

Maternal conjugated linoleic acid modulates TAG metabolism in adult rat offspring

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Abstract

Conjugated linoleic acid (CLA) might regulate the lipid depots in liver and adipose tissue. As there is an association between maternal nutrition, fat depots and risk of offspring chronic disease, the aim was to investigate the effect of maternal CLA consumption on TAG regulation and some inflammatory parameters in adult male rat offspring receiving or not receiving CLA. Female Wistar rats were fed control (C) or CLA-supplemented (1%, w/w) diets during 4 weeks before and throughout pregnancy and lactation. After weaning, male offspring of CLA rats were fed C or CLA diets (CLA/C and CLA/CLA groups, respectively), whereas C male rat offspring were fed a C diet (C/C group) for 9 weeks. Serum TAG levels were increased in the CLA/CLA and CLA/C groups, associated with a reduction of lipoprotein lipase activity and weights of adipose tissue. The liver TAG levels were decreased in the CLA/CLA group, related to a significant reduction of fatty acid synthase (FAS), acetyl-CoA carboxylase (ACC) and glucose-6-phosphate dehydrogenase enzyme activities, as well as to the mRNA levels of FAS, ACC, stearoyl-CoA desaturase-1 and sterol regulatory element-binding protein-1c. Even though normal TAG levels were found in the liver of CLA/C rats, a reduction of lipogenesis was also observed. Thus, these results demonstrated a programming effect of CLA on the lipid metabolic pathways leading to a preventive effect on the TAG accretion in adipose tissue and the liver of male rat offspring. This knowledge could be important to develop some dietary strategies leading to a reduced incidence of obesity and fatty acid liver disease in humans.

Key words: Conjugated linoleic acid: Maternal programming: Offsprings: Lipid metabolism: TAG

Conjugated linoleic acid (CLA) refers to a group of dienoic derivatives of linoleic acid with conjugated double bonds arranged in different combinations of *cis* and *trans* configuration⁽¹⁾. The bioactive properties of *c9t11*-CLA and *t10c12*-CLA isomers have been extensively studied. CLA is incorporated into tissues⁽²⁾, increases energy expenditure, modulates TAG metabolism in liver, adipose tissue and skeletal muscle, modifies inflammatory markers and regulates oxidative status^(3,4). Studies in different experimental animal models have shown controversial results on hepatic TAG content. CLA might attenuate or prevent the development of hepatic steatosis in rats^(5,6), whereas a lipotrophic effect, characterised by a severe reduction of adipose tissue, elevated fat accretion in the liver and hepatomegaly has been described in mice⁽⁷⁾. The differential CLA response on TAG metabolism depends on the type of isomers, the level of CLA and fat in the diets and species⁽⁸⁾, among other factors.

The relationship between the intra-uterine environment and the propensity to lipid metabolic alterations in adult life was first noted in response to maternal food restriction⁽⁹⁾. Similarly, maternal high-fat diet consumption was associated with metabolic disorders including obesity⁽¹⁰⁾, hepatic steatosis⁽¹¹⁾, glucose intolerance and inflammation in later life⁽¹²⁾. Even though the type and amount of the maternal diet is very important, maternal body fat accumulation during early pregnancy plays a determining role in fetus development and propension or prevention of metabolic alterations in adult life⁽¹³⁾. There is limited information about the effects of compounds or agents, such as CLA, that by regulating body fat accretion could modulate the lipid metabolism in offspring animals. In this sense, maternal CLA supplementation in rats fed high-fat diets reverted some negative effects on inflammation, glucose intolerance, insulin insensitivity, hyperlipidaemia and other associated metabolic disorders^(14–16) in offspring. Moreover, we have previously

Abbreviations: ACC, Acetyl-CoA carboxylase; C, control; CLA, conjugated linoleic acid; CPT-1a, carnitine palmitoyltransferase 1a; EWAT, epididymal white adipose tissue; FA, fatty acids; FAME, fatty acid methyl esters; FAS, fatty acid synthase; G6PDH, glucose-6 phosphate dehydrogenase; LPL, lipoprotein lipase; SCD-1, stearoyl-CoA desaturase-1; SREBP-1c, sterol regulatory element-binding protein-1c; TAG-SR, TAG secretion rate.

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shown⁽¹⁷⁾ that mice fed high-fat diets increased fetal resorption, decreased the IL-4 placental gene expression in the first generation and increased the TNF α gene expression in the second generation. In this generation, CLA normalised *Tnfa* decreased *Il4* and *transforming growth factor β 1*. Nevertheless, at least to the best of our knowledge, there is no information on the metabolic effects of CLA at recommended fat levels. Therefore, the aim of this study was to investigate the effect of maternal CLA consumption on TAG regulation and some inflammatory parameters related with TAG accretion in adult male rat offspring receiving or not receiving CLA.

Methods

Materials and diets

Nutrients and other chemical compounds, vitamins and minerals for the preparation of the diets, were of chemical grade or better, with the exception of soyabean oil (Cada Día), sucrose, cellulose and maize starch, which were obtained from local sources. CLA-mix oil was obtained from Lipid Nutrition B.V. and consisted of an equimolecular mixture of *c9t11*-CLA and *t10c12*-CLA. All solvents and reagents used for the fatty acid (FA) quantification were of chromatography grade purchased from Merck. Internal standard glyceryl tritridecanoate (13:0-TAG) and external standards GLC-463 containing fifty-two fatty acids methyl esters (FAME) mixture (purity >99%) were purchased from Nu-Chek. CLA, *cis/trans* mix (catalogue no. 05507), was purchased from Sigma. Other FAME standards were provided by the International CYTED Net (208RT0343). Primers used for PCR analysis were synthesised by Invitrogen. For the enzyme assays, the materials were from Sigma. All the other chemicals and reagents used were at least American Chemical Society degree or molecular grade and purchased from Merck, Invitrogen and Applied Biosystems. The TAG test kit was obtained from the Sociedad de Bioquímicos. For all the experiments, two isoenergetic diets (Table 1) containing 7% (w/w) of different

dietary fats were used. The control (C) diet was based on the American Institute of Nutrition Ad-Hoc Committee recommendation (AIN-93G) formulated for growth, pregnancy and lactation phases of rodents⁽¹⁸⁾ and contained 7% of soyabean oil as source of fats. The CLA diet was identical to the C diet, with the exception that 1% (w/w) of the CLA-mix oil replaced 1% (w/w) of the soyabean oil. The FA composition of CLA-mix oil (of total FAME) was as follows: 16:0, 5.85%; 18:0, 1.20%; *c9*-18:1, 9.05%; *c11*-18:1, 0.41%; *c9t11*-CLA, 38.99%; *t10c12*-CLA, 38.76%; *c9,c12*-18:2, 1.08%; and other FA, 4.66%. The FA composition of diets (Table 2) was determined by GC as indicated below. The diets were freshly prepared every 3 d, gassed with N and stored at 0–4°C.

Animals and experimental design

All the experiments were conducted in compliance with the regulations of the School of Biochemistry (Universidad Nacional del Litoral) and Guide to the Care and Use of Experimental Animals of Laboratory⁽¹⁹⁾. Male and virgin mature female Wistar rats were provided from the facilities at our University. The animals were housed in collective cages under controlled conditions (23 \pm 2°C and 12 h light–12 h dark) and free access to standard food and water. After 1 week of adaptation period, the female rats were randomly divided into two groups (*n* 6/group) and fed C or CLA diets (initial weights: 274.1 (SEM 8.6) and 269.9 (SEM 5.7) g, respectively) for 4 weeks before mating and throughout pregnancy and lactation. At parturition time, litter size was standardised to eight pups per dam in both C- and CLA-fed mothers. The pups were maintained with their own mothers until weaning. In all, twelve post-weaning male pups were randomly selected from C-fed mothers and received the same diet (C/C group). A total of twenty-four post-weaning male pups from CLA-fed mothers were randomly divided into two subgroups. One subgroup of twelve pups was fed the CLA diet (CLA/CLA group) and the other subgroup of twelve pups was fed the C diet (CLA/C group). The animals from the C/C, CLA/C and CLA/CLA groups were fed the respective diets for 9 weeks.

After the dietary treatment, one set of 6 animals per group was fasted overnight and killed (09.00–11.00 hours) under

Table 1. Composition of the experimental diets

| g/kg dry diet | C* | CLA* |
|------------------|-------|-------|
| Soyabean oil | 70.0 | 60.0 |
| CLA-mix oil | – | 10.0 |
| Casein | 200.0 | 200.0 |
| Maize starch | 529.5 | 529.5 |
| Sucrose | 100.0 | 100.0 |
| Fibre | 50.0 | 50.0 |
| Vitamin mixture† | 10.0 | 10.0 |
| Mineral mixture‡ | 35.0 | 35.0 |
| L-Cys, L-Met | 3.0 | 3.0 |
| Choline | 2.5 | 2.5 |

C, control diet; CLA, conjugated linoleic acid diet.

* Homemade prepared based on the American Institute of Nutrition Ad-Hoc Committee recommendation (AIN-93G) formulated for growth, pregnancy and lactation phases of rodents⁽¹⁸⁾.

† Vitamin mixture (per kg of diet): nicotinic acid, 30.0 mg; pantothenate, 15.0 mg; pyridoxine, 6.0 mg; thiamin, 5.0 mg; riboflavin, 6.0 mg; folic acid, 2.0 mg; vitamin K, 750.0 μ g; *D*-biotin, 200.0 μ g; vitamin B₁₂, 24.0 μ g; retinyl acetate, 1.2 mg; cholecalciferol, 0.025 mg; DL- α -tocopheryl acetate, 50 mg.

‡ Mineral mixture (mg/kg of diet): Ca, 5000.0; P, 1561.0; K, 3600.0; S, 300.0; Na 1019.0; Cl, 1571.0; Mg, 507.0; Fe, 35.0; Zn, 30.0; Mn, 10.0; Cu, 6.0; iodine, 0.2; Mo 0.15; Se, 0.15; Si, 5.0; Cr 1.0; F, 1.0; Ni, 0.5; B, 0.5; Li, 0.1; and V, 0.1.

Table 2. Fatty acid composition of experimental diets

| Fatty acids (% of total fatty acids methyl esters) | C | CLA |
|--|-------|-------|
| 16:0 | 10.84 | 10.12 |
| 18:0 | 4.50 | 4.01 |
| <i>c9</i> -18:1 | 17.60 | 16.27 |
| <i>c11</i> -18:1 | 4.10 | 3.56 |
| <i>c9c12</i> -18:2 | 54.30 | 46.53 |
| <i>c6c9c12</i> -18:3 | 0.31 | 0.26 |
| <i>c5</i> -20:1 | 0.16 | 0.13 |
| <i>c9c12c15</i> -18:3 | 7.42 | 6.36 |
| <i>c11</i> -20:1 | 0.14 | 0.12 |
| <i>c9t11</i> -CLA | ND | 5.57 |
| <i>t10c12</i> -CLA | ND | 5.53 |
| <i>c9c12c15</i> -20:3 | 0.39 | 0.33 |
| 24:0 | 0.11 | 0.09 |
| NI | 0.13 | 1.12 |

C, control diet; CLA, conjugated linoleic acid diet; ND, not detected; NI, not identified.

anaesthesia (1 mg azepromazine + 100 mg ketamine/kg body weight) by cardiac exsanguination. Blood was collected and serum was obtained after centrifugation (1000 **g** for 10 min at 4°C). Liver, gastrocnemius muscle (GM) and epididymal white adipose tissue (EWAT) were dissected, weighed and immediately frozen. All samples were stored at –80°C until analysis.

Fatty acid composition in liver, epididymal white adipose tissue, serum and dietary lipids

The CLA levels in liver, EWAT and serum, and the FA composition of experimental diets, were determined by GC using a Shimadzu (GC 2014) chromatograph equipped with a flame ionisation detector. Analyses were carried out with a capillary column CP Sil 88 (100 m, 0.25 µm film thickness; Varian). The carrier gas was H₂ with a split ratio of 1:10. The column temperature was held at 75°C for 2 min after injection, and then 5°C/min to 170°C, held for 40 min, 5°C/min to 220°C and held for 40 min. Injection volume was 0.5 µl and the column flow was 0.8 ml/min. Total fat in tissues, serum and diets was extracted using the method described by Bligh & Dyer⁽²⁰⁾. The FAME were formed by transesterification with methanolic potassium hydroxide solution as an interim stage before saponification (ISO 5509:2000, Point 5 International Union of Pure and Applied Biochemistry method 2.301). The FAME were identified by comparison of their retention times relative to those of commercial standards. Values of FA content were expressed as percentage of total FA. Further GC details could be obtained from previous publications^(21,22).

TAG levels in serum and liver

TAG levels in serum were determined by spectrophotometric methods using a commercially available test kit. Portions of frozen tissue were powdered and homogenised in saline solution (10%, w/v) for TAG determination⁽²³⁾.

Lipoprotein lipase activity in epididymal white adipose tissue and gastrocnemius muscle

The adipose tissue lipoprotein lipase (LPL) enzyme activity was quantified in EWAT acetone powder by the fluorometric method of Del Prado⁽²⁴⁾. To assess muscle LPL activity, GM samples were homogenised in a NH₄Cl/NH₄OH-Heparin buffer and the LPL activity quantification was performed by the technique of Del Prado⁽²⁴⁾. Values were expressed as nmol fluorescein/min per total tissue or as nmol fluorescein/min per g tissue. Further methodological details about the assessment have been previously published⁽²²⁾.

Lipogenic and β-oxidative enzyme activities

In liver homogenates, acetyl-CoA carboxylase (ACC; *EC* 6.4.1.2), fatty acid synthase (FAS; *EC* 2.3.1.85) and glucose-6-phosphate dehydrogenase (G6PDH; *EC* 1.1.1.49) activities, as lipogenic enzymes, were measured according to the methods of Zimmerman⁽²⁵⁾, Lynen⁽²⁶⁾ and Kuby & Noltmann⁽²⁷⁾, respectively.

The enzyme activities were expressed either as mU/mg of protein, where 1 mU was 1 nmol NADH consumed (ACC), 1 nmol NADPH consumed (FAS) or 1 nmol NADPH produced (G6PDH) per min. Carnitine palmitoyltransferase Ia (CPT-Ia; *EC* 1.3.99.3) activity, as β-oxidative enzyme, was assessed in the liver mitochondrial fraction by the method described by Bieber *et al.*⁽²⁸⁾ and expressed as mU/mg of protein (1 mU = 1 nmol CoA/min). Protein content was determined using bovine serum albumin as standard by the Lowry method⁽²⁹⁾.

Hepatic mRNA levels of lipogenic and β-oxidative enzymes, transcriptional factors and inflammatory cytokines

Total RNA was isolated from liver using Trizol according to the manufacturer's instructions. RNA samples were then treated with a DNA-free kit (Applied Biosystems) to remove any contamination with genomic DNA. The yield and quality of the RNA were assessed by measuring absorbance at 260, 270, 280 and 310 nm and by electrophoresis on 1.3% agarose gels. A measure of 1.0 µg of total RNA of each sample was reverse-transcribed to first-strand complementary DNA (cDNA) using a M-MLV RT. Relative mRNA levels were quantified using real-time PCR with a StepOne 18 TM Real-Time PCR Detection System (Applied Biosystems). Sequence-specific primers were used (Genbank: ACC, NM_022193.1; FAS, NM_017332.1; sterol regulatory element-binding protein-1c (SREBP-1c), NM_001276708.1; stearoyl-CoA desaturase-1a (SCD-1a), NM_139192.2; CPT-Ia, NM_031559.2; PPARα, NM_013196.1; IL-6, NM_012589.2; IL-1β, NM_031512.2; PPARγ, NM_013124.3; β-actin, NM_031144.3; ubiquitin C (UBC), NM_017314.1 and hypoxanthine phosphoribosyltransferase 1 (HPRT1), NM_012583.2), commercially synthesised (Invitrogen Custom Primers) and the sequences were as follows: ACC: 5'-AAC AGT GTA CAG CAT CGC CA-3' (forward), 5'-CAT GCC GTA GTAG GTT GAG GT-3' (reverse); FAS: 5'-CAG AAC TCT TCC AGG ATAG TCA ACA -3' (forward), 5'-GTC GCC CTAG TCA AGG TTC AG -3' (reverse); SREBP-1c, 5'-GGA GCC ATAG GAT TAGC ACA TT-3' (forward), 5'-GCT TCC AGA GAG GAG CCC AG-3' (reverse); SCD-1, 5'-CAC ACG CCG ACC CTC ACA ACT-3' (forward), 5'-TCC GCC CTT CTC TTT GAC AGC C-3' (reverse); CPT1, 5'-ACG TAGA GTAG ACT GGT GGG AAG AAT-3' (forward), 5'-TCT CCA TAGG CGT AGT AGT TAGC TAGT-3' (reverse); PPARα, 5'-CCC CAC TTAG AAG CAG ATAG ACC-3' (forward), 5'-CCC TAA GTA CTAG GTA GTC CGC-3' (reverse); IL-6, 5'-CCT TCT TAGG GAC TAGA TAGT TAGT TAGA C-3' (forward), 5'-GGG TAGG TAT CCT CTAG TAGA AGT CTC C-3' (reverse); IL-1β, 5'-AGG CAG TAGT CAC TCA TTAG TAGG C-3' (forward), 5'-TCA CAT GGG TCA GAC AGC ACG-3' (reverse); PPARγ, 5'-CGG AGT CCT CCC AGC TAGT TCG CC-3' (forward), 5'-GGC TCA TAT CTAG TCT CCG TCT TC-3' (reverse); β-actin 5'-CAT GAA GAT CAA GAT CAT TAGC TCC T-3' (forward), 5'-CTAG CTT GCT GAT CCA CAT CTAG-3' (reverse); UBC 5'-ACACCAAGAAGGTCAAACAGGA-3' (forward), 5'-CA CCTCCCATCAAACCCAA-3' (reverse); HPRT1 5'-TCCTCCTCA GACCGCTT TTC-3' (forward), 5'-ATCACTAATCACGACGCT GGG-3' (reverse). Standard curves for each primer were generated on separate runs using several serial dilutions

(1/10–1/1000) of pooled cDNA samples. The corresponding primer efficiency (E) of one cycle in the exponential phase was calculated according to equation $E = 10^{(-1/\text{slope})(30)}$. All the efficiencies of the primers were 100 (SEM 10)%. Target genes were normalised with the geometric mean of three housekeeping genes: β -actin, UBC and HPRT1⁽³¹⁾. Relative expression ratios were calculated using the recommended $2^{-\Delta\Delta C_t}$ method⁽³²⁾.

Hepatic TAG secretion rate

Another set of six animals per group was used; the animals were fasted overnight and anaesthetised as mentioned above. Then, 600 mg/kg body weight of triton WR 1339 in saline solution, an agent known to inhibit the peripheral removal of TAG-rich lipoproteins, was injected intravenously⁽³³⁾. Blood samples were taken immediately before and 120 min after the injection of the triton solution for the estimation of TAG accumulation in serum. Hepatic TAG secretion rate (TAG-SR) was estimated based on serum TAG concentration at 0 and 120 min, plasma volume and body weight. Further details have been previously reported⁽³⁴⁾.

Statistical analysis

Values were expressed as means with their standard errors of six animals per group. The minimum sample size needed to detect a statistically significant difference ($P < 0.05$) was calculated⁽³⁵⁾. A sample size of six had an 80% power ($P = 0.05$). The statistical analysis was performed by one-way ANOVA (1 × 3) followed by Scheffe's test ($P < 0.05$), using SPSS version 17.0 (SPSS Inc.), except for the comparison between CLA isomers shown in Table 4 for which Student's *t* test was used. All *post hoc* multiple comparisons were made using Tukey's critical range test. Significant differences were considered at $P < 0.05$.

Results

Nutritional parameters and TAG content in serum and liver

During all the experiment in dams and offspring, the diets were well accepted and the animals showed a healthy status without any pathological manifestation owing to the treatments received. The body weights (g) of dams after 4 weeks of dietary treatment before mating were C: 357.2 (SEM 14.3) and CLA: 347.4 (SEM 15.7) and the mean food intakes (kJ/d) were C: 255.9 (SEM 12.0) and CLA: 253.3 (SEM 5.8). Body and tissue weights of offspring and serum and liver TAG content are shown in Table 3. The final body weights decreased in groups CLA/CLA and CLA/C, parallel to a significant decrease in the EWAT weights. The absolute liver weight was lower in the CLA/CLA and CLA/C groups, but their relative weights were not affected by the dietary treatments. Serum TAG levels were increased in the CLA/CLA and CLA/C groups, and liver TAG levels were decreased in the CLA/CLA group, but not in the CLA/C group.

Percentage of individual conjugated linoleic acid in serum, liver and epididymal white adipose tissue

The percentage of individual CLA isomers in serum and tissues of offsprings was expressed as % of total FAME (Table 4). Both

Table 3. Body and tissue weights and serum and liver TAG content of offspring (Mean values with their standard errors; *n* 6 animals/group)

| | C*/C† | | CLA*/CLA† | | CLA*/C† | |
|---------------------------|--------------------|------|--------------------|------|--------------------|------|
| | Mean | SEM | Mean | SEM | Mean | SEM |
| Final BW (g) | 324.4 ^a | 6.2 | 299.9 ^b | 2.4 | 301.6 ^b | 7.1 |
| Liver weight (g) | 8.19 ^a | 0.14 | 7.57 ^b | 0.11 | 7.64 ^b | 0.12 |
| Liver weight (g/100 g BW) | 2.52 | 0.05 | 2.56 | 0.04 | 2.54 | 0.02 |
| EWAT (g) | 3.48 ^a | 0.60 | 2.74 ^b | 0.06 | 2.41 ^b | 0.13 |
| EWAT (g/100 g BW) | 1.07 ^a | 0.02 | 0.91 ^b | 0.02 | 0.80 ^c | 0.05 |
| Serum TAG (mmol/l) | 1.06 ^a | 0.07 | 1.77 ^b | 0.09 | 2.03 ^b | 0.19 |
| Liver TAG (μmol/g) | 11.09 ^a | 0.45 | 8.41 ^b | 0.75 | 11.40 ^a | 0.43 |

C, control diet; CLA, conjugated linoleic acid diet; BW, body weight; EWAT, epididymal white adipose tissue.
^{a,b,c} Mean values in the same row with unlike superscript letters were significantly different ($P < 0.05$). Statistical analyses between different groups were established by one-way ANOVA (1 × 3), followed by Scheffe's test.
 * Diet fed to the dam.
 † Diet fed to the male offspring.

Table 4. Percentage of individual conjugated linoleic acid (CLA) (% of total fatty acids methyl esters) in serum, liver and epididymal white adipose tissue (EWAT) of offspring (Mean values with their standard errors; *n* 6 animals/group)

| | C†/C‡ | | CLA†/CLA‡ | | CLA†/C‡ | |
|--------------|-----------------|-----|--------------------|-------------------|-----------------|-----|
| | Mean | SEM | Mean | SEM | Mean | SEM |
| Serum | | | | | | |
| c9t11-CLA | ND ^a | – | 0.76 ^{b*} | 0.07 | ND ^a | – |
| t10c12-CLA | ND ^a | – | 0.12 ^b | 0.02 | ND ^a | – |
| Liver | | | | | | |
| c9t11-CLA | ND ^a | – | 1.12 ^{b*} | 0.19 | ND ^a | – |
| t10c12-CLA | ND ^a | – | 0.38 ^b | 0.13 | ND ^a | – |
| EWAT | | | | | | |
| c9t11-CLA | ND ^a | – | 2.95 ^{b*} | 0.10 | ND ^a | – |
| t10c12-CLA | ND ^a | – | 1.39 | 0.13 ^b | ND ^a | – |

C, control diet; ND, not detected.
^{a,b} Mean values in the same row with unlike superscript letters were significantly different ($P < 0.05$). Statistical analyses between different groups were established by one-way ANOVA (1 × 3), followed by Scheffe's test.
 * Statistical differences between c9t11-CLA and t10c12-CLA isomers were established by Student's *t* test ($P < 0.05$).
 † Diet fed to the dam.
 ‡ Diet fed to the male offspring.

CLA isomers were incorporated in serum, liver and EWAT of the CLA/CLA group, but we did not detect any CLA isomer in CLA/C rats. The percentage of individual CLA in the tissues of CLA/CLA rats was EWAT > liver > serum. In addition, all tissues showed higher levels of c9t11-CLA than the t10c12-CLA isomer.

Serum and liver TAG regulation

The serum fasting TAG levels were mainly modulated by liver TAG secretion and peripheral TAG removal (Table 5). The hepatic TAG-SR did not change in the CLA/CLA group; however, it was significantly decreased in the CLA/C group. The serum TAG removal rate was estimated by the EWAT and GM LPL activities. The EWAT LPL activity, expressed either by total tissue or by g of tissue, was diminished in the CLA/CLA and CLA/C groups; however, this activity was not modified by any dietary treatment in the GM.

Table 5. Serum and liver TAG regulation of offspring (Mean values with their standard errors; *n* 6 animals/group)

| | C*/C† | | CLA*/CLA† | | CLA*/C† | |
|--|--------------------|------|--------------------|------|--------------------|------|
| | Mean | SEM | Mean | SEM | Mean | SEM |
| Hepatic TAG-SR nmol/min/100 g | 209.0 ^a | 3.3 | 210.6 ^a | 7.5 | 166.8 ^b | 5.9 |
| EWAT LPL activity nmol/min per EWAT | 958.0 ^a | 89.8 | 446.3 ^b | 92.4 | 251.3 ^b | 13.6 |
| nmol/min per µg protein | 151.2 ^a | 18.4 | 54.3 ^b | 4.7 | 35.1 ^b | 4.6 |
| GM LPL activity nmol/min per GM | 3.2 | 0.2 | 3.4 | 0.1 | 3.3 | 0.2 |
| nmol/min per µg protein | 46.7 | 3.4 | 47.4 | 2.2 | 43.6 | 1.1 |
| Liver enzyme activities | | | | | | |
| ACC (mU/mg protein) | 84.3 ^a | 3.0 | 51.0 ^b | 2.6 | 59.0 ^b | 6.1 |
| FAS (mU/mg protein) | 5.2 ^a | 0.4 | 3.3 ^b | 0.2 | 3.3 ^b | 0.2 |
| G6PDH (mU/mg protein) | 54.1 ^a | 4.0 | 36.3 ^b | 2.6 | 21.6 ^c | 1.6 |
| CPT-1a (mU/mg protein) | 1.8 | 0.2 | 1.5 | 0.1 | 1.7 | 0.2 |

C, control diet; CLA, conjugated linoleic acid diet; TAG-SR, TAG secretion rate; EWAT, epididymal white adipose tissue; GM, gastrocnemius muscle; LPL, lipoprotein lipase; ACC, acetyl-CoA carboxylase; FAS, fatty acid synthase; G6PDH, glucose-6-phosphate dehydrogenase; CPT-1a, carnitine palmitoyltransferase-1a; 1 mU, 1 nmol NADH consumed/min (ACC); 1 nmol NADPH consumed/min (FAS); 1 nmol NADPH produced per min (G6PDH); 1 nmol CoA/min (CPT-1a).

^{a,b,c} Mean values in the same row with unlike superscript letters were significantly different (*P* < 0.05). Statistical analyses between different groups were established by one-way ANOVA (1 × 3), followed by Scheffe's test.

* Diet fed to the dam.

† Diet fed to the male offspring.

Table 6. Hepatic mRNA levels of lipogenic and β-oxidative enzymes, transcriptional factors and inflammatory cytokines of offspring (Mean values with their standard errors; *n* 6 animals/group (relative units))

| | C*/C† | | CLA*/CLA† | | CLA*/C† | |
|----------|--------------------|-------|--------------------|-------|--------------------|-------|
| | Mean | SEM | Mean | SEM | Mean | SEM |
| ACC | 1.000 ^a | 0.033 | 0.551 ^b | 0.097 | 0.446 ^b | 0.049 |
| FAS | 1.000 ^a | 0.062 | 0.502 ^b | 0.096 | 0.936 ^a | 0.065 |
| SCD-1 | 1.000 ^a | 0.020 | 0.325 ^b | 0.100 | 0.656 ^c | 0.042 |
| SREBP-1c | 1.000 ^a | 0.021 | 0.643 ^b | 0.152 | 1.206 ^a | 0.030 |
| CPT-1a | 1.000 ^a | 0.025 | 0.881 ^a | 0.063 | 0.509 ^b | 0.029 |
| PPARα | 1.000 ^a | 0.018 | 1.050 ^a | 0.065 | 0.733 ^b | 0.020 |
| IL-6 | 1.000 | 0.250 | 0.823 | 0.099 | 1.025 | 0.156 |
| IL-1β | 1.000 ^a | 0.056 | 0.613 ^b | 0.041 | 0.666 ^b | 0.060 |
| PPARγ | 1.000 ^a | 0.032 | 0.716 ^b | 0.049 | 0.559 ^b | 0.014 |

C, control diet; CLA, conjugated linoleic acid diet; ACC, acetyl-CoA carboxylase; FAS, fatty acid synthase; SCD-1, stearoyl-CoA desaturase-1; SREBP-1c, sterol regulatory element-binding protein 1c; CPT-1a, carnitine palmitoyltransferase 1a.

^{a,b} Mean values in the same row with unlike superscript letters were significantly different (*P* < 0.05). Statistical analyses between different groups were established by one-way ANOVA (1 × 3), followed by Scheffe's test.

* Diet fed to the dam.

† Diet fed to the male offspring.

The fasting hepatic TAG levels were mostly regulated by lipogenesis, β-oxidation and hepatic TAG-SR (Table 5). The ACC, FAS and G6PDH activities were diminished in the CLA/CLA and CLA/C groups. Meanwhile, the activity of the main enzyme involved in the hepatic FA oxidation, CPT-1a, was not modified in the CLA/CLA and CLA/C groups.

The liver expression of some key genes and transcriptional factors related to FA biosynthesis (*Acc*, *Fas*, *Scd1* and *Srebp1c*) and β-oxidation (*Cpt1a*, *Ppara*), as well as inflammation (*Il6*, *Il1b*, *Pparg*), are shown in Table 6. The levels of mRNA of ACC,

FAS, SCD-1a and SREBP-1c were diminished in the CLA/CLA group. The ACC, but not FAS, SCD-1 and SREBP-1c, gene expression was decreased in the CLA/C group. As regards fatty acid oxidation, CPT-1a and PPARα mRNA levels were unchanged in CLA/CLA group, but decreased in the CLA/C group. In addition, some of the genes related to inflammation were assayed, showing a significant decrease in the *Pparg* and *Il1b* expression in both CLA/CLA and CLA/C. No changes were observed in IL-6 mRNA levels.

Discussion

Dietary CLA reduces body fat accumulation and modulates lipid metabolism, as well as the inflammatory response, which could have a potential key role during the gestation period. The CLA-mix oil contains mainly *c9t11*-CLA and *t10c12*-CLA in equimolecular proportion; however, their biological effects are different. Even though both isomers are involved in lipid metabolism regulation, *t10c12*-CLA is principally responsible for increasing lipolysis and the β-oxidation/lipogenesis ratio modulating fat accretion⁽⁷⁾, whereas the *c9t11*-CLA has an anti-inflammatory response⁽³⁶⁾. Taking into account that the reduction in lipid depots might induce metabolic changes in the offspring, increasing the risk of metabolic alteration in adult life⁽¹⁵⁾, the aim of this study was to investigate the effect of maternal CLA consumption on nutritional parameters, TAG levels in serum and liver, as well as their regulation in adult offspring rats that received CLA or not. In addition, some gene expression factors related to the inflammatory process were studied. To the best of our knowledge, this is the first study to investigate the influence of maternal CLA on TAG metabolism in adult offspring rats fed normo-energetic diets.

It is known that CLA reduces the fat mass by different mechanisms including decreased fatty acid uptake, increased fatty acid oxidation, increased overall energy expenditure and others⁽¹⁾. In agreement with several studies in different experimental animal models fed CLA^(8,34,37), the present results in offspring male rats fed CLA (CLA/CLA group) showed a lower final body weight related to reduced fat depots. It is important to note that the results on rat body composition are controversial, depending on the type of isomer and level of CLA, nutritional status and time of feeding^(3,8). There are few studies showing the effect of CLA on adult offspring rats fed the same diet as the dams. In a previous study from our laboratory⁽¹⁷⁾, CLA reduced the EWAT weights at first and second generation of mice, and similar results have been obtained in rats of the first generation⁽¹⁴⁾. In the present study, the effects on fat depots were observed in the presence of high CLA levels in adipose tissue (CLA/CLA group) and in the absence of CLA (CLA/C group). It is well known that CLA isomers are incorporated and metabolised into the tissues, and the rate of CLA metabolism is sensitive to experimental conditions⁽³⁸⁾. However, to the best of our knowledge, there is no study showing the disappearance rate of CLA in the offspring of CLA-fed mothers; therefore, the measurement of CLA levels in tissues of the CLA/C group was performed to warrant the absence of CLA in tissues of offspring. Thus, the metabolic effects of the diets in the offspring could be specifically attributed to maternal CLA, which was administered

to the dams during pregnancy and lactation. Interestingly, those adult offspring from CLA-fed rats that were fed C diets after weaning (CLA/C group) also showed a reduction of body and EWAT weights. It is noteworthy that in both offspring from CLA dams the reduction of EWAT was associated with a lower LPL enzyme activity. These results might hypothesise that a programming mechanism of CLA during early life could be involved in controlling the adiposity in offspring through the fatty acid uptake by the EWAT.

The plasma TAG levels were regulated by the circulating TAG removal rate from the adipose tissue and muscle, as well as by the hepatic VLDL-TAG secretion. As the hepatic VLDL-TAG-SR was not increased and the muscle LPL enzyme activity did not change by the dietary treatments, the reduced LPL activity in EWAT could explain the higher plasma TAG levels in both CLA/C and CLA/CLA groups. A similar association between the increase of plasma TAG and the reduction of adipose tissue LPL activity was found in mice fed different types of oils supplemented with 1% CLA-mix oil at normal fat levels⁽³⁹⁾ and hamsters fed high-fat diets supplemented with a similar commercial CLA⁽⁴⁰⁾. However, compared with results previously observed in mice⁽³⁹⁾, in the present work, the VLDL-TAG-SR did not contribute to the hypertriglycerolaemia. Other authors found a reduction in mice fed high-fat diets^(34,41) or no changes in serum TAG in rats fed diets rich in saturated fats⁽⁴²⁾. Thus, the differences observed in the effect of CLA-mix oil are not only owing to the species but also to the type of isomer and dietary experimental protocol, among other reasons.

Liver is an organ that plays a crucial role in lipid metabolism and has been reported to be a key target for CLA⁽¹⁾. As shown in EWAT, CLA was incorporated into the liver of CLA/CLA, but was not detected in the CLA/C group. In the CLA/CLA group, the percentage of *c9/t11*-CLA was higher than that of the *t10,c12*-CLA isomer. Similar results have been previously shown in animals fed CLA diets⁽⁴³⁾. The effect of CLA on hepatic TAG has been proposed to be species specific⁽⁴⁴⁾. In contrast to the effects described by CLA inducing a significant liver TAG accretion in mice^(7,34), neither liver steatosis nor hepatomegaly was observed in both groups of offspring rats from CLA-fed dams. Furthermore, CLA/CLA rats showed a reduction of hepatic TAG content. The lowering effect of CLA on hepatic TAG has been previously shown in rats under different conditions of liver steatosis^(43,45–49). Purushotham *et al.*⁽⁴³⁾ proposed that CLA attenuated hepatic steatosis in adult Wistar rats by modulating hepatic genes involved in lipogenesis and lipid oxidation. In addition, these authors suggested that the dissimilar effects of CLA on hepatic steatosis and insulin sensitivity may depend on differences in fat-to-lean partitioning between adipose and liver tissues in numerous animal models. Interestingly, in both groups of rats from CLA-fed dams, the lipogenesis was significantly reduced. Specifically, in the CLA/CLA group, the lower ACC, FAS and G6PDH activities and reduced mRNA levels of ACC, FAS, SCD-1 and SREBP-1c, with normal CPT-1a activity and mRNA levels clearly demonstrated an unbalance between lipogenesis and β -oxidation. In agreement with our results, CLA decreased the hepatic FAS⁽⁵⁰⁾ and SCD-1⁽⁵¹⁾ activities and FAS⁽⁵²⁾ and SCD-1^(51,52) mRNA levels. In addition, an

unchanged FA oxidation has also been observed⁽⁵³⁾. However, some reports have shown increased^(54,55) and reduced⁽⁵⁶⁾ FA oxidation. A reduced lipogenesis was also observed in the CLA/C group, but in contrast to the CLA/CLA group two different compensatory mechanisms might be involved to balance the hepatic TAG levels. Specifically, a decreased β -oxidation, as a result of a lower CPT-1a gene expression, and a reduction of hepatic TAG secretion contributed to maintain the normal hepatic TAG content. The most interesting and original result is that a lower ACC, FAS and G6PDH enzyme activity and a diminished mRNA level of ACC was also observed even in the absence of CLA in the diet. Reinforcing the above-mentioned effects observed in body composition, CLA might programme the lipid metabolism driving to a prevention of liver TAG accretion. There are no studies reporting data to contrast with the present findings.

The beneficial effects of CLA on immune and inflammatory responses in different animal models, describing the mechanism of action of CLA, particularly the *c9,t11-18:2*, have recently been reviewed⁽⁵⁷⁾. Thus, after CLA exposure, a reduction of pro-inflammatory cytokine production has been described, mainly IL-6, TNF α , IFN γ , and IL-1 β , which could play an important role in the pathogenesis of many chronic-inflammation-mediated diseases^(58,59). Reynolds *et al.*⁽¹⁴⁾ and Segovia *et al.*⁽¹⁵⁾ demonstrated in different experimental animal models that CLA reversed maternal meta-inflammation induced by high-fat diet and prevented programming of adverse offspring metabolic outcomes. Our results also show that CLA might down-regulate the liver gene expression of some pro-inflammatory factors such as IL-1 β and PPAR γ in offspring, even at recommended dietary fat levels.

In brief, these results demonstrate a programming effect of CLA on lipid metabolic pathways leading to a preventive effect on the TAG accretion in adipose tissue and liver of male rat offspring. This knowledge could be important to develop some dietary strategies to reduce the incidence of obesity and fatty acid liver disease in humans.

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