of $ER\alpha$ and $ER\beta$ genes during aging and after a high fat diet (HFD) in rat brain. In particular, we evaluated the $ER\alpha$ and $ER\beta$ expression profiles in the cerebral cortex and hippocampus of pubertal young (2-months), adult (4- and 8-month), and middle-aged (16-month) rats. Our results demonstrate that $ER\alpha$ and $ER\beta$ subtypes have specific tissue- and age-related expression patterns that are unbalanced by an HFD that leads to the obese phenotype.

2. Materials and methods

2.1. Animal and experimental design

Male Wistar rats (Charles River, Calco, Como, Italy) of 2 months of age (adolescent), with the same starting body weight $(160 \pm 10 \text{ g})$, were individually caged in a temperature-controlled room $(23 \pm 1 \,^{\circ}\text{C})$ with a 12 h light/12-h dark cycle. Animals were housed in the Animal Care Facility at the Department of Biology, with *ad libitum* access to water and to a standard diet (SD, 10.6% fat J/J, 60,4% carbohydrate, 15.47 KJ/g) (Mucedola 4RF21; Settimo Milanese, Milan, Italy) up to 4 (social maturity, group 1), 8 (adulthood, group 2), or 16 (middle-aged, group 3) months. Six animals for each group

were anesthetized with chloral hydrate (40 mg/100 g body wt) and killed by decapitation. Group 4 was made of eight adolescent rats sacrificed four days after their arrival in the animal facility.

In parallel, another group of rats (referred to as HFD, n = 12) received a high-fat diet rich in lard (40% fat J/J, 31% carbohydrate, 19.23 KJ/g). The HFD was formulated to differ from the standard diet in the fat and carbohydrate contribution to the energy value but to be identical in terms of proteins, vitamins, minerals and fibers [38]. Six animals of both SD and HFD groups were sacrificed after 4 weeks and other six individuals after 12 weeks. Throughout the experimental period, body weights and food intakes were monitored daily to calculate the body-weight gain. Spilled food was collected and compensated in readjusting the calculation of food intake.

At the end of the experimental period, animals of each group were anesthetized by chloral hydrate (40 mg/100 g body wt) and killed by decapitation. The brains were quickly removed and the cerebral cortex and hippocampus were dissected on ice. Samples of each brain region were snap frozen in liquid nitrogen immediately and stored at -80 °C for subsequent RNA isolation. The study was performed in strict accordance with the criteria established by the National Institutes of Health. The protocol for animal care and use was approved by the Committee on the Ethics of Animal Experiments of the University of Naples Federico II.

2.2. RNA purification and cDNA synthesis

Total RNA was extracted according to the TRI-Reagent (Sigma Aldrich) protocol. The quality of each total RNA was checked by electrophoresis on 2% agarose gel stained with ethidium bromide and measuring the optical density at 260/280 nm. A ratio of 1.8–2.0 was accepted for further reverse transcription. The QuantiTect Reverse Transcription Kit (Qiagen) was used for the removal of genomic DNA contamination and for the subsequent cDNA synthesis. Approximately 1 µg of total RNA was used, according to the kit's protocol.

2.3. Quantitative Real-Time PCR analysis

The Real-Time PCR reactions were carried out in triplicate for each sample in an Applied Biosystems 7500 Real-Time System by using the Power SYBR Green Master Mix PCR (Applied Biosystems) following the procedures recommended by the manufacturer [39]. Each SYBR Green reaction (20-µl total volume) contained 12 µL of Real-Time PCR Master Mix, 1 µL of each of the forward and reverse primer (10 μ M), 2 μ L of cDNA diluted 1:1 and 4 µL of nuclease free water. For internal standard control, the expression of the β -actin gene was quantified [40]. A single pair of specific primers for both $ER\alpha$ and $ER\beta$ isoforms was designed on the nucleotide sequences of *Rattus norvegicus* $ER\alpha$ (NM012689.1) and $ER\beta$ (AF042058.1); β -actin primers were designed on the R. norvegicus template (NM031144.2). Primer sequences are reported in Table 1. The lengths of the obtained cDNA fragments were: $ER\alpha$, 216 bp; $ER\beta$, 262 bp; β -actin, 128 bp.

 Table 1

 Specific primers used for Real-Time PCR analyses.

Name	Nucleotiode sequence (5'-3')	Length (mer)	
ERa_F	GCACATTCCTTCCTTCCGTC	20	
$ER\alpha_R$	CTCGTTCCCTTGGATCTGGT	20	
ERβ_F	ACAGTCCTGCTGTGATGAAC	20	
ERβ_R	ACTAGTAACAGGGCTGGCAC	20	
β-act_F	ACCCGCCACCAGTTCGCCAT	20	
β-act_R	CGGCCCACGATGGAGGGGAA	20	

The PCR was performed under the following conditions: holding stage of 95 °C per 10 min; cycling stage (45 cycles): 95 °C × 10 s-60 °C × 10 s-72 °C × 10 s; melting stage: 95 °C × 5 s-65 °C × 1 min-95 °C × 30 s-40 °C × 30 s. The melting curve analysis of PCR products was performed in order to ensure gene-specific amplification. Changes in the gene expression relative to the different samples were calculated according to the standard $2^{-\Delta\Delta Ct}$ method described by Livak and Schmittgen [41].

2.4. Statistical analysis

Data are presented as mean \pm standard error of the mean (SEM) from four separate experiments of each sample. Statistical analyses were carried out by GraphPad Prism software. The differences between the mean values were analyzed by one-way analysis of variance (ANOVA), followed by Fisher's LSD test. The differences were considered significant when p < 0.05.

3. Results

3.1. Expression of ER α and ER β in cortical and hippocampal areas of 2 month-old rats

First, we investigated the relative abundance of $ER\alpha$ and $ER\beta$ in the cerebral cortex and hippocampus of young rats. We found that $ER\beta$ was more abundant in the cerebral cortex, whereas in the hippocampus $ER\alpha$ was dominant compared to the cortex (Fig. 1). In general, however, low



Fig. 1. Real-Time PCR analysis of $ER\alpha$ and $ER\beta$ expression in cerebral cortex and hippocampus of 2 month-old rats. $ER\alpha$ and $ER\beta$ mRNA levels are shown relative to the β -actin mRNA. The data represent the mean \pm s.e.m.

but significant amounts of both $ER\alpha$ and $ER\beta$ were found in these brain regions.

Real-Time PCR analysis carried out in the cerebral cortex revealed a general decrease in $ER\alpha$ and $ER\beta$ expression during aging (Fig. 2). In particular, $ER\alpha$ transcripts decreased by half in 4 month-old rats, then the amount of transcripts remained constant; the $ER\beta$ transcript amount rapidly decreased to about one-tenth in 4 month-old rats, before rising in middle-aged rats at a fifth of the value measured in young rats (Fig. 2).

In the hippocampus, the expression of $ER\alpha$ greatly increased with aging, whereas $ER\beta$ expression generally decreased (Fig. 2). Compared to 2 month-old rats, the level of $ER\alpha$ transcript was almost unchanged in 4-month-old rats and sharply increased by about 5 and 25 times in adult (8-month-old) and middle-aged (16 month-old) rats, respectively (Fig. 2). As regarding $ER\beta$, a significant decrease in expression was observed in 4- and 8 monthold rats, with mRNA levels 1/5 lower than in 2 month rats; in middle-aged rats, the $ER\beta$ expression level slightly increased, although remaining below the value observed in young rats (Fig. 2).

3.2. HFD-related changes of ER and ER β expression in rat brain

As no data are available on the effect of a HFD on the brain expression of *ERs*, we evaluated $ER\alpha$ and $ER\beta$ mRNA levels in cortex and hippocampus of rats fed a HFD for 4 or 12 weeks.

First of all, we determined the weight gain of HFD-fed rats. Both control and HFD rats showed a significantly weight gain during housing (Table 2). The increase in body weight was affected by the duration of the housing and diet administration. In control animals, the weight gain was of 18% after 4-week and 35% after 12-week housing, respectively. In HFD rats, the weight gain was 47% after 4 weeks and about 80% after 12 weeks, with an increase by 25% between the two 4-week groups and of 48% in the 12-week group.

ER expression, as detected by Real-Time PCR, was significantly affected by the diet. In the cortex of HFD rats receiving the diet for 4 weeks, the level of $ER\alpha$ transcripts did not change, whereas the level of $ER\beta$ transcripts was lower than in control animals (about 1/5, Fig. 3). Interestingly, after 12 weeks of HF diet, both $ER\alpha$ and $ER\beta$ transcription was dramatically up-regulated (54-fold and 141-fold, respectively) (Fig. 3).

In the hippocampus of rats receiving the HF diet for 4 weeks, the level of $ER\alpha$ transcripts increased about twofold with respect to control rats, but decreased below the value determined in the control rats, when the treatment was extended for 12 weeks (Fig. 3). As regarding $ER\beta$, the expression level was not affected by the HFD, no matter how long it was (Fig. 3).

4. Discussion

Over the past 25 years, evidence shows that estradiol modulates cognitive function in mammals, including humans. The E2 action begins in utero when estrogens



Fig. 2. Effect of aging on $ER\alpha$ and $ER\beta$ gene expression in rat cerebral cortex (A) and hippocampus (B). The $ER\alpha$ and $ER\beta$ mRNA expression was normalized to that of β -actin mRNA and converted in fold change, compared with the 2 month-old rats. The data represent the mean \pm s.e.m. Significance of differences is shown. *p < 0.05, **p < 0.001, a: significance vs. 2 months; aa: significance vs. 4 and 8 months; aaa: significance vs. 8 month.



Fig. 3. Effect of high fat diet on $ER\alpha$ and $ER\beta$ gene expression in rat cerebral cortex (A) and hippocampus (B). The $ER\alpha$ and $ER\beta$ mRNA expression was normalized to that of β -actin mRNA and converted in fold change, compared with rats fed with a standard diet. The data represent the mean \pm s.e.m. Significance of differences is shown. *p < 0.05, **p < 0.001. a: significance vs. standard diet; aa: significance vs. high fat diet for 4 weeks.

 Table 2

 Average body weight in control and HFD rats.

	Control			High fat diet	
	0 weeks	4 weeks	12 weeks	4 weeks	12 weeks
Body weight (g) Weight gain (%)	311.47 ± 14.3	369.22 ± 15.0 18.5	$\begin{array}{c} 422.69 \pm 18.9 \\ 35.7 \end{array}$	$\begin{array}{c} 460.17\pm32.1\\ 47.7\end{array}$	$\begin{array}{c} 615.4 \pm 41.2 \\ 97.5 \end{array}$

direct sexual differentiation of the brain regions controlling reproduction and some cognitive functions linked to it. They reach the highest levels during puberty; with aging, levels of circulating estrogens collapse, contributing to the age-related declines in learning and memory function.

The aim of this study was to assess whether changes in *ERs* occur with aging in *rattus* cerebral cortex and hippocampus. Investigations were performed in males to avoid the influence of the fluctuations of steroids, mainly of E2, on the expression and localization of these receptors in brain [42–44].

First, our results demonstrate that the two *ER* subtypes are differentially expressed in the two brain regions examined. Indeed, regardless of the rats' age, in cerebral cortex the *ER* β subtype is more abundant with respect to the hippocampus; conversely, in the latter, $ER\alpha$ is more abundant. However, our results show also a very low transcription rate of both $ER\alpha$ and $ER\beta$ in rat brain. From these data, we can speculate that in male rats, $ER\beta$ is involved in the control of the complex mental functions of the cerebral cortex, whereas $ER\alpha$ may have a role in the hippocampus activities, such as learning and long-term memory. This hypothesis is corroborated by the findings showing that the learning performance in $ER\beta$ -KO mice was higher to that observed in $ER\alpha$ -KO, indicating that removal of $ER\beta$ preserves the learning abilities [44].

Many studies reported age-related changes in the transcript level of several genes in the mammalian nervous system [45,46]. We found that the relative abundance of $ER\alpha$ and $ER\beta$ transcripts changes during aging in both

cortex and hippocampus. In the cortex, we observed a general and significant decrease in *ER* content from 2- to 16-month-old rats, and these findings are in agreement with that found in the cerebral cortex of mice, where *ER* α and *ER* β transcription rate decreases in males with increasing age [42]. The *ER* expression trend in the hippocampus is different, as with aging *ER* α is upregulated, while *ER* β is down-regulated. In particular, in the male 4-month-old rats, the level of ERs does not change significantly compared to 2 month-old rats, but in adult (8-month) and middle-aged (16-month) rats, the *ER* α expression considerably increases by about 5 and 25 times, and the *ER* β decreases by about 1/5 and 1/2, respectively.

On the basis of the current knowledge, the *ER* downregulation observed in the cortical area could be linked to a neuroprotective effect [47]. It is known that Folstein test scores in women in AD terminal stage show an increase in *ER* α levels in the frontal cortex [48], and that estrogen and ER agonists are capable of altering the transcription of genes involved in neuroinflammatory responses in astroglia, interneurons and microglia of the frontal cortex [49–51].

Also in the hippocampus, the observed $ER\alpha$ upregulation could have a neuroprotective role against the onset of neurodegenerative diseases, contributing to the effectiveness of the hormonal therapy ameliorating the memory loss that often accompanies advanced age [52]. In transgenic mice and in humans, it has been demonstrated that cytoplasmatic $ER\alpha$ inhibits the development of AD pathology [53,54], whereas $ER\beta$ increased in the hippocampal cells of AD patients [28]. Moreover, the expression of $ER\alpha$ seems to restore the protective action of estrogen against amyloid- β peptide, supporting the key role of $ER\alpha$ in estrogen-mediated neuroprotection [55].

Because $ER\beta$ may act as a negative regulator of $ER\alpha$ mediated transcription [7,22], a general low level of $ER\beta$ transcripts during aging in brain is not surprising.

Another objective of this work was to evaluate the expression of $ER\alpha$ and $ER\beta$ in brain of rats fed with an HFD for 4 and 12 weeks, a condition that is able to induce systemic and peripheral inflammation, insulin resistance, hepatic steatosis, higher serum glucose and leptin levels [56], as well as inflammation and decrease in the haptoglobin level in the brain [38]. Once again, from our data emerges a different response in the two brain regions. In cortex, the 4-week HFD treatment affects only $ER\beta$ expression, which significantly decreased; conversely, the 12 weeks of HFD induced a huge increase (by about 100-fold) in both $ER\alpha$ and $ER\beta$ expression. In the hippocampus, the $ER\beta$ transcripts level did not change following HFD, whereas $ER\alpha$ transcript level showed a two-fold increase after 4 weeks of treatment and a halving after 12 weeks.

Evidence exists that circulating estrogen levels are able to induce the expression of leptin receptors, probably through estrogen responsive elements present in the gene promoter region [57]. A crosstalk between ER and leptin receptors has also been evidenced [58–60]. For this reason, the weight gain observed in postmenopausal women is often associated with the decrease in estrogen levels occurring in this period. It has also been demonstrated that estrogen is involved in the regulation of ghrelin expression, so the *ER* down-regulation may jointly cause a reduced response to leptin and an increased susceptibility to ghrelin.

Further, it is interesting to note that obesity increases the risk of age-related cognitive decline and is accompanied by peripheral inflammation [61–63]. In particular, dietary obesity impaired hippocampus-dependent memory and reduced long-term potentiation [63].

In this scenario, it is conceivable that the huge increase in *ER* expression observed in the cortex of rats fed for 12 weeks with a hypercaloric diet may represent a defense mechanism to counterbalancing the excessive energy intake, increasing sensitivity to leptin and reducing hunger. Conversely, the lack of *ER* responsiveness to HFD in hippocampus might be one of the motives for which the dietary composition negatively affects hippocampal functions, making this brain area, crucial for learning and memory, one of the most vulnerable sites in inflammation, in early AD and in other neurodegenerative diseases development.

Taken together, our data add a new slice to identify factors that significantly change in the brain following aging and HFD administration and unveil another possible molecular mechanism involved in the brain consequences of dietary fats.

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