

Early redox imbalance is associated with liver dysfunction at weaning in overfed rats

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Key points

- Childhood obesity is associated with precocious oxidative stress, which can contribute to future diseases.
- Early overfed rats have increased oxidative stress in liver and plasma at weaning.
- The model of litter size reduction causes adipocyte hypertrophy and lower PPAR γ at weaning, suggesting a decrease in adipocyte proliferation.

Abstract Neonatal overfeeding induced by litter size reduction leads to further obesity and other metabolic disorders, such as liver oxidative stress and microsteatosis at adulthood. We hypothesized that overfeeding causes an early redox imbalance at weaning, which could programme the animals to future liver dysfunction. Thus, we studied lipogenesis, adipogenesis, catecholamine status and oxidative balance in weaned overfed pups. To induce early overfeeding, litters were adjusted to three pups at the 3rd day of lactation (SL group). The control group contained 10 pups per litter until weaning (NL group). Peripheral autonomic nerve function was determined *in vivo* at 21 days old. Thereafter, pups were killed for further analysis. Differences were considered significant when $P < 0.05$. The SL pups presented with a higher visceral adipocyte area, higher content of lipogenic enzymes (ACC, FAS) and with a lower content of adipogenic factors (CEBP, PPAR γ) in visceral adipose tissue (VAT). Although autonomic nerve activity and adrenal catecholamine production were not significantly altered, catecholamine receptor (β 3ADR) content was lower in VAT. The SL pups also presented with higher triglyceride, PPAR γ , PPAR α and PGC1 α contents in liver. In plasma and liver, the SL pups showed an oxidative imbalance, with higher lipid peroxidation and protein oxidation. The SL group presented with a higher serum alanine aminotransferase (ALT). The early increase in lipogenesis in adipose tissue and liver in weaned overfed rats suggests that the higher oxidative stress and lower catecholamine content in VAT are associated with the early development of liver dysfunction and adipocyte hypertrophy.

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Abbreviations ACC, acetyl-CoA carboxylase; β 2ADR, β 2 adrenergic receptor; β 3ADR, β 3 adrenergic receptor; ALT, alanine aminotransferase; AST, aspartate aminotransferase; CAT, catalase; C/EBPs, CCAAT-enhancer-binding proteins; FAS, fatty acid synthase; GPx, glutathione peroxidase; MDA, malondialdehyde; NL, normal litter; PGC1 α , peroxisome proliferator-activated receptor-gamma coactivator 1 α ; PPAR γ , nuclear receptor peroxisome proliferator-activated receptor γ ; ROS, reactive oxygen species; SL, small litter; SOD, superoxide dismutase; TG, triglyceride; TH, tyrosine hydroxylase; VAT, visceral adipose tissue.

Introduction

Obesity is an epidemic health problem (Ng *et al.* 2014) associated with metabolic syndrome, which is characterized by central obesity, dyslipidaemia, type 2 diabetes, cardiovascular diseases, and non-alcoholic fatty liver disease (Shin *et al.* 2013). Childhood obesity is especially concerning because it causes precocious metabolic syndrome (Ozturk & Soylu, 2014) and high rates of long-standing obesity (Tounian, 2011). Currently it is known that disturbances arising from the obesity are influenced by pro-oxidative conditions (Codoñer-Franch *et al.* 2011; Aroor & DeMarco, 2014).

Experimental studies have shown that precocious nutritional interventions are able to modulate tissue-specific endocrine and physiological functions, as an adaptation mechanism. This physiological phenomenon is named ontogenetic plasticity or metabolic programming (de Moura & Passos, 2005; Hanson & Gluckman, 2014). It has been observed in overnutrition during lactation by litter size reduction, leading to a higher milk offer (Cunha *et al.* 2009) and long-term obesity and their comorbidities (Habbout *et al.* 2013*b*). This precocious obesity increases serum triacylglycerol, visceral and total fat body content and produces high serum insulin at weaning (Rodrigues *et al.* 2011).

The development of obesity is related to adipocyte hypertrophy and ectopic lipid accumulation. The adipocyte differentiation and lipid storage is stimulated by specific transcription factors, such as CCAAT-enhancer-binding proteins (C/EBPs) and a nuclear receptor, peroxisome proliferator-activated receptor gamma (PPAR γ) (Ali *et al.* 2013). Another factor involved in lipid metabolism is the peroxisome proliferator-activated receptor alpha (PPAR α); its reduced activation is related to reduction of fatty acid oxidation and increased lipid accumulation (Gao *et al.* 2015). It has been demonstrated that the proliferation capacity of adipose tissue from obese mammals can reach a limit, and the hypertrophied adipocytes release high amounts of free fatty acids, which are stored in the liver (Ameer *et al.* 2014). Donnelly and colleagues (2005) showed that patients with liver steatosis present with 59% of triglycerides in the livers originating from adipocyte lipolysis, 26% from *de novo* lipogenesis and only 15% from the diet. This suggests that a dysfunction of the visceral adipose tissue may be the first step to liver steatosis. In the liver, *de novo* lipogenesis is stimulated by PPAR γ expression, which overexpression is related to lipid accumulation (Sanders & Griffin, 2015). Acetyl CoA carboxylase (ACC) and fatty acid synthase (FAS) are essential enzymes for lipogenesis; the first enzyme catalyses the malonyl-CoA synthesis while the second enzyme catalyses the synthesis of long chains fatty acid from acetyl-CoA and malonyl-CoA (Sanders & Griffin, 2015). Adipocyte hypertrophy and

liver lipid accumulation are associated with increased expression and activity of these enzymes in both tissues (Sanders & Griffin, 2015).

The increase in lipid influx into the liver from visceral adipocytes might be favoured by increased activity of the catecholaminergic system, leading to higher sympathetic noradrenaline (norepinephrine) or adrenal medullar adrenaline release, or through increased catecholamine sensitivity in adipocytes (Nielsen *et al.* 2014). Similarly, liver lipid storage might be favoured by reduced catecholamine sensitivity in this tissue. Since the model of litter size reduction leads to an adrenal catecholamine dysfunction at adulthood (Conceição *et al.* 2013*b*), it is important to identify whether this disruption occurs early in life, contributing to the long-term metabolic disturbances found in this programming model of obesity.

Some studies associate metabolic illness, such as obesity and their comorbidities, with an increase in oxidative stress (Bullon *et al.* 2000; Tariq *et al.* 2014). However, there are few reports showing the close association between neonatal oxidative stress and the development of diseases at adulthood (Holloway *et al.* 2005; Bruin *et al.* 2008; Niu *et al.* 2013).

Habbout and colleagues (2012 and 2013*a*) showed that increased plasma and cardiac oxidative stress were associated with higher susceptibility to cardiac injury at adulthood. Our group demonstrated that rats that were early overfed due to reducing litter size presented with long-term obesity, hyperphagia, hepatic microsteatosis and oxidative stress at adulthood (Conceição *et al.* 2013*a*). However, there are no reports on whether the early obesity induced by litter size reduction is related to precocious oxidative imbalance. Thus, the aim of the present study is to determine if early postnatal overnutrition is able to induce fat accumulation and oxidative stress in liver and adipocytes during the weaning period. We also evaluated visceral adipocyte morphology, the expression of adipogenic/lipogenic proteins in visceral adipose tissue and/or liver and autonomic nervous system activity and catecholamine status in 21-day-old rats.

Methods

Experimental model

Our protocol was approved by the Animal Care and Use Committee of the Biology Institute of the State University of Rio de Janeiro. The Wistar rats employed in the experiment were housed under controlled temperature ($23 \pm 3^\circ\text{C}$), light (12 h light–dark cycle), and had free access to water and food. Twenty nulliparous female Wistar rats were placed with (10) male rats (both groups 120 days old) at a 2-to-1 ratio for 5 days. After the

mating, pregnant females were housed in individual cages until delivery. After birth, all litters were adjusted to 10 male pups for each dam. Female pups were substituted by male pups from other litters. To induce early over-feeding (SL group, $n = 10$), at the third day of life, the litter size was reduced to three male pups per dam. The control group (NL group, $n = 10$) was kept with ten pups per dam until weaning at post-natal day 21 (PN21). We carried out the experiments with male pups only because it had already been demonstrated that there are no gender differences at weaning with this model of litter size reduction, especially in regard of body weight, fat mass, lean mass, leptinaemia, energy expenditure and respiratory exchange ratio (Stefanidis & Spencer, 2012).

At weaning, the pups were fasted for 2 h and then anesthetized (pentobarbital sodium, 90 mg (kg body weight)⁻¹) to allow *in vivo* autonomic nerve activity assessment. After the experiments, the pups were killed by exsanguination. Visceral fat tissue (epididymal, mesenteric and retroperitoneal depots) was dissected and weighed. Blood samples were centrifuged (1000 × g, 4°C, 20 min) to collect the plasma and stored (-20°C) for later analysis. Liver and visceral adipose tissue (VAT) was collected for further analysis.

Autonomic nerve activity assessment

To evaluate the autonomic activity, a parasympathetic nerve, the left superior branch of the upper vagus nerve, and a sympathetic nerve, the left intrascapular nerve, were exposed under a dissection microscope. The branches were placed on a pair of hook platinum electrodes connected to an electronic device (Bio-Amplificator, Insight, Ribeirão Preto, SP, Brazil) to record electrical signals. The nerve activity was amplified (10,000) and filtered (1–60 kHz). Data were analysed using Insight software (Insight). All nerve acquisition was processed inside a Faraday cage to avoid electromagnetic interference and the rats were kept under a warming light. After 10 min of stabilization, the average of number of spikes during 15 min was counted and the nerve firing rate was expressed as number of spikes (5 s)⁻¹ (Scomparin *et al.* 2009).

Morphological evaluation of the adipose tissue

Adipose tissue samples were fixed in formaldehyde 0.1 M phosphate-buffered saline (pH 7.2). The tissues were then dehydrated, cleared and paraffin-embedded. Non-consecutive slices 10 μm thick were obtained and stained with Haematoxylin and Eosin to assess morphology. Digital images were obtained from histological laminas with the Olympus BX40 microscope (Olympus, Tokyo, Japan) using a ×40 objective. Three laminas were obtained from each rat, and 10 pictures of

different fields were taken from each lamina. From each picture, the area of five adipocytes was measured using the software Image-Pro Plus 5.0 (Media Cybernetics, Silver Spring, MD, USA) (Conceição *et al.* 2011).

Triglyceride content

Triglyceride (TG) content was determined by colorimetric assay after total liver lipid content extraction with isopropanol (Vetec, Duque de Caxias, RJ, Brazil) and centrifugation (740 × g, 10 min, 4°C) (Folch *et al.* 1957). The TG content in the supernatant was evaluated using a commercial kit (Bioclin, BH, MG, Brazil), and the absorbance was measured at 500 nm (Hidex, Turku, Finland).

Western blotting analysis

The protein content of peroxisome proliferator-activated receptor α and γ (PPARα and PPARγ; ab8934 from Abcam, Cambridge, MA, USA and sc-7273 from Santa Cruz Biotechnology Inc., Dallas, Texas, USA. Wembley, UK, respectively), peroxisome proliferator-activated receptor-gamma coactivator 1 α (PGC1; Santa Cruz, sc-13067), acetyl-CoA carboxylase (ACC; Santa Cruz, sc-26817), fatty acid synthase (FAS; Abcam, ab150508), β2 adrenergic receptor (β2ADR; Abcam, ab36956), β3 adrenergic receptor (β3ADR; Santa Cruz, sc-50436), tyrosine hydroxylase (TH; Sigma-Aldrich Inc., MO, USA, T2928), superoxide dismutase 1 and 2 (SOD1 and SOD2; Santa Cruz, sc-11407 and sc-133134, respectively), glutathione peroxidase (GPx; Santa Cruz, sc-133152), catalase (CAT; Sigma-Aldrich, C0979), 4-hydroxynonenal protein adduct (4-HNE; Santa Cruz, sc-130083), and 3-nitrotyrosin (Santa Cruz, sc-55256) were evaluated by Western blotting in liver and VAT. Briefly, tissues were homogenized in RIPA buffer (50 mM Tris-HCl (pH 7.4), 1% NP-40, 150 mM NaCl, 1 mM EDTA, 1 mM PMSE, 1 mM Na₃VO₄, 1 mM NaF), and protease inhibitor cocktail (F. Hoffmann-La Roche Ltd, Basel, Switzerland). Homogenates were centrifuged for 5 min (1,120 × g, 4°C). The protein concentration in the supernatants was determined using the Pierce BCA Protein Assay Kit (Thermo Scientific, Carlsbad, CA, USA) and aliquots of homogenate containing 10 mg total protein were then analysed by SDS-PAGE. Samples were electroblotted onto nitrocellulose membranes (Hybond ECL; Amersham Pharmacia Biotech, Amersham, London, UK). Membranes were incubated with Tris-buffered saline (TBS) containing 2% albumin for 90 min. Membranes were then washed with TBS and incubated with a specific primary antibody (dilution 1:500) overnight at 4°C, after which they were washed and incubated with secondary antibody diluted to 1:5000 (anti-mouse,

B8520 from Sigma-Aldrich; anti-goat, 62–6540, and anti-rabbit, 65–6140, from Invitrogen, Carlsbad, CA, USA) and then conjugated with HRP at an adequate dilution for 1 h at room temperature. The protein bands were visualized by chemiluminescence (Kit ECL plus, Amersham Biosciences, London, UK) followed by exposure to ImageQuant LAS (GE Healthcare, Buckinghamshire, Chalfont St Giles, UK). Area and density of the bands were quantified by Image J software (Wayne Rasband National Institute of Health, Bethesda, MA, USA). Results were expressed relative (%) to the control group.

Reverse transcription polymerase chain reaction (RT-PCR) analysis

Total RNA was extracted from 100 mg of VAT, under RNase-free conditions, with TRIzol Reagent (Invitrogen), and quantified via Nano Vue™ Plus Spectrophotometer (GE Healthcare, Buckinghamshire, UK). The cDNA was prepared from total RNA using the Moloney murine leukemia virus reverse transcriptase (M-MLV RT) for RT-PCR and oligo(dT)15 primer (Promega, WI, USA). The mRNA levels of C/EBP β (forward (F) primer: 5'-ATGCAATCCGGATCAAACGT-3'; reverse (R) primer: 5'-CCGCAGGAACATCTTTAAGTGA-3') and PPAR γ (forward (F) primer: 5'-ATTCTGGCCCACTTCGG-3'; reverse (R) primer: 5'-TGGAAGCCTGATGCTTTATCCCCA-3') in visceral fat mass were amplified on Applied Biosystems 7500 Real-Time PCR System (Life Technologies Co. Carlsbad, CA, USA) using SYBR Green PCR Master Mix (Applied Biosystems, Foster City, CA, USA) according to the recommendations of the manufacturer. The oligonucleotide primers and probes were designed and prepared by ABI Applied Biosystems. The co-amplification of mouse 36 β -4 Mrna (F primer: 5'-TGTTTGACAACGGCAGCATTT-3'; R primer: 5'-CCGAGGCAACAGTTGGGT A-3'), a variant internal control, was performed in all of the samples. Assays were performed in triplicate, and the results were normalized to the 36 β -4 mRNA levels using the $2^{-\Delta\Delta CT}$ method (Livak & Schmittgen, 2001).

Adrenal catecholamine measurement

Total catecholamines were quantified by the trihydroxyindole method (Conceição *et al.* 2013b). Right adrenal glands were homogenized in 10% acetic acid and centrifuged (1120 \times g, 5 min). Briefly, 50 μ l of adrenaline standard and the adrenal supernatant were mixed with 250 μ l of buffer phosphate (0.5 M, pH 7.0) and 25 μ l of potassium ferricyanate (0.5%), and incubated for 20 min

in an ice bath. The reaction was stopped with 500 μ l of ascorbic acid (60 mg ml⁻¹)–NaOH (5 N) solution, and diluted with 2 ml of water distilled. The fluorescence was determined at 420 nm to excitation and 510 nm to emission (Hidex, Turku, Finland).

Antioxidant enzymes activities

Liver aliquots (200 mg) were homogenized in potassium phosphate buffer with EDTA. After centrifugation (3398 \times g, 10 min, 4°C), homogenates were stored at –80°C until each oxidative parameters assay. The total protein content was determined using the Pierce BCA Protein Assay Kit (Thermo Scientific, CA, USA). Total superoxide dismutase (SOD) activity was assayed by measuring the inhibition of adrenaline auto-oxidation as absorbance at 480 nm (Bannister & Calabrese, 1987). Glutathione peroxidase (GPx) activity was evaluated according to Flohé & Günzler (1984) by measuring the oxidation of NADPH in the presence of H₂O₂ at 340 nm. Catalase (CAT) activity was measured by the rate of decrease in H₂O₂ at 240 nm according to the Aebi method (1984). Enzymatic activities were expressed as U (mg of protein)⁻¹.

The SOD/GPx and SOD/(GPx+CAT) ratio activities were calculated, since previous data show that the ratio of the activities of these enzymes is more important than the absolute activity and is related to cellular oxidative status (Amstad *et al.* 1991; de Haan *et al.* 1996).

Antioxidant capacity assay

The antioxidant capacity of the sample was measured by the reduction of the DPPH^{*} (2,2-diphenyl-1-picrylhydrazyl) radical at 517 nm (Amarowicz *et al.* 2000). Plasma or liver homogenate samples (400 μ l) were mixed with 400 μ l of DPPH^{*} (0.25 mM) in ethanol solution and incubated (30 min) at darkroom and room temperature. They were then centrifuged (212 \times g, 5 min, 4°C), and the supernatant absorbance (Abs) was read in spectrophotometer (517 nm). The sample scavenging capacity for DPPH^{*} free radicals was evaluated by the following equation:

$$AA\% = 100 - \{[(Abs_{\text{sample}} - Abs_{\text{blank}}) \times 100] / Abs_{\text{control}}\},$$

where AA is antioxidant activity. The results were expressed as percentage DPPH^{*} scavenging.

MDA content

The lipid peroxidation was estimated by quantification of malondialdehyde (MDA) using the thiobarbituric acid reactive substances method (Vynche, 1970). The liver was

homogenized in phosphate buffer with EDTA. The liver homogenate was mixed with 10% trichloroacetic acid (1:2) and centrifuged ($212 \times g$, 10 min, 4°C). After this, $500 \mu\text{l}$ of supernatant was incubated with $500 \mu\text{l}$ of thiobarbituric acid (0.67%) for 30 min (95°C). The absorbance was read in a spectrophotometer (532 nm). The results were expressed as $\mu\text{mol} (\text{mg protein})^{-1}$.

Plasma transaminases

The plasma activities of alanine aminotransferase (ALT) and aspartate aminotransferase (AST) were quantified by the Reitman & Frankel method (1957) using colorimetric kits (Doles, Goiânia, GO, Brazil).

Statistical analysis

The statistical analyses were carried out using Graph Pad Prism 5.0 for Windows statistical software (GraphPad Software, La Jolla, CA, USA). Comparisons between the groups were performed with Student's unpaired *t* test. For all analyses, the data were given as means and standard error of the mean (SEM) and were considered significant when $P < 0.05$.

Results

As expected, the SL group presented with higher body mass from PN07 until PN21 (PN21: NL = 35.3 ± 0.5 g vs. SL = 44.3 ± 1.0 g; $P < 0.0001$), higher visceral fat mass (PN21: NL = 180.0 ± 17.0 mg vs. SL = 390.0 ± 34.0 mg; $P < 0.0001$) and higher relative visceral fat mass (PN21: NL = 4.76 ± 0.3 vs. SL = 8.25 ± 0.6 ; mg tissue (g body weight) $^{-1}$), $P < 0.0001$) compared with NL group. These pups also had a higher adipocyte area (+148%; $P < 0.05$; Fig. 1).

In VAT, we detected lower adipogenic transcription factor C/EBP β mRNA (-57% ; $P = 0.0082$; Fig. 2A), lower PPAR γ mRNA (-62% ; $P = 0.0003$; Fig. 2B) and lower PPAR γ protein content (-29% ; $P = 0.025$; Fig. 2C) in SL pups compared with the NL pups. The lipogenic enzymes ACC and FAS were higher ($+252\%$; $P = 0.01$; Fig. 2D and $+223\%$; $P = 0.002$; Fig. 2E, respectively) in VAT in the SL group.

At weaning, the liver mass of the SL group was not different from the NL group (Fig. 3A), although they presented with a higher triglyceride liver content ($+22\%$; $P < 0.0001$; Fig. 3B). Whereas PPAR γ content was higher ($+56\%$; $P = 0.034$; Fig. 3C) in SL pups, FAS content was unchanged (Fig. 3D). The lipolytic transcription factors, PPAR α and PGC1 α , were higher in SL group ($+98\%$; $P = 0.03$; Fig. 3E and $+82\%$; $P = 0.0004$; Fig. 3F, respectively) than in controls.

The parameters related to autonomic nervous function, catecholamine production and receptors in rats at weaning are depicted in Table 1. Both the sympathetic and the parasympathetic nerve activities were unaltered in the SL group. In the adrenal glands, neither the catecholamine content nor the tyrosine hydroxylase content was different

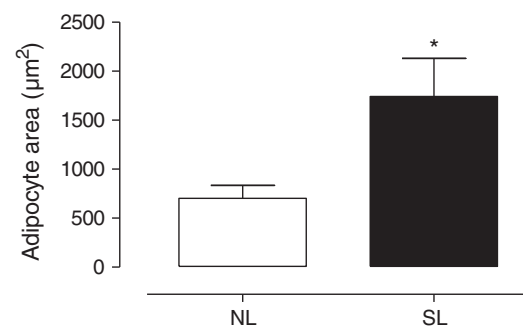
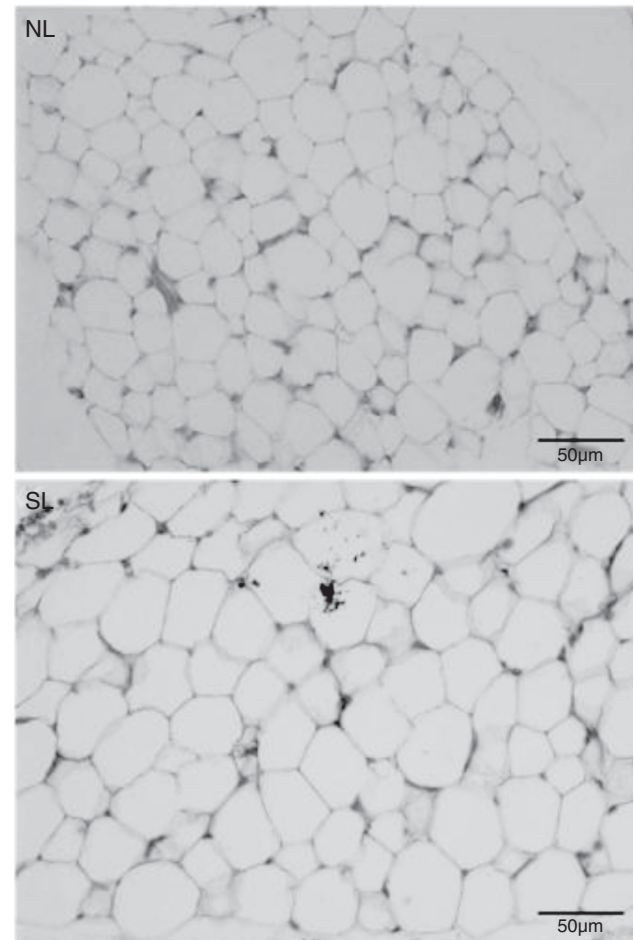


Figure 1. The effect of reducing litter size on adipocyte morphology at 21 days old

Representative photomicrography of the visceral adipocyte from rats raised in normal litters (NL) and small litters (SL) during lactation. The quantitative analysis of the adipocyte area is shown below. Values are expressed as means \pm SEM; $n = 4$ per group; * $P < 0.05$.

between groups. Regarding the adrenergic receptor analysis, the $\beta 3$ ADR content was lower in VAT in SL pups than in NL pups, while the $\beta 2$ ADR content in liver was unchanged.

In the liver, SOD1, SOD2 and CAT contents were increased but the GPx content was decreased (Fig. 4A). In liver, SOD and CAT activities were unchanged, and GPx activity was decreased in SL group (-33% ; $P = 0.04$; Table 2). The total antioxidant capacity assayed as the DPPH^{*} scavenging capacity was decreased (-50% ; $P = 0.03$; Table 2). The ratio between the antioxidant enzyme activities, SOD/GPx (Table 2) and SOD/(GPx+CAT) (Table 2), were unchanged. The MDA content and protein carbonyl content were increased in SL pups ($+77\%$, $P = 0.02$, and $+50\%$, $P = 0.02$, respectively, Table 2), as were the 4-HNE protein adducts and 3-nitrotyrosin liver content (Fig. 4B and C, respectively).

Regarding the plasma transaminase analysis, the SL group showed an increase in ALT activity ($+25\%$; $P = 0.006$; Fig. 5B) compared with the NL group, despite having normal AST activity (Fig. 5A).

In the plasma analysis (Table 2), SOD and GPx activities were unaltered (Table 2) in SL rats, while CAT activity was lower than controls (-42% ; $P = 0.04$). These animals also had higher SOD/GPx ($+126\%$; $P = 0.006$) and SOD/(GPx+CAT) ($+106\%$; $P = 0.01$), indicating an increased cellular oxidative condition. Both MDA and protein carbonyl contents were increased in the SL group compared with the NL group ($+130\%$, $P = 0.02$, and $+162\%$, $P = 0.01$, respectively).

Discussion

As expected, at weaning the SL pups showed higher body weight and visceral adipose tissue. The increases in central adiposity and adipocyte hypertrophy were associated with a higher content of the lipogenic enzymes ACC, which converts acetyl-CoA to malonil-CoA, and FAS, which catalyses the synthesis of long chain fatty acid from acetyl-CoA and malonil-CoA, in the VAT of SL pups (Ameer *et al.* 2014). Aubert and colleagues (1980) showed that litter size reduction caused adipocyte

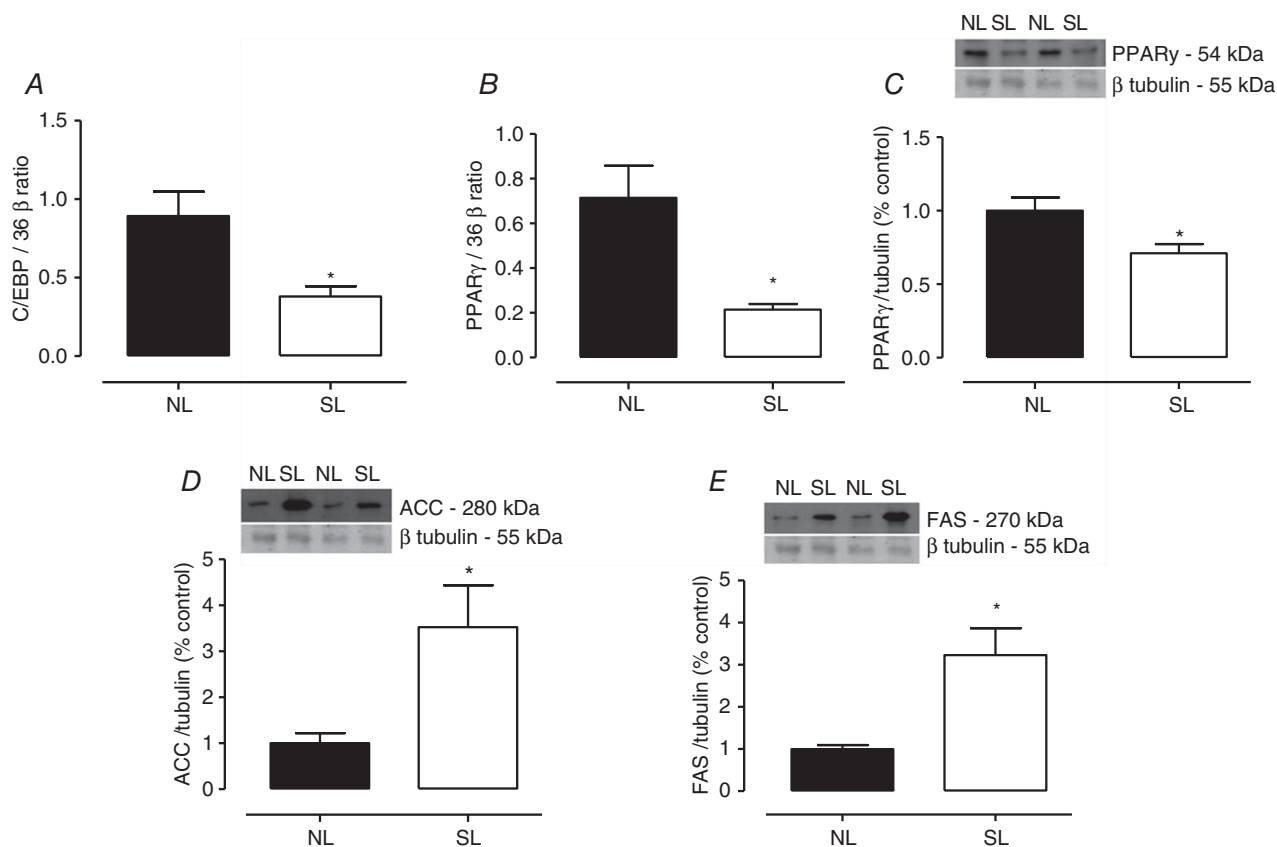


Figure 2. The effect of reducing litter size on adipogenic and lipogenic biomarkers in visceral adipose tissue at 21 days old

RNA level of C/EBP β (A) and PPAR γ (B), protein content of PPAR γ (C), ACC (D) and FAS (E) in visceral adipose tissue from rats raised in normal litters (NL) and small litters (SL) during lactation. The representative blots of proteins are shown above graphs C, D and E. β -Tubulin content was used as control loading. Results are expressed as relative (%) to the control group and as means \pm SEM; $n = 4$ –7 per group; * $P < 0.05$. The PCR analyses were normalized to the 36 β mRNA level.

hypertrophy and hyperplasia in adult mice. However, the present study shows that both the mRNA level and protein content of the adipogenic factors *C/EBPβ* and *PPARγ* were lower in visceral adipose tissue from

the SL group. In agreement, Akyürek and colleagues (2013) reported lower serum *PPARγ* levels in obese 8- to 12-year-old children. The maternal obesity model also is associated with lower *PPARγ* levels in visceral

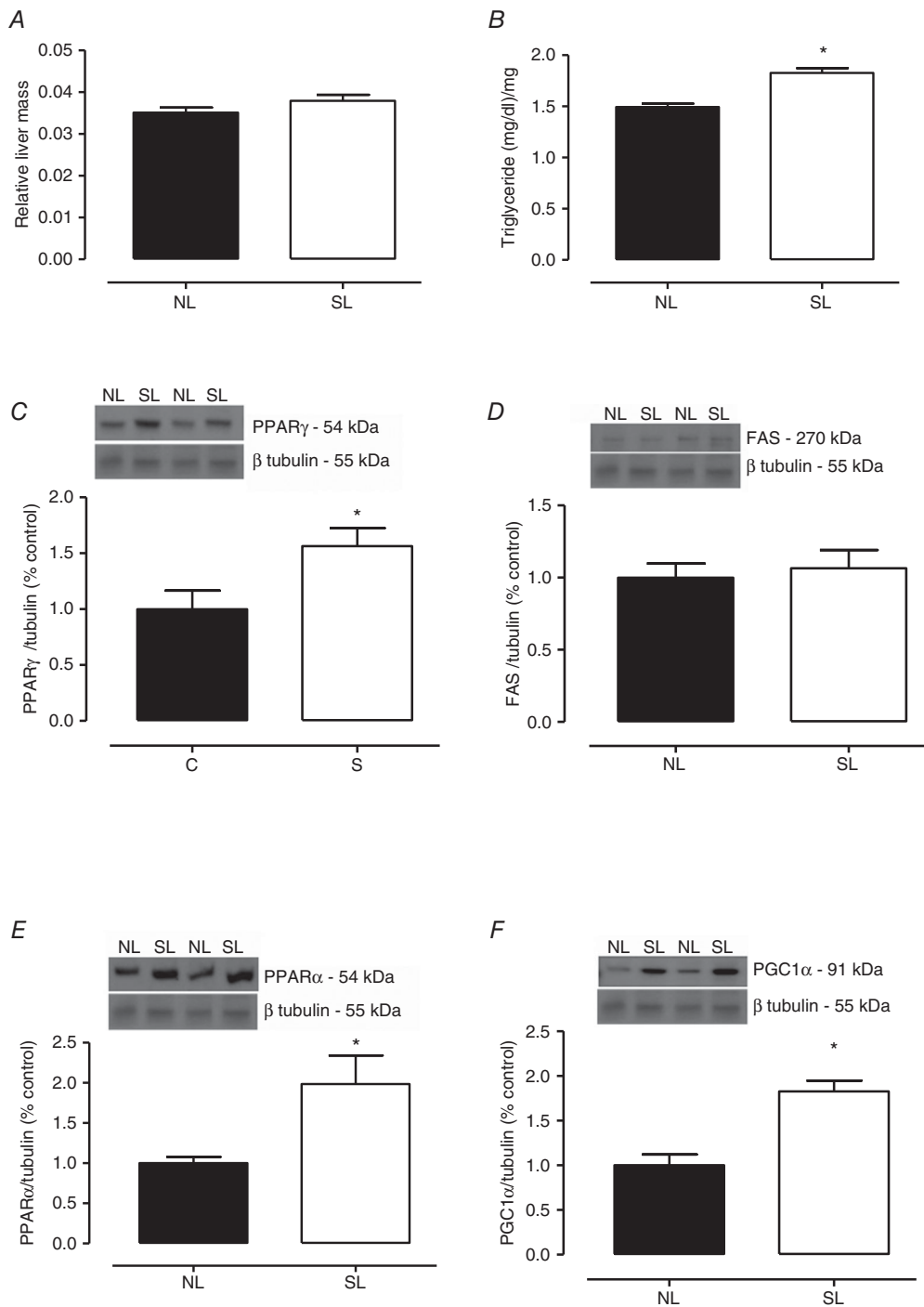


Figure 3. The effect of reducing litter size on lipid metabolism in liver at 21 days old
 Relative mass (A), triglyceride content (B), *PPARγ* (C), *FAS* (D), *PPARα* (E) and *PGC1α* (F) protein contents from rats raised in normal litters (NL) and small litters (SL) during lactation. The representative blots of proteins are shown above graphs C, D, E and F. β -Tubulin content was used as control loading. Results are expressed as relative (%) to the control group and as means \pm SEM; $n = 6-10$ per group; * $P < 0.05$.

Table 1. Effect of reducing litter size on parameters related to autonomic nervous, catecholamine production and receptors in rats at PN21

	NL	SL	P
Sympathetic nerve activity (spikes (5 s) ⁻¹)	23.14 ± 2.9	20.53 ± 3.2	0.55
Parasympathetic nerve activity (spikes (5 s) ⁻¹)	18.13 ± 2.1	18.80 ± 2.3	0.83
Catecholamine content in adrenal (μM mg ⁻¹)	5.78 ± 1.4	8.36 ± 1.4	0.20
TH content in adrenal medulla (% of control)	1.00 ± 0.1	0.90 ± 0.1	0.70
β3ADR content in VAT (% of control)	1.00 ± 0.1	0.55 ± 0.1*	0.01
β2ADR content in liver (% of control)	1.00 ± 0.2	1.17 ± 0.3	0.58

Data are presented as means ± SEM of 5–10 pups for NL and SL groups with level of significance (*) set at $P < 0.05$. NL, normal litter; SL, small litter; TH, tyrosine hydroxylase; β3ADR, β3 adrenergic receptor; β2ADR, β2 adrenergic receptor; VAT, visceral adipose tissue.

adipose tissue of adult mice offspring (Magliano *et al.* 2013). PPARγ activation causes mature adipocyte apoptosis and increases adipocyte proliferation, which is more sensitive to insulin anti-lipolytic action (Okuno *et al.* 1998). However, Akyürek *et al.* (2013) and Magliano *et al.* (2013) support the conclusion that PPARγ is not necessary for mature adipocyte function in obesity. Apparently, increased lipogenesis in the adipocytes is associated with a reduction in PPARγ and C/EBP in the SL group, which suggests that the adipocyte storage capacity reaches a limit, because there is less stimulus for adipocyte proliferation, favouring adipocyte hypertrophy and hypertriglyceridaemia (Rodrigues *et al.* 2011), and ectopic lipid storage, as observed in the higher triglyceride content in the liver.

The liver triglyceride content was increased in SL rats, followed by increased PPARγ protein content. The association between liver lipid accumulation and increases in PPARγ was previously reported in an experimental model of obesity and diabetes (Satoh *et al.* 2013; Alfaradhi *et al.* 2014). The precocious triglyceride accumulation may be due to increased *de novo* hepatic lipogenesis, since the increase in milk intake in the litter size reduction model provides elevated amounts of carbohydrate for fatty acid synthesis (Moreira *et al.* 2009; Cunha *et al.* 2009). The lipolytic factors PPARα and PGC1 were increased and the FAS content was unchanged in the SL group, suggesting increased oxidative metabolism in the liver. These results have already been reported (Lin *et al.* 2005); a high fat diet intake was able to increase the PGC-1β content in mice liver. These findings suggest that in short-term obesity the liver is able preferentially to use fatty acids as an energy source, which could decrease the contribution of glycolysis to the insulin resistance observed in this model at this age (Rodrigues *et al.* 2011).

Some studies show that autonomic dysfunction makes an important contribution to obesity development (Lima *et al.* 2007; Messina *et al.* 2013). Brown adipose tissue (BAT) has a thermogenic function, dissipating energy as heat, stimulated by sympathetic nerve activity. A reduction in sympathetic activity is reported in obesity models

(Lima *et al.* 2007; Scomparin *et al.* 2009). Here, we report evidence that the firing rates of BAT (sympathetic nerve) and vagus (parasympathetic nerve) nerves were not changed in the SL group at weaning. The long-term outcomes of reducing litter size upon the sympathetic and parasympathetic nerve activity have already been reported (de Almeida *et al.* 2013). At PN90, the SL rats showed an increase in vagal nerve superior branch activity, which was related to higher insulin secretion, but these SL rats did not show a change in BAT sympathetic nerve activation. In addition, the adrenal medulla catecholamine content and TH content were not affected by early post-natal obesity. These findings suggest that precocious onset of obesity may be not related to changes in autonomic peripheral nerve activity as demonstrated previously in other obese models. Scomparin and colleagues (2009) showed a reduction in sympathetic firing rate in MSG obese model adult rats. Besides alterations in firing rate activity, the sympathetic function can also be modulated by adrenergic tissue sensitivity (Lima *et al.* 2007). In the present study, SL rats had lower β3ADR in VAT, suggesting reduced adrenergic sensitivity, which leads to a decrease in lipolytic action and accords with higher central adiposity. In the liver, the SL rats showed unchanged β2ADR at weaning. Thus, the adipocyte changes can, at least in part, be explained by the reduction in adrenergic action, and not by changes seen in the liver. Curiously, on reaching adulthood (PN180) these SL animals had a lower β2ADR content in liver, but showed no changes in β3ADR content in VAT (Conceição *et al.* 2013b). These liver alterations are more remarkable, when we consider the higher liver steatosis in the adult SL animal. Thus, we suggest that the programming effect is seen first in the adipocytes, rather than in the liver, and this effect is aggravated during development.

Obesity and a high-fat diet are related to an increase in reactive oxygen species production (Roberts *et al.* 2006; Aroor & DeMarco, 2014). In our plasma analysis, SOD and GPx activities were unchanged, CAT activity was decreased and SOD/GPx and SOD/(GPx+CAT) ratios were increased in the SL group. The SOD/GPx

activity ratio is associated with higher production of hydrogen peroxide and lipid peroxidation (Amstad *et al.* 1991; de Haan *et al.* 1996). Accordingly, the lipid peroxidation and protein carbonyl content were increased in the plasma of the SL group; this oxidative condition in plasma has already been observed as an outcome of reducing litter size at adulthood (PN180 and PN210) (Conceição *et al.* 2013a; Habbout *et al.* 2012 and 2013b).

The common hepatic manifestation of metabolic syndrome is non-alcoholic steatosis. Its development involves increased free fatty acid influx and *de novo* lipogenesis followed by oxidative stress conditions (Rolo *et al.* 2012). Besides lipid accumulation in the liver, early overfeeding is able to induce oxidative stress at weaning. The SOD and CAT content in the liver was higher in our SL group, despite no change in their activities, and GPx content and activity were decreased. These

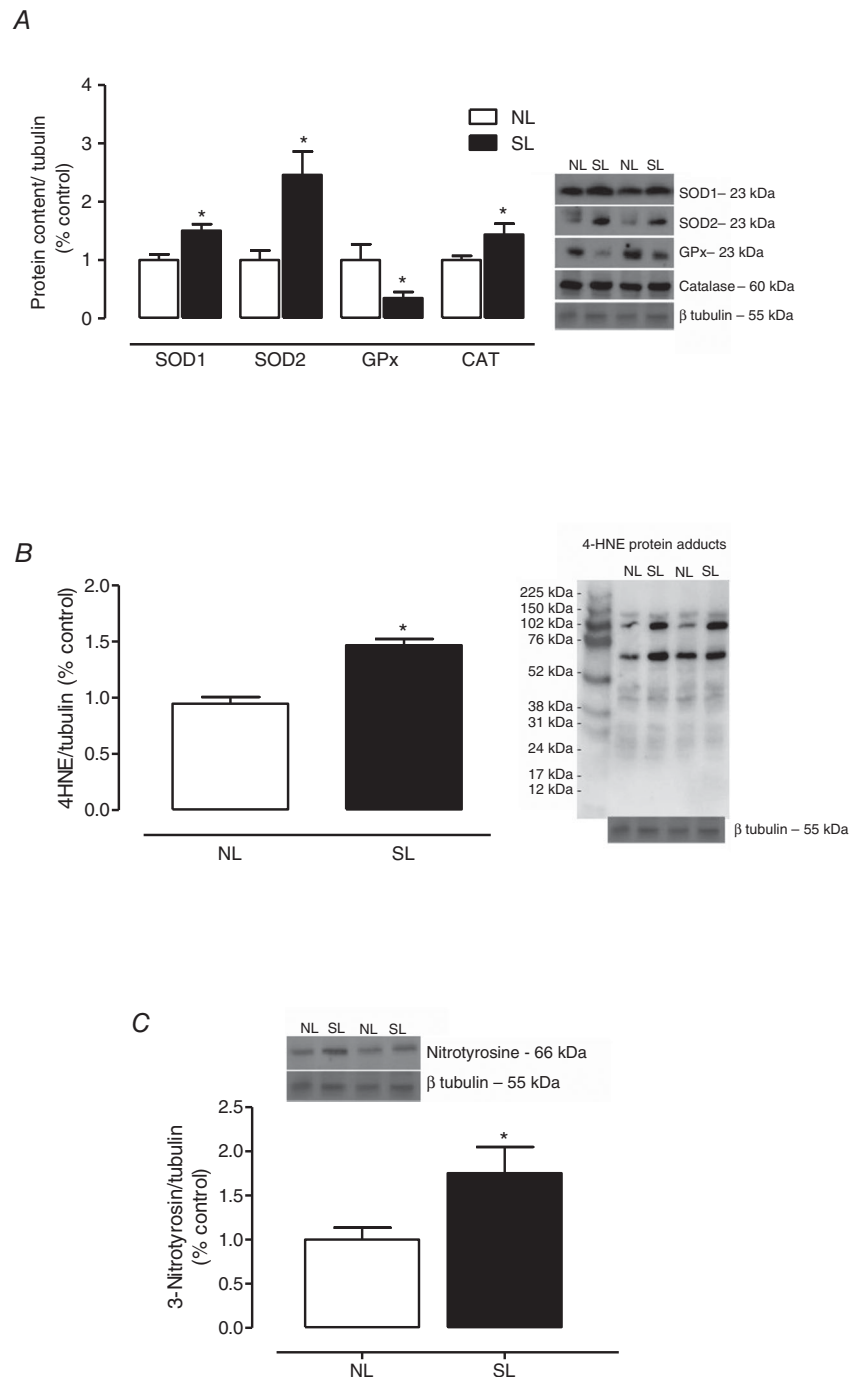


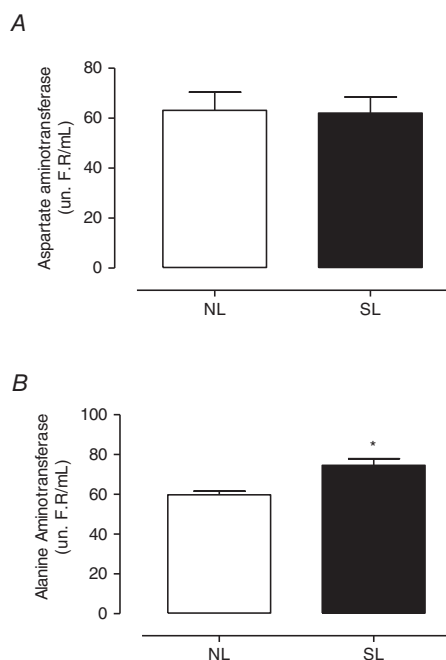
Figure 4. The effect of reducing litter size on antioxidant protein content and oxidative biomarkers in liver at 21 days old
 SOD1, SOD2, GPx and CAT protein contents (A), 4HNE protein adducts (B), and 3-nitrotyrosine content (C) from rats raised in normal litters (NL) and small litters (SL) during lactation. The representative blots of proteins are shown next to the graphs. β -Tubulin content was used as control loading. Results are expressed as relative (%) to the control group and as means \pm SEM; $n = 6-7$ per group; * $P < 0.05$.

Table 2. Effect of reducing litter size on oxidative parameters in liver and plasma of rats at PN21

		NL	SL	P
Liver	SOD (U (mg protein) ⁻¹)	2138 ± 280	1975 ± 222	0.67
	GPx (U (mg protein) ⁻¹)	0.332 ± 0.05	0.220 ± 0.01*	0.04
	CAT (U (mg protein) ⁻¹)	0.107 ± 0.01	0.104 ± 0.01	0.82
	SOD/GPx ratio	8187 ± 1708	8099 ± 757	0.96
	SOD/(GPx+CAT) ratio	5772 ± 1033	5493 ± 427	0.82
	DPPH [•] scavenging (%)	15.68 ± 2.1	7.81 ± 2.6*	0.03
	MDA (μmol mg ⁻¹)	73.0 ± 11.0	129.0 ± 17.0*	0.02
Plasma	Total protein bound carbonyl (nmol (mg protein) ⁻¹)	150.7 ± 14.0	226.7 ± 24.0*	0.02
	SOD (U (mg protein) ⁻¹)	1861 ± 203	1790 ± 191	0.8
	GPx (U (mg protein) ⁻¹)	0.037 ± 0.01	0.041 ± 0.01	0.72
	CAT (U (mg protein) ⁻¹)	0.048 ± 0.01	0.028 ± 0.01*	0.04
	SOD/GPx ratio	3316 ± 805	7496 ± 821*	0.006
	SOD/(GPx+CAT) ratio	3091 ± 781	6363 ± 697 *	0.01
	MDA plasma (μmol mg ⁻¹)	0.09 ± 0.01	0.22 ± 0.04*	0.02
	Total protein bound carbonyl (nmol (mg protein) ⁻¹)	310.4 ± 60.4	813.5 ± 140.9*	0.01

Data are presented as means ± SEM of 10 pups for NL and SL groups with level of significance (*) set at $P < 0.05$. SOD, superoxide dismutase; GPx, glutathione peroxidase; CAT, catalase; MDA, malondialdehyde; NL, normal litter; SL, small litter; DPPH[•], 2,2-diphenyl-1-picrylhydrazyl.

findings may contribute to the decrease in antioxidant capacity confirmed by a decrease in DPPH[•] scavenging activity, indicating oxidative damage in the liver, with

**Figure 5. The effect of reducing litter size on liver function enzymes at 21 days old**

Plasma activity of aspartate aminotransferase (AST; A) and alanine aminotransferase (ALT; B) from rats raised in normal litters (NL) and small litters (SL) during lactation. Results are expressed as means ± SEM; $n = 6-10$ per group; * $P < 0.05$. un R.F./mL refers to the Reitman Frankel unit used to quantify plasma activity of aspartate aminotransferase and alanine aminotransferase through the Reitman and Frankel method.

higher MDA content and 4-HNE protein adducts, which are parameters of lipid peroxidation, followed by carbonylated protein content and 3-nitrotyrosin, which are biomarkers of protein oxidation by free radical species. Furthermore the SL rats showed a higher plasma level of ALT activity. This finding is indicative of cellular liver damage (Rinella, 2015; Sanal, 2015), in agreement with our present data. Additionally, it is known that high ROS content is correlated to apoptosis (Bruin *et al.* 2008), immune system activation (Cohen *et al.* 2011) and epigenetic modifications (Milagro *et al.* 2012). This early disturbance may contribute to increased lipid accumulation at weaning and the other disturbances developed by this programming model at adulthood, such as microsteatosis, oxidative stress and insulin resistance in the liver (Conceição *et al.* 2013a).

Early obesity also may be result of an imbalance in formula feeding prescription, which may cause a high caloric intake. In fact both litter size reduction and formula feeding increase the body fat content and oxidative stress, but each condition has its own particular physiological profile. The lack of maternal hormones in formula feeding may prejudice normal adipose tissue and neuronal development (Melnik, 2014); formula feeding has no antioxidants which are important for adaptation to the hyperoxygenated environment after birth (Friel *et al.* 2011). This precocious exposure to a hyperoxidative status affects the physiological mechanism related to epigenetic modifications of important metabolism-related genes, such as the lipogenic, adipogenic and antioxidant genes, which can cause future diseases (Yara *et al.* 2015).

Our data help to elucidate the sequence of mechanisms by which early overfeeding induces both adipocyte

hypertrophy and liver dysfunction. The reduction in adrenergic sensitivity, despite an apparently normal sympathetic activation, may contribute to early adipocyte lipid accumulation. The postnatal obesity we observed was related to an increase in lipogenic capacity in adipose tissue and liver. Furthermore, the present results provide evidence that early overfeeding is able to induce the precocious onset of oxidative imbalance. This early exposure to a pro-oxidative condition might favour liver dysfunction and obesity. Thus further experimental studies must be designed to help us understand whether the early prevention of oxidative stress might be a good strategy to prevent future disease.

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Additional information

Competing interests

The authors declare that there is no competing interest that could be perceived as prejudicing the impartiality of the research reported.

Author contributions

Conception and design: E.P.S.C., E.O., E.G.M., P.C.L. Animal treatment, collection and measurements: E.P.S.C., J.C.C. Analysis and interpretation of data: E.P.S.C., E.O., J.C.C., E.G.M., P.C.L. Drafting and/or revising the article critically for important intellectual content: E.P.S.C., E.O., J.C.C., E.G.M., P.C.L. All authors read and approved the final version of the manuscript.

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