

Effects of carnosine supplementation on markers for the pathophysiological development of metabolic dysfunction-associated steatotic liver disease in a diet-induced model

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Abstract

Consumption of diets high in sugar and fat is related to the development of Metabolic dysfunction-associated steatotic liver disease (MASLD). Carnosine (CAR) is a dipeptide with antioxidant and anti-inflammatory action and has been studied for treating diseases. This work aimed to evaluate the effects of CAR on diet-induced MASLD in rats. Male Wistar rats were distributed into 2 groups (17 weeks): normocaloric (Co, n=12), and hypercaloric diet rich in lipids and simple carbohydrates (MASLD, n=12). After, the animals were redistributed to begin the treatment with CAR (4 weeks): Co (n=6), Co + CAR (n=6), MASLD (n=6), and MASLD + CAR (n=6), administered intraperitoneally (250mg/kg). Evaluations included nutritional, hormonal and metabolic parameters; hepatic steatosis, inflammatory and oxidative markers. MASLD group had a higher adiposity index, systolic blood pressure, glucose, plasma and liver triglycerides and cholesterol, insulin, hepatic steatosis, oxidative markers, and lower PPAR- α (Peroxisome Proliferator-activated receptor α), compared to the Co. CAR attenuated plasma and hepatic triglyceride and cholesterol levels, hepatic steatosis, CD68+ macrophages, and hepatic oxidative markers, in addition to increasing HDL cholesterol levels and PPAR- α , compared to the untreated MASLD group. CAR acts in important pathophysiological processes of MASLD and may be a therapeutic compound to control the disease.

Keywords: Metabolic dysfunction-associated steatotic liver disease, carnosine, obesity, hypercaloric diet.

1. Introduction

Non-alcoholic fatty liver disease (NAFLD) is the most common liver injury, which affects about a third of the world's population and is defined by the accumulation of fat in liver tissue without alcohol consumption and other liver damage (Younossi et al., 2016). In 2023, a new terminology and definition were proposed, starting to be called Metabolic dysfunction-associated steatotic liver disease (MASLD). Therefore, the MASLD is based on hepatic steatosis with one of the five cardiometabolic risk factors: overweight or obesity, type 2 diabetes mellitus or hyperglycemia, hypertension or antihypertensive drug treatment, and dyslipidemia (Rinella et al., 2023).

MASLD has a spectrum of evolution from simple steatosis and can involve Metabolic dysfunction-associated steatohepatitis (MASH) and even liver cancer (Carneros et al., 2020; Francisqueti-Ferron et al., 2022; Rinella et al., 2023). According to a survey based on data from Europe, North America, and Asia, the prevalence of patients with MASLD is around 39% (Lim et al., 2021), while MASH affects around 5% of the world's population (Younossi et al., 2016). Its pathogenesis is still not well defined, but there are two theories for its development: the first is the “two hits” theory, where it is believed that the first aggression is the imbalance caused by insulin resistance (IR) between the hepatic pathways of acquisition and elimination of lipids, and the second attack involves the processes of inflammation and oxidation (Day and James, 1998). The second theory is that its development takes place from a “multiple hits” model, where several aggressive agents trigger this injury, being genetic, epigenetic, nutritional, and inflammatory, including changes in the microbiota, IR, and oxidative stress (Buzzetti et al., 2016; Day and James, 1998).

The Western diet, rich in simple carbohydrates and fats, is a crucial precursor factor of MASLD, which can cause harmful effects on the liver by accumulating these macronutrients (Rives et al., 2020; Vancells Lujan et al., 2021). Data in the literature indicate that women who consume a diet high in sugar and fat have a 119% greater risk (OR, 2.19; 95% CI, 1.40–3.46) of developing fatty liver disease (Jia et al., 2015); similar data were found for adolescents aged 17 years who had the same Western diet consumption pattern (Oddy et al., 2013). The accumulation of fatty acids in the liver generates hepatic steatosis. It favors the excessive production of ROS (reactive oxygen species), causing an imbalance in the antioxidant system and, consequently, oxidative

stress, which can cause various cellular damage (Arroyave-ospina et al., 2021; Yki-Järvinen et al., 2021).

Given the complexity of MASLD and its consequences, the search for forms of prevention and treatment is necessary. Currently, there is no specific pharmacological treatment for this disease, only drugs that treat specific metabolic alterations, such as dyslipidemia and hyperglycemia, are used (Nassir, 2022; Salvoza et al., 2022). On the other hand, the search for bioactive compounds that can be used to improve these conditions emerges, and one of these compounds is carnosine (CAR), a natural dipeptide composed of β -alanine and L-histidine, which can be found in the human body, in higher concentrations in skeletal muscle and brain (Aldini et al., 2021).

For many years CAR was the focus of studies only related to improving the performance of physical exercises, mainly due to its buffering capacity in the muscles (Matthews et al., 2019). However, CAR has been identified as a potential therapeutic agent in treating diseases due to its antioxidant, anti-inflammatory, and ROS-sequestering activity (Chmielewska et al., 2020). Some studies demonstrated that CAR was able to attenuate the development of type 2 diabetes in ob/ob mice (Albrecht et al., 2017), improve parameters related to metabolic syndrome induced by diet in Wistar rats (Al-Sawalha et al., 2019), as well as improve hepatic steatosis in mice that consumed a high-fat diet (Mong et al., 2011) and in a model of type 2 diabetes in db/db mice (Schwank-Xu et al., 2021). Faced with the need to clarify the biological effects of CAR and the scarcity of studies relating this dipeptide to MASLD, the present study aimed to verify the effects of CAR supplementation on markers of the MASLD pathophysiology induced by diet in rats.

2. Material and Methods

2.1. Animals and Experimental Protocol

This study was approved by the Ethics Committee on Animal Use (n° 1322/2019). It was carried out following the Guide for the Care and Use of Laboratory Animals published by the U.S. National Institutes of Health. Male Wistar rats from the Central Animal Facility of the Faculty of Medicine of Botucatu, approximately 21 days old, were kept in individual cages in an environment with temperature ($24\pm 2^\circ\text{C}$), humidity ($55\pm 5\%$) and light (light cycle-dark 12h) controlled. Initially, they were distributed into two experimental groups, fed different diets for MASLD induction: normocaloric ($n=12$) and hypercaloric rich in sugar and fat ($n=12$). On the 17th week, after the difference between plasma triglycerides, defined as an inclusion criterion, the animals were redistributed into

four groups for treatment: a control group that continued receiving the normocaloric diet (co, n=6), a control group that received the normocaloric diet treated with carnosine (Co+CAR, n=6), a group that continued receiving the hypercaloric diet (MASLD, n=6) and a group that received the hypercaloric diet treated with carnosine (MASLD+CAR, n=6), for four weeks. Feed and water were offered ad libitum. The animals that received the hypercaloric diet also received sucrose in their drinking water at a concentration of 25% throughout the experiment. Both diets were produced according Francisqueti et al. (2017), and are described in Table 1. The groups received CAR supplementation (250 mg/kg/day) or vehicle (sterile saline solution, 0.9% NaCl) intraperitoneally for four weeks. At the end of the 21 weeks, the animals fasted for 8 hours and then were anesthetized (Thiopental 120 mg/kg/i.p.) and euthanized by decapitation. Blood samples, adipose tissue, and liver were collected.

Table 1. Diet composition and nutritional values.

Components	Normocaloric	Hypercaloric
Soybean meal (g/kg)	335	340
Sorghum (g/kg)	278	80
Soy hulls (g/kg)	188,5	116,7
Dextrin (g/kg)	146,5	20
Maize starch (g/kg)	278	80
Sucrose (g/kg)	-	80
Fructose (g/kg)	-	180
Soybean oil (g/kg)	23	-
Lard (g/kg)	-	154,3
Minerals (g/kg)	25	25
Salt (g/kg)	4	4

2.2. Carnosine Supplementation

CAR (L-carnosine) was administered intraperitoneally (250 mg/kg) for four weeks after induction. The dose was defined based on studies where CAR presented effective results and did not generate toxic and/or adverse effects on the animals (Aydın et al., 2017; Kalaz et al., 2016). The infusion was performed using a 30G needle, with a liquid volume close to 500 µL being injected. The product CarnoPure™ (Flamma S.p.A., Chignolo d'Isola, Bergamo, Italy) was used exclusively for scientific experiments and used in studies abroad. This product was kindly provided by this company, free of charge, for carrying out this experiment.

2.3. Nutritional characterization and obesity

Nutritional characterization was evaluated based on feed, water, and caloric intake (feed and water). Food and water intake were calculated daily from the individual leftovers of each animal. Caloric intake was calculated through food intake multiplied by the energy value of each diet. The characterization of obesity was based on weight gain and adiposity index. Weight gain was calculated by the difference between the final weight and the initial weight of the animals [weight gain (g)=final weight (g) - initial weight (g)]. The adiposity index represents the ratio between the sum of fat deposits (visceral, epididymal, and retroperitoneal) and the final weight of the animals (adiposity index = sum of fat deposits (g)/final weight (g) x 100) (Costa et al., 2019).

2.4. Systolic Blood Pressure

Systolic blood pressure (SBP) was measured by the indirect method of tail plethysmography using an electric sphygmomanometer Narco Bio-System® PE 300, model 709-0610 (International Biomedical, Inc, Houston, Texas, USA). For this purpose, the animals were kept for 5 minutes in a wooden box heated between 38°C and 40°C to stimulate arterial vasodilation. Afterward, the cuff was attached to the tail of each animal and inflated (200 mmHg), and deflated, allowing the recording of arterial pulses collected in a Gould RS 3200 polygraph (Gould Instrumenta Valley View, Ohio, USA). The mean of the three PAS readings was recorded for each animal (Siqueira et al., 2022).

2.5. Biochemical and Hormonal Analysis

Glucose was determined using a glucometer (Accu-Check Performa; Roche Diagnostics, Indianapolis, IN, USA). The concentrations of urea, creatinine, uric acid, triglycerides, total cholesterol and its fractions, HDL cholesterol and non-HDL cholesterol (n-HDL) were analyzed by the enzymatic colorimetric method with commercial kits (BioClin®, Belo Horizonte, MG, Brazil) in automated biochemical equipment (Chemistry Analyzer BS-200, Mindray Medical International Limited, Shenzhen, China).

The insulin level was measured using the indirect enzyme immunoassay (ELISA) method using commercial kits (Linco Research Inc., R&D Systems, Millipore, and B-Bridge International Inc.), and the reading was performed on a Spectramax 190 microplate spectrophotometer (Molecular Devices®, Sunnyvale, CA, USA). The IR assessment was performed using the Homeostasis Model Index - Insulin Resistance (HOMA-IR) using glycemia and plasma insulin parameters. The calculation of the HOMA index obeyed the following formula: insulin ($\mu\text{U/mL}$) x blood glucose (mmol/L) / 22.5 (Francisqueti-Ferron et al., 2022).

2.6. Liver Tissue Analysis

2.6.1. Histological Parameters

After collection, liver tissue was stored in 4% paraformaldehyde with 0.1M phosphate buffer (pH=7.4) during the first 24 hours. Afterward, the tissue was transferred

to ethyl alcohol, and the samples were embedded in paraffin to obtain five μm -thick sections. Such sections (one per animal) were stained with Hematoxylin & Eosin (HE) and evaluated for steatosis, taking into account the macro and microvesicular aspects, according to the NAFLD Activity Score (NAS) parameter (Liang et al., 2014). Then, ten fields were randomly selected (20x objective), which were evaluated in each section, and the following scores were considered: 0 (<5% of the microscopic field), 1 (5-33%), 2 (33-66%), or 3 (>66%). Other liver samples were frozen and included in an Optimal Cutting Temperature medium (Tissue-Trek, USA), and 10 μm sections were obtained and stained with Oil red (Sigma Aldrich, USA) to obtain representative photos. To assess the presence and deposition of collagen sections samples were stained with picosirius red with subsequent detection by refringent images on optical microscope. The METAVIR score system was used to evaluate the level of fibrosis: F0: no fibrosis; F1: portal fibrosis without septa; F2: portal fibrosis with septa; F3: septa without cirrhosis; F4: cirrhosis.

2.6.2. Immunohistochemistry for CD68

Other 5 μm thick liver sections were submitted to antigen retrieval in 0.01M citrate buffer (pH 6.0, 120°C, five min.) in a Pascal pressure chamber (Dako Cytomation, Denmark). After blocking endogenous peroxidase with 10% H_2O_2 in phosphate-buffered saline (PBS) (15 minutes), the slides were treated with skimmed milk for 60 minutes to block nonspecific binding. Then, the histological sections were incubated in a humid chamber (4°C overnight) with a primary anti-CD68 antibody solution (macrophages, ab125212, 1:500 dilution, Abcam, UK). Then, the slides were incubated with HRP polymer (Horseradish peroxidase) in a single step (EasyPath - Erviegas, Brazil) (20 minutes at room temperature). The reactions were revealed with the chromogen 3'-diaminobenzidine (DAB) (Sigma Aldrich, USA), and the slides were counterstained with Harris hematoxylin. CD68+ macrophages were counted in 10 randomly selected fields (20x) and divided by field area (mm^2) using Image J software (NIH, USA) (Romualdo et al., 2022).

2.6.3. Oxidative Stress markers

Malondialdehyde (MDA) level was used to assess lipid peroxidation. Briefly, 250 μL of supernatant from hepatic tissue was added to 750 μL of 10% trichloroacetic acid for protein precipitation. The samples were centrifuged (3000 rpm, for 5 minutes; Eppendorf® Centrifuge 5804-R, Hamburg, Germany), and the supernatant was removed. Thiobarbituric acid (ATB) was added at a ratio of 0.67% (1:1), and the samples were heated for 15 minutes at 100°C. After cooling, the reading was performed at 535 nm in the Spectra Max 190 microplate reader (Molecular Devices®, Sunnyvale, CA, USA). The MDA concentration was obtained by the molar extinction coefficient ($1.56 \times$

105 M⁻¹cm⁻¹) and the absorption of the samples, and the final result was reported in nmol/g of protein (Siqueira et al., 2023).

To quantify protein carbonylation (CBO), 100 µL of liver tissue homogenate was used for 100 µL of 2,4-dinitrophenylhydrazine (DNPH, 10mM in 2M HCl). The samples were incubated for 10 minutes at room temperature, and then 50 µL of sodium hydroxide (6M NaOH) was added and incubated again for 10 minutes at room temperature. Reading at 450 nm was performed on a Spectramax 190 microplate reader (Molecular Devices®, Sunnyvale, CA, USA). The results were obtained by the molar extinction coefficient (22,000 M⁻¹ cm⁻¹) of DNPH and expressed in nmol/g of protein. Protein carbonylation levels were expressed in nmol of DNPH/mg of protein (Garcia et al., 2022).

Advanced oxidation protein products (AOPP) were detected in liver tissue. Tissue samples were homogenized in PBS buffer at 1:10 (sample:PBS). AOPP quantification was performed by diluting 1:5 of 200 µl of the sample (homogenate). 200 µl of Chloramine-T (0-100 µmol/L) was used for the calibration curve, and 200 µl of PBS as a blank. 10 µl of 1.16M KI (potassium iodide) and 20 µl of acetic acid were added to the samples. Absorbance was read immediately at 340 nm (Multiskan Ascent spectrophotometer, Labsystems, Vantaa, Finland). The final concentration of AOPP is expressed in chloramin units (µmol/L), and the result in the liver tissue is corrected for the amount of protein (Ferron et al., 2019).

2.6.4. Hepatic Levels of Triglycerides and Cholesterol

Intracellular levels of hepatic triglycerides and cholesterol were determined by a colorimetric method (Colorimetric Triglycerides and Cholesterol Assay Kit; Cayman Chemical, Ann Arbor, MI, USA). The results were corrected for the amount of tissue (Francisqueti-Ferron et al., 2022).

2.7. Gene Expressions

Gene expression of peroxisome proliferator-activated receptor α (PPAR-α; gene *Ppara*, primers F-AGATGGGCTGGTTCTTCCTC and R-CAGTGCATTTGCCAGG), very long-chain acyl-CoA dehydrogenase (VLCAD, gene *Acadvl*, primers F-ATGGAATTCATGGCCAGTGAGGCCACTC and R-TTTACGAATTCTCAGACTCTAAGGGGGT) and β-actin (gene *Actb*, primers F-GGGAAATCGTGCGTGACAT and R-CAGGAGGAGCAATGATCTT) were performed using real time PCR (polymerase chain reaction). Total RNA (ribonucleic acid) was extracted from liver tissue using the TRIzol® reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's specifications. The RNA was subjected to reverse transcription for conversion to cDNA (complementary deoxyribonucleic acid) (High-Capacity cDNA Reverse Transcription Kit, Applied Biosystems, CA, USA) using the

provided primers (Exxtend Biotecnologia Ltda, Campinas-SP, Brazil). The real-time PCR kit used the cDNA (TagMan® Gene Expression Assays, Applied Biosystems, Foster City, CA, USA) to evaluate expressions. The β -actin was used to normalize expressions of PPAR- α and VLCAD.

2.8. Statistical Analysis

Results are expressed as mean $\pm$ standard deviation. Two-way ANOVA determined differences between groups, and data were subjected to Tukey's post hoc test to compare all groups. Histological parameters are expressed as medians, and Kruskal-Wallis was used with Tukey's post hoc test. Statistical analyzes were performed using Sigma Plot 12.0 software for Windows (Systat Software Inc., San Jose, CA, USA). A p-value <0.05 was considered statistically significant and all performed statistical analyzes were tested to ensure the power to detect differences among groups and the sample size.

3. Results

3.1. Nutritional characterization and obesity

The nutritional characterization and obesity is presented in Table 2. It is possible to observe that the MASLD and MASLD+CAR groups had lower food intake, higher water, caloric intake, and more significant weight gain when compared to the Co and Co+CAR groups, respectively. There was no difference in the initial body weight between groups. The MASLD group had a higher final weight when compared to the Co group. The adiposity index determined the presence of obesity, and it can be seen that the MASLD induction groups (MASLD and MASLD+CAR) had higher adiposity indices when compared to the control groups (Co and Co+CAR, respectively).

Table 2. Nutritional characterization and obesity.

	Groups				Effects		
	Co	Co+CAR	MASLD	MASLD+CAR	Diet	CAR	Interactions
Food intake (g/day)	37.0 $\pm$ 2.31	36.3 $\pm$ 2.07	25.6 $\pm$ 0.904*	25.3 $\pm$ 1.72†	<0.001	0.528	0.858
Water intake (ml/day)	53.2 $\pm$ 3.79	52.0 $\pm$ 7.17	68.0 $\pm$ 4.02*	64.9 $\pm$ 3.48†	<0.001	0.320	0.564
Caloric intake (kcal/day)	133 $\pm$ 8.28	130 $\pm$ 7.42	166 $\pm$ 7.84*	168 $\pm$ 5.44†	<0.001	0.849	0.584
Initial body weight (g)	324 $\pm$ 25.0	324 $\pm$ 26.8	320 $\pm$ 16.4	314 $\pm$ 27.1	0.450	0.775	0.753
Final body weight (g)	502 $\pm$ 57.2	477 $\pm$ 25.8	569 $\pm$ 38.0*	530 $\pm$ 68.5	0.008	0.133	0.743
Weight gain (g)	145 $\pm$ 48.9	137 $\pm$ 23.2	250 $\pm$ 34.6*	198 $\pm$ 48.6†	<0.001	0.085	0.201
Adiposity index (%)	4.15 $\pm$ 1.60	3.12 $\pm$ 1.11	8.51 $\pm$ 1.03*	7.68 $\pm$ 2.56†	<0.001	0.19	0.883

Data are presented as mean $\pm$ standard deviation. Effects are the comparison by two-way ANOVA with Tukey's post-hoc test, with $p < 0.05$ as statistically significance. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine. * vs Co; † vs Co+CAR.

3.2. Systolic Blood Pressure

Figure 1 shows the Systolic Blood Pressure (SBP) levels. The MASLD group had a higher SBP compared to the Co group.

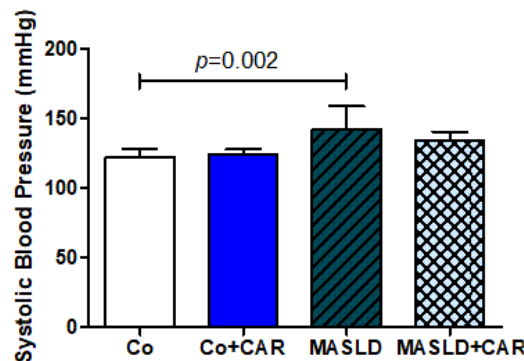


Figure 1. Systolic Blood Pressure Levels. Data are presented as mean $\pm$ standard deviation. Comparison by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.3. Biochemical and hormonal analysis

Table 3 shows the biochemical and hormonal markers. The MASLD and MASLD+CAR groups had lower urea levels when compared to the Co and Co+CAR groups, respectively. There was no significant difference uric acid, and creatinine. ALT levels were significantly lower in control treated with CAR and MASLD induction groups (MASLD and MASLD+CAR) had higher blood glucose levels when compared to the control groups (Co and Co+CAR, respectively). Co+CAR showed lower concentrations of total cholesterol compared to the Co group, while MASLD+CAR showed higher concentrations of total cholesterol compared to the Co+CAR group. Regarding triglycerides, the MASLD and Co+CAR groups had higher and lower concentrations when compared to the Co group. In contrast, the MASLD+CAR group had lower and higher levels when compared to the MASLD and Co+CAR groups, respectively. In addition, the MASLD+CAR group had higher HDL cholesterol (high-density lipoprotein) levels than the MASLD group. There was no statistical difference for non-HDL cholesterol levels. The MASLD and MASLD+CAR groups had higher insulin and HOMA-IR values when compared to the Co and Co+CAR groups, respectively.

Table 3. Biochemical and hormonal markers.

	Groups				Effects		
	Co	Co+CAR	MASLD	MASLD+CAR	Diet	CAR	Interactions
AST (U/l)	155 $\pm$ 34.3	125 $\pm$ 18.6	114 $\pm$ 21.0	130 $\pm$ 32.4	0.128	0.532	0.058
ALT (U/l)	104 $\pm$ 14.4	69.5 $\pm$ 16.1*	83.0 $\pm$ 26.5	87.5 $\pm$ 21.5	0.858	0.084	0.028
Uric acid (mg/dl)	0.967 $\pm$ 0.175	0.800 $\pm$ 0.420	1.27 $\pm$ 0.761	1.67 $\pm$ 1.69	0.150	0.768	0.476

Urea (mg/dl)	58.8 ± 8.66	61.0 ± 9.63	35.0 ± 15.5 [*]	44.2 ± 4.58 [†]	<0.001	0.195	0.417
Creatinine (mg/dl)	4.48 ± 3.08	4.59 ± 5.01	4.48 ± 1.68	4.36 ± 0.985	0.931	0.998	0.930
Blood Glucose (mg/dl)	78.0 ± 8.29	86.3 ± 11.4	88.8 ± 6.74 [*]	97.3 ± 4.50 [†]	0.004	0.020	0.980
Total Cholesterol (mg/dl)	62.3 ± 7.12	46.5 ± 10.1 [*]	52.7 ± 9.63	62.0 ± 8.32 [†]	0.430	0.380	0.002
Triglycerides (mg/dl)	70.4 ± 16.9	41.0 ± 8.32 [*]	205 ± 34.6 [*]	112 ± 27.4 ^{#†}	<0.001	<0.001	0.004
HDL (mg/dl)	14.5 ± 3.02	14.0 ± 1.79	11.4 ± 2.33	15.5 ± 3.27 [#]	0.471	0.114	0.047
n-HDL (mg/dl)	47.8 ± 4.67	40.5 ± 10.9	40.2 ± 6.59	46.5 ± 5.79	0.785	0.870	0.035
Insulin (ng/ml)	0.224 ± 0.031	0.175 ± 0.039	0.306 ± 0.039 [*]	0.293 ± 0.073 [†]	<0.001	0.132	0.379
HOMA - IR	1.24 ± 0.199	1.07 ± 0.227	1.93 ± 0.304 [*]	2.03 ± 0.535 [†]	<0.001	0.800	0.342

Data are presented as mean ± standard deviation. Effects are the comparison by two-way ANOVA with Tukey's post-hoc test, with p<0.05 as statistically significance. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine; AST: aspartate aminotransferase; ALT: alanine aminotransferase; HDL: high density lipoprotein; n-HDL: non-high-density lipoprotein; HOMA-IR: homeostasis model assessment – insulin resistance. * vs Co; # vs MASLD; † vs Co+CAR.

3.4. Liver Tissue Analysis

3.4.1. Histological Parameters

Figure 2 shows the determination of hepatic steatosis by histological parameters. The MASLD induction group supplemented with carnosine (MASLD+CAR) attenuated the steatosis with macro and microvesicular aspects developed by the untreated MASLD induction group (MASLD). Also, picosirius red staining demonstrate septal fibrosis deposition (black arrow) in MASLD that is significantly reduced in MASLD treated with CAR.

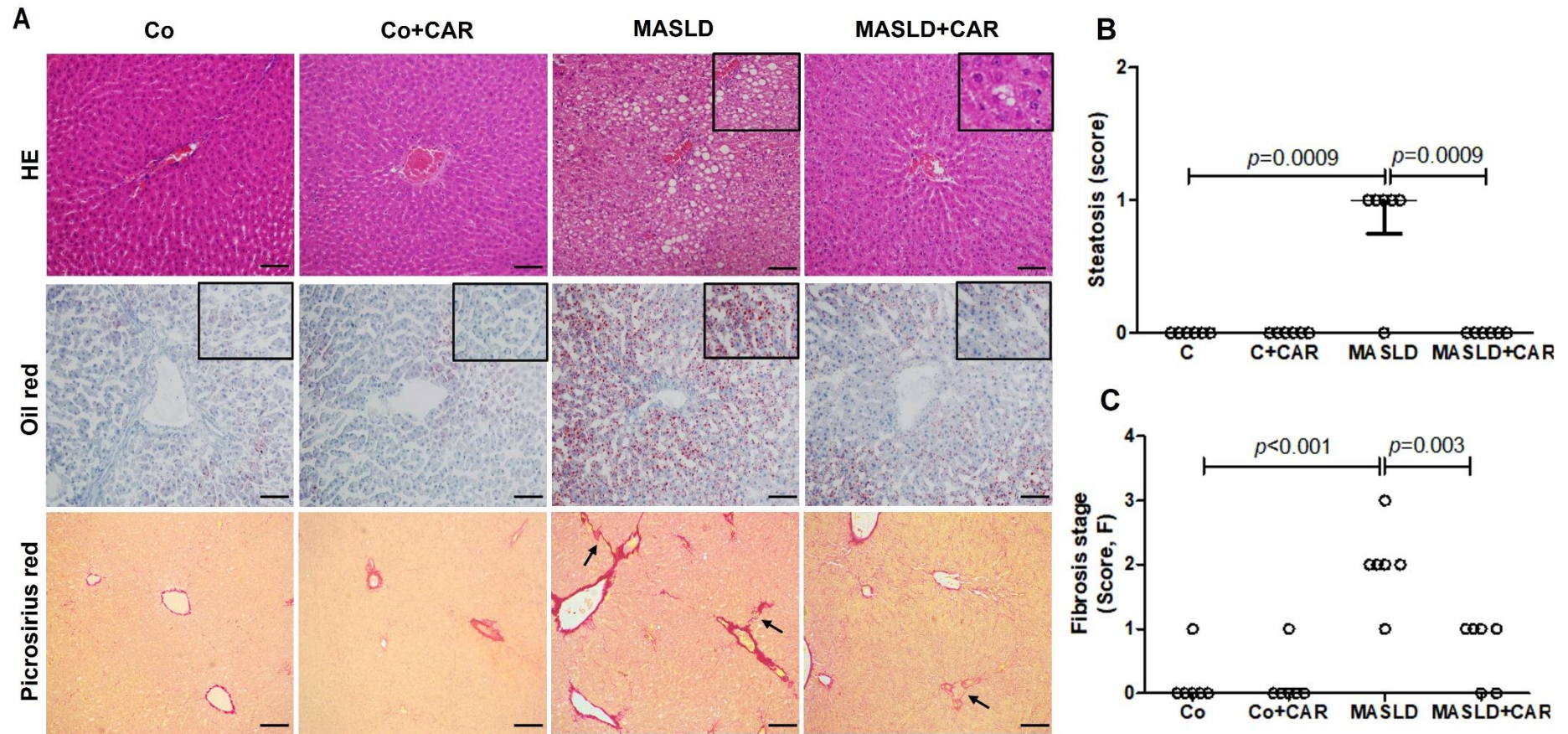


Figure 2. Determination of hepatic steatosis by histological parameters. A) Representative photomicrographs of liver sections stained with HE, Oil red and Picrosirius red (scale bar: 50 μ m); B) Steatosis Score; C) Fibrosis Stage. Data are presented as median (max – min) and individual data points, and were analyzed by Kruskal-Wallis with Tukey's post-hoc test. $p < 0.05$ statistically significant; $n = 6$ animals/group. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.4.2. Immunohistochemistry for CD68+

Figure 3 shows the effects of CAR supplementation on the density of CD68+ macrophages. The density of macrophages is lower in the MASLD+CAR group compared to the MASLD group, pointing to an attenuation of the inflammation generated by CAR.

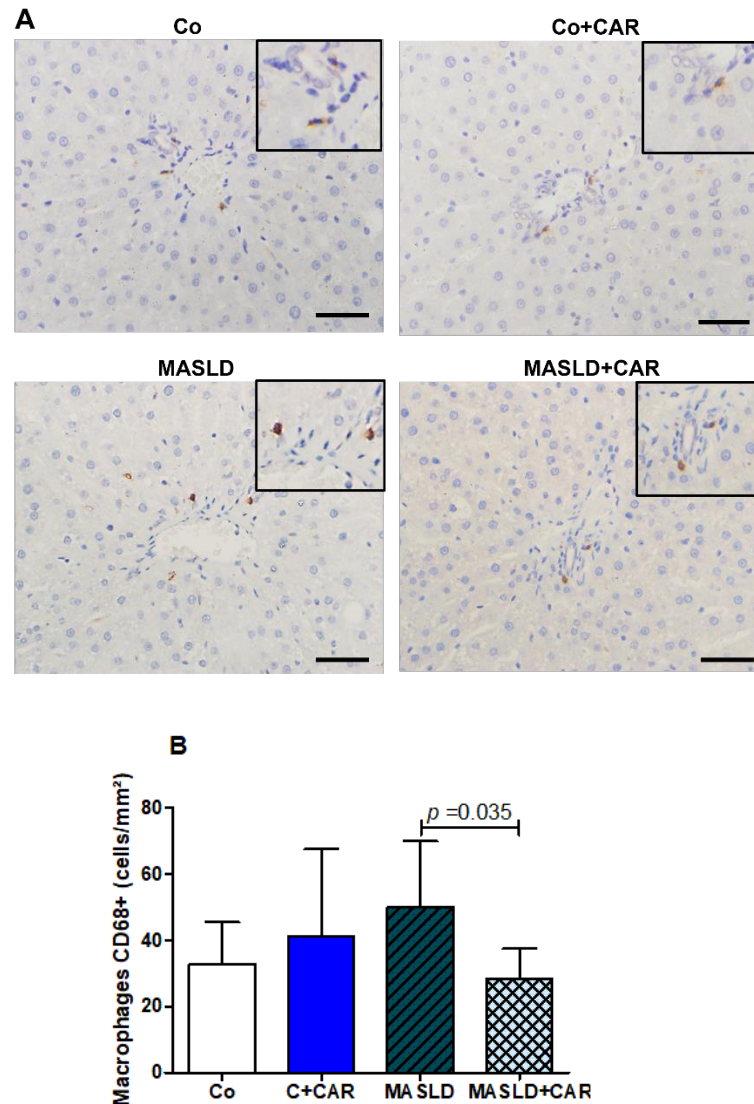
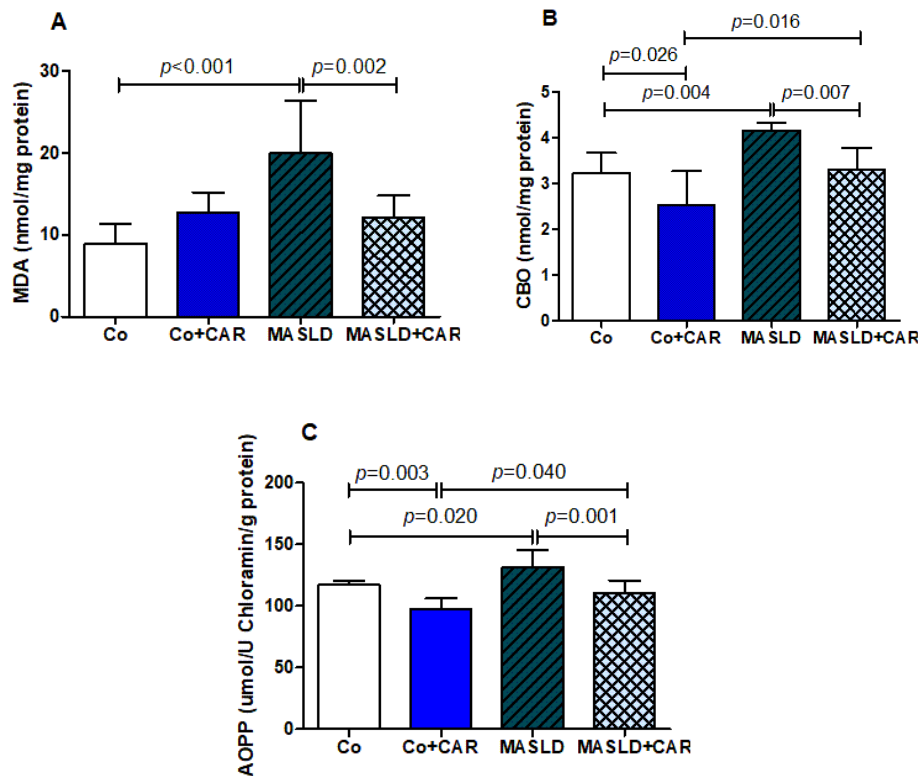


Figure 3. Effects of CAR treatment on CD68+ expression density. A) Representative photomicrographs of immunostained liver sections (details, scale bar: 100 μ m). B) Semi-quantitative analysis. Data are presented as mean $\pm$ standard deviation and individual data points, and were analyzed by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant; $n = 6$ animals/group. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.4.3. Oxidative Stress Markers

Figure 4 shows levels of MDA (Figure 4A), CBO (Figure 4B) and AOPP (Figure 4C) in liver tissue. The MASLD group showed higher levels of MDA, CBO, and AOPP compared to the Co group, and in contrast, the MASLD+CAR group attenuated MDA,

325 CBO, and AOPP levels compared to the MASLD group. Levels of CBO and AOPP of the
 326 MASLD+CAR group are elevated compared to the Co+CAR group.



328

329 **Figure 4.** Oxidative Stress Markers. A) Liver levels of MDA; B) Liver levels of CBO; C) Liver levels of AOPP. Data are
 330 presented as mean $\pm$ standard deviation. Comparison by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically
 331 significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group;
 332 MASLD+CAR: MASLD induction group treated with carnosine; MDA: Malondialdehyde; CBO: Protein carbonylation;
 333 AOPP: Advanced protein oxidation products.

334 3.4.4. Liver levels of triglycerides and cholesterol

335 Figure 5 shows the hepatic levels of triglycerides and cholesterol. The MASLD
 336 group had higher triglyceride and cholesterol concentrations in the liver than the Co
 337 group. The CAR-treated MASLD induction group (MASLD+CAR) had elevated
 338 triglyceride concentrations relative to the Co+CAR group but attenuated hepatic
 339 triglyceride and cholesterol levels compared to the untreated MASLD induction group.

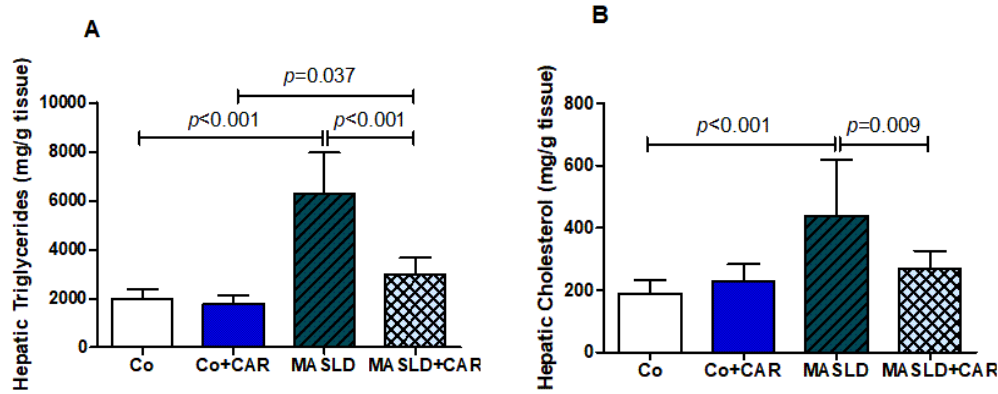


Figure 5. Liver levels of triglycerides and cholesterol. A) Hepatic levels of triglycerides; B) Hepatic cholesterol levels. Data are presented as mean $\pm$ standard deviation. Comparison by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.5. PPAR- α and VLCAD Gene Expressions

Figure 6 shows gene expression by real-time PCR in liver tissue. It can be seen that the MASLD group had lower PPAR- α and VLCAD mRNA levels compared to the Co group. In contrast, the MASLD+CAR group had higher levels of PPAR- α than MASLD group. MASLD+CAR also showed higher VLCAD mRNA levels than MASLD but not enough to get statistical significance.

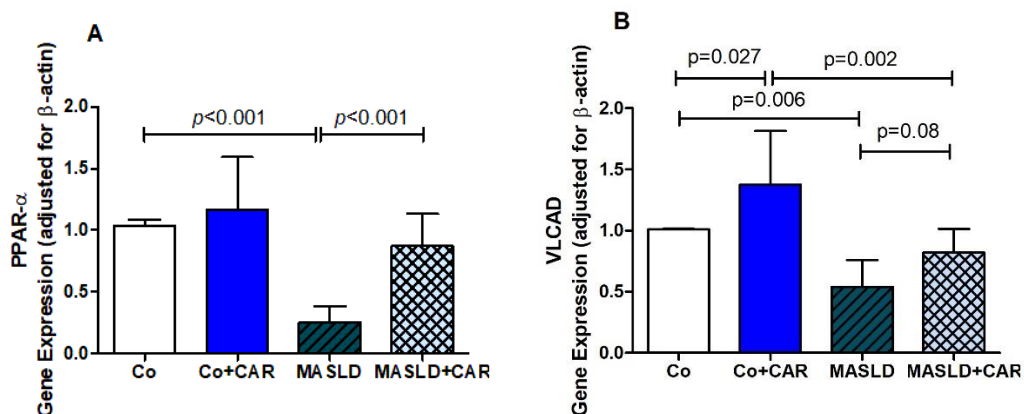


Figure 6. Gene expressions in liver tissue. A) PPAR- α Gene Expression; B) VLCAD Gene Expression. Comparison by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

4. Discussion

The present study aimed to verify the effects of CAR supplementation on diet-induced MASLD in rats. Studies have shown that the high consumption of diets rich in sugars and fats, as well as sugary drinks, favors the development of MASLD, a condition

associated with several morbidities, such as metabolic syndrome, insulin resistance, inflammation and oxidative stress (Carneros et al., 2020). In this condition, the present study demonstrated beneficial effects of CAR in the treatment of MASLD by influencing activity of key modulators of lipid metabolism, reflecting in lower lipid deposition and lower inflammatory and oxidative injuries in liver tissue.

MASLD was recently defined by the presence of hepatic steatosis associated with one of the five cardiometabolic risk factors: overweight or obesity, type 2 diabetes mellitus or hyperglycemia, hypertension or antihypertensive drug treatment, and dyslipidemia (Rinella et al., 2023). In this study, the MASLD group presented obesity (higher adiposity index), hypertension, insulin resistance and dyslipidemia, evidencing the presence of the condition. Higher water and caloric intakes were observed in the MASLD group, with a consequent increase in final body weight and weight gain, as observed in other similar studies (Garcia et al., 2022; Siqueira et al., 2022). Although the animals in the MASLD group had lower food intake than the Co group, they had higher sugar-sweetened-water (25% sucrose) consumption, justifying the weight gain and higher final weight of these animals. This dietary habit in association with physical inactivity generates a positive energy balance, which results in hypertrophy and hyperplasia of adipocytes. This change in adipose tissue causes accumulation of fatty acids as well as changes in the production of inflammatory cytokines (Garcia et al., 2019; White and Ravussin, 2019). When hepatocytes reach their lipid-storing capacity, lipotoxicity and hepatocellular stress lead to apoptosis, and then, chemokines are released to activate resident macrophages, kupffer cells, and circulating monocyte-derived macrophages (Li et al., 2020). Macrophages are essential factors in the progression of MASLD. Their activation and proliferation promote several stimuli for inflammatory perpetuation with consequent progression to MASH (Ju and Tacke, 2016). Oxidative stress is also a critical alteration that favors the development of MASLD since the hepatic accumulation of lipids favors the overproduction of ROS, which at high concentrations leads to a decrease in the activity of antioxidant systems, causing several structural and oxidative changes in the metabolism of cellular macromolecules (Bayarsaikhan et al., 2022; Gabbia et al., 2021; Hong et al., 2021). Hepatic storage of triglycerides and cholesterol is an initial mechanism to prevent toxicity; however, excess steatosis can induce lipotoxicity and glucotoxicity in the liver, producing ROS in various organelles (Sabir et al., 2022; Yang and Jo, 2018). Once established the pathophysiology of MASLD, the present study aimed to verify the effects of CAR treatment on markers for the development of this condition.

The treatment with CAR did not influence food, water and caloric intake, weight gain, and final weight of the animals in the treated MASLD induction group (MASLD+CAR), corroborating data found in the literature, where CAR supplementation did not influence food and water intake, as well as animal weight parameters (Everaert et al., 2021; Giriş et al., 2014). Treatment with CAR did not influence levels of adiposity index, hypertension, and total cholesterol in the MASLD group. Instead, it was observed attenuated plasma triglyceride levels, as well as improved HDL cholesterol concentrations. Al-Sawalha et al. (2019) conducted a study in rats with metabolic syndrome and CAR supplementation for 16 weeks, where changes similar to those found in our work on the lipid profile were also observed. The positive effects of CAR on lipid parameters were also observed in humans with type 2 diabetes (Houjehani et al., 2018) who received CAR for 12 weeks, demonstrating that CAR can act as an anti-dyslipidemic agent and may have protective effects in models of obesity-related complications.

The mechanism by which CAR intervenes on lipid profile is not well understood. In the present work, along with a better plasma lipid profile and lower deposition of lipids in liver tissue it was observed important modulatory activity on PPAR- α gene expression. PPAR- α is a key transcription factor that regulates transcription of genes involved in fatty acid β -oxidation and thermogenesis (Todisco et al., 2022). Moreover, in response of PPAR- α activation or not, it was observed a modulatory effect on mRNA expression of VLCAD, an enzyme responsible for the catalysis of the first reaction of the mitochondrial very long chain fatty acids β -oxidation (Koo, 2013). The oxidation of fatty acids, mainly regulated by PPAR- α , reduces the levels of fatty acids available in the liver, using them as an energy source. However, in the face of mitochondrial dysfunction, the β -oxidation process is impaired, thus generating fat accumulation, as well as the production of ROS (Arroyave-ospina et al., 2021; Chen et al., 2020; Mong et al., 2011). In the present study, the MASLD group showed lower PPAR- α gene expression than the Co group. CAR supplementation in the MASLD+CAR increased mRNA levels compared to the MASLD group, pointing to a possible role on this component. Additionally, mRNA levels of VLCAD showed similar behavior, demonstrating high response in Co+CAR and a restoring effect in the MASLD+CAR, even still not statistically significant. There are no data in the literature reporting the role of CAR on PPAR- α , but there are studies where β -alanine, a precursor of CAR, was able to stimulate the expression of PPAR- β/δ , a key regulator of muscle lipid balance, suggesting that histidyl dipeptides can be natural ligands of PPARs (Schnuck et al., 2016). In line with this, these findings suggest a possible role of CAR on dysfunctions of lipid metabolism, in view of the reduction of

plasma and hepatic triglycerides, hepatic cholesterol, steatosis and oxidative stress, as well as an increase in PPAR- α gene expression in MASLD.

Although no significant difference was observed between the MASLD group and the Co group for CD68 expression, CAR significantly attenuated their activation compared to the MASLD group. The reduction of macrophage infiltration in MASLD+CAR is in line with the reduction of collagen deposition marked by Sirius red staining and fibrosis score in the same group, suggesting a possible role of CAR in the attenuation of MASLD progression to MASH. Similar data were found in the literature, where CAR attenuated inflammatory infiltrates (monocytes and macrophages) (Chen et al., 2017). Another study reported a significant decrease in CD68 macrophages in mice with atherosclerotic lesions treated with CAR (Barski et al., 2013). CAR's action on macrophages may be related to its potential to reduce oxidative stress, which is directly related to inflammation. However, CAR may also be acting on MIF, generating a decrease in the activity of CD68 macrophages in the obesity model. Aiello et al., 2022 observed that CAR increased MIF in a 3D skin model, associating this effect with the antioxidant action of CAR and its power to reduce oxidative stress and its positive effect on aging.

In line with this hypothesis, the present work showed that the MASLD group had higher hepatic levels of MDA, carbonyl proteins (CBO), and AOPP in comparison to the control group, but, the treatment with CAR (MASLD+CAR group) attenuated the hepatic concentrations of MDA, CBO, and AOPP in comparison to the MASLD group. It is also possible to notice lower concentrations of CBO and AOPP in the Co+CAR group compared to the Co group, which may be related to the health and quality of the animals used in the study. These data corroborate those described in the literature, where CAR reduced the plasmatic and hepatic levels of AOPP, as well as the hepatic levels of MDA and CBO in rats with diabetes induced by a hypercaloric diet and streptozotocin, possibly due to its protective effect against stress oxidative, detoxifying reactive components and producing compounds less harmful to the oxidant system, as is the case of carnosinyl proteins, which inhibit the oxidative conversion of glycated proteins into advanced glycation end-products (AGEs) (Fatih Aydın et al., 2017; Hipkiss et al., 2001). Yılmaz et al., 2017 reported that CAR decreased plasma and hepatic levels of CBO and AOPP in rats with oxidative stress induced by methylglyoxal, associating this activity with its potential antioxidant effect, as an ion chelator and ROS scavenger, preventing modifications in lipids, proteins and nucleic acids caused by metal ions to generate free radicals or by ROS themselves. Studies have described that the imidazole ring present in the structure of CAR has an essential role in inhibiting the formation of reactive

species, in addition to the anti-glycation potential of CAR, preventing the formation of abnormal proteins that can generate AGEs (Boldyrev et al., 2013; Chmielewska et al., 2020).

5. Conclusion

The administration of CAR promoted changes in markers for the MASLD pathophysiology. The supplementation of CAR in MASLD was involved with the attenuation of plasma and hepatic lipid deposition, improvement of plasma HDL cholesterol and attenuation of oxidative damage. It was also observed modulation of the PPAR- α mRNA expressions, a key component in the control of lipid metabolism and oxidative functions in liver tissue. These findings suggest that the PPAR- α transcription factor can be a potential target for the CAR effects in MASLD. More studies are still needed to better elucidate and determine the exact way by means CAR induces PPAR- α modulation. For instance, we conclude that supplementation with CAR is effective for treating lipid metabolism dysfunctions and oxidative stress in MASLD.

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CRedit authorship contribution statement

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Effects of carnosine supplementation on markers for the pathophysiological development of metabolic dysfunction-associated steatotic liver disease in a diet-induced model

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Abstract

Consumption of diets high in sugar and fat is related to the development of Metabolic dysfunction-associated steatotic liver disease (MASLD). Carnosine (CAR) is a dipeptide with antioxidant and anti-inflammatory action and has been studied for treating diseases. This work aimed to evaluate the effects of CAR on diet-induced MASLD in rats. Male Wistar rats were distributed into 2 groups (17 weeks): normocaloric (Co, n=12), and hypercaloric diet rich in lipids and simple carbohydrates (MASLD, n=12). After, the animals were redistributed to begin the treatment with CAR (4 weeks): Co (n=6), Co + CAR (n=6), MASLD (n=6), and MASLD + CAR (n=6), administered intraperitoneally (250mg/kg). Evaluations included nutritional, hormonal and metabolic parameters; hepatic steatosis, inflammatory and oxidative markers. MASLD group had a higher adiposity index, systolic blood pressure, glucose, plasma and liver triglycerides and cholesterol, insulin, hepatic steatosis, oxidative markers, and lower PPAR- α (Peroxisome Proliferator-activated receptor α), compared to the Co. CAR attenuated plasma and hepatic triglyceride and cholesterol levels, hepatic steatosis, CD68+ macrophages, and hepatic oxidative markers, in addition to increasing HDL cholesterol levels and PPAR- α , compared to the untreated MASLD group. CAR acts in important pathophysiological processes of MASLD and may be a therapeutic compound to control the disease.

Keywords: Metabolic dysfunction-associated steatotic liver disease, carnosine, obesity, hypercaloric diet.

1. Introduction

Non-alcoholic fatty liver disease (NAFLD) is the most common liver injury, which affects about a third of the world's population and is defined by the accumulation of fat in liver tissue without alcohol consumption and other liver damage (Younossi et al., 2016). In 2023, a new terminology and definition were proposed, starting to be called Metabolic dysfunction-associated steatotic liver disease (MASLD). Therefore, the MASLD is based on hepatic steatosis with one of the five cardiometabolic risk factors: overweight or obesity, type 2 diabetes mellitus or hyperglycemia, hypertension or antihypertensive drug treatment, and dyslipidemia (Rinella et al., 2023).

MASLD has a spectrum of evolution from simple steatosis and can involve Metabolic dysfunction-associated steatohepatitis (MASH) and even liver cancer (Carneros et al., 2020; Francisqueti-Ferron et al., 2022; Rinella et al., 2023). According to a survey based on data from Europe, North America, and Asia, the prevalence of patients with MASLD is around 39% (Lim et al., 2021), while MASH affects around 5% of the world's population (Younossi et al., 2016). Its pathogenesis is still not well defined, but there are two theories for its development: the first is the "two hits" theory, where it is believed that the first aggression is the imbalance caused by insulin resistance (IR) between the hepatic pathways of acquisition and elimination of lipids, and the second attack involves the processes of inflammation and oxidation (Day and James, 1998). The second theory is that its development takes place from a "multiple hits" model, where several aggressive agents trigger this injury, being genetic, epigenetic, nutritional, and inflammatory, including changes in the microbiota, IR, and oxidative stress (Buzzetti et al., 2016; Day and James, 1998).

The Western diet, rich in simple carbohydrates and fats, is a crucial precursor factor of MASLD, which can cause harmful effects on the liver by accumulating these macronutrients (Rives et al., 2020; Vancells Lujan et al., 2021). Data in the literature indicate that women who consume a diet high in sugar and fat have a 119% greater risk (OR, 2.19; 95% CI, 1.40–3.46) of developing fatty liver disease (Jia et al., 2015); similar data were found for adolescents aged 17 years who had the same Western diet consumption pattern (Oddy et al., 2013). The accumulation of fatty acids in the liver generates hepatic steatosis. It favors the excessive production of ROS (reactive oxygen species), causing an imbalance in the antioxidant system and, consequently, oxidative

stress, which can cause various cellular damage (Arroyave-ospina et al., 2021; Yki-Järvinen et al., 2021).

Given the complexity of MASLD and its consequences, the search for forms of prevention and treatment is necessary. Currently, there is no specific pharmacological treatment for this disease, only drugs that treat specific metabolic alterations, such as dyslipidemia and hyperglycemia, are used (Nassir, 2022; Salvoza et al., 2022). On the other hand, the search for bioactive compounds that can be used to improve these conditions emerges, and one of these compounds is carnosine (CAR), a natural dipeptide composed of β -alanine and L-histidine, which can be found in the human body, in higher concentrations in skeletal muscle and brain (Aldini et al., 2021).

For many years CAR was the focus of studies only related to improving the performance of physical exercises, mainly due to its buffering capacity in the muscles (Matthews et al., 2019). However, CAR has been identified as a potential therapeutic agent in treating diseases due to its antioxidant, anti-inflammatory, and ROS-sequestering activity (Chmielewska et al., 2020). Some studies demonstrated that CAR was able to attenuate the development of type 2 diabetes in ob/ob mice (Albrecht et al., 2017), improve parameters related to metabolic syndrome induced by diet in Wistar rats (Al-Sawalha et al., 2019), as well as improve hepatic steatosis in mice that consumed a high-fat diet (Mong et al., 2011) and in a model of type 2 diabetes in db/db mice (Schwank-Xu et al., 2021). Faced with the need to clarify the biological effects of CAR and the scarcity of studies relating this dipeptide to MASLD, the present study aimed to verify the effects of CAR supplementation on markers of the MASLD pathophysiology induced by diet in rats.

2. Material and Methods

2.1. Animals and Experimental Protocol

This study was approved by the Ethics Committee on Animal Use (n° 1322/2019). It was carried out following the Guide for the Care and Use of Laboratory Animals published by the U.S. National Institutes of Health. Male Wistar rats from the Central Animal Facility of the Faculty of Medicine of Botucatu, approximately 21 days old, were kept in individual cages in an environment with temperature ($24\pm 2^\circ\text{C}$), humidity ($55\pm 5\%$) and light (light cycle-dark 12h) controlled. Initially, they were distributed into two experimental groups, fed different diets for MASLD induction: normocaloric (n=12) and hypercaloric rich in sugar and fat (n=12). On the 17th week, after the difference between plasma triglycerides, defined as an inclusion criterion, the animals were redistributed into

four groups for treatment: a control group that continued receiving the normocaloric diet (co, n=6), a control group that received the normocaloric diet treated with carnosine (Co+CAR, n=6), a group that continued receiving the hypercaloric diet (MASLD, n=6) and a group that received the hypercaloric diet treated with carnosine (MASLD+CAR, n=6), for four weeks. Feed and water were offered ad libitum. The animals that received the hypercaloric diet also received sucrose in their drinking water at a concentration of 25% throughout the experiment. Both diets were produced according Francisqueti et al. (2017), and are described in Table 1. The groups received CAR supplementation (250 mg/kg/day) or vehicle (sterile saline solution, 0.9% NaCl) intraperitoneally for four weeks. At the end of the 21 weeks, the animals fasted for 8 hours and then were anesthetized (Thiopental 120 mg/kg/i.p.) and euthanized by decapitation. Blood samples, adipose tissue, and liver were collected.

Table 1. Diet composition and nutritional values.

Components	Normocaloric	Hypercaloric
Soybean meal (g/kg)	335	340
Sorghum (g/kg)	278	80
Soy hulls (g/kg)	188,5	116,7
Dextrin (g/kg)	146,5	20
Maize starch (g/kg)	278	80
Sucrose (g/kg)	-	80
Fructose (g/kg)	-	180
Soybean oil (g/kg)	23	-
Lard (g/kg)	-	154,3
Minerals (g/kg)	25	25
Salt (g/kg)	4	4

2.2. Carnosine Supplementation

CAR (L-carnosine) was administered intraperitoneally (250 mg/kg) for four weeks after induction. The dose was defined based on studies where CAR presented effective results and did not generate toxic and/or adverse effects on the animals (Aydın et al., 2017; Kalaz et al., 2016). The infusion was performed using a 30G needle, with a liquid volume close to 500 µL being injected. The product CarnoPure™ (Flamma S.p.A., Chignolo d'Isola, Bergamo, Italy) was used exclusively for scientific experiments and used in studies abroad. This product was kindly provided by this company, free of charge, for carrying out this experiment.

2.3. Nutritional characterization and obesity

Nutritional characterization was evaluated based on feed, water, and caloric intake (feed and water). Food and water intake were calculated daily from the individual leftovers of each animal. Caloric intake was calculated through food intake multiplied by the energy value of each diet. The characterization of obesity was based on weight gain and adiposity index. Weight gain was calculated by the difference between the final weight and the initial weight of the animals [weight gain (g)=final weight (g) - initial weight (g)]. The adiposity index represents the ratio between the sum of fat deposits (visceral, epididymal, and retroperitoneal) and the final weight of the animals (adiposity index = sum of fat deposits (g)/final weight (g) x 100) (Costa et al., 2019).

2.4. Systolic Blood Pressure

Systolic blood pressure (SBP) was measured by the indirect method of tail plethysmography using an electric sphygmomanometer Narco Bio-System® PE 300, model 709-0610 (International Biomedical, Inc, Houston, Texas, USA). For this purpose, the animals were kept for 5 minutes in a wooden box heated between 38°C and 40°C to stimulate arterial vasodilation. Afterward, the cuff was attached to the tail of each animal and inflated (200 mmHg), and deflated, allowing the recording of arterial pulses collected in a Gould RS 3200 polygraph (Gould Instrumenta Valley View, Ohio, USA). The mean of the three PAS readings was recorded for each animal (Siqueira et al., 2022).

2.5. Biochemical and Hormonal Analysis

Glucose was determined using a glucometer (Accu-Check Performa; Roche Diagnostics, Indianapolis, IN, USA). The concentrations of urea, creatinine, uric acid, triglycerides, total cholesterol and its fractions, HDL cholesterol and non-HDL cholesterol (n-HDL) were analyzed by the enzymatic colorimetric method with commercial kits (BioClin®, Belo Horizonte, MG, Brazil) in automated biochemical equipment (Chemistry Analyzer BS-200, Mindray Medical International Limited, Shenzhen, China).

The insulin level was measured using the indirect enzyme immunoassay (ELISA) method using commercial kits (Linco Research Inc., R&D Systems, Millipore, and B-Bridge International Inc.), and the reading was performed on a Spectramax 190 microplate spectrophotometer (Molecular Devices®, Sunnyvale, CA, USA). The IR assessment was performed using the Homeostasis Model Index - Insulin Resistance (HOMA-IR) using glycemia and plasma insulin parameters. The calculation of the HOMA index obeyed the following formula: insulin ($\mu\text{L U/mL}$) x blood glucose (mmol/L) / 22.5 (Francisqueti-Ferron et al., 2022).

2.6. Liver Tissue Analysis

2.6.1. Histological Parameters

After collection, liver tissue was stored in 4% paraformaldehyde with 0.1M phosphate buffer (pH=7.4) during the first 24 hours. Afterward, the tissue was transferred

to ethyl alcohol, and the samples were embedded in paraffin to obtain five μm -thick sections. Such sections (one per animal) were stained with Hematoxylin & Eosin (HE) and evaluated for steatosis, taking into account the macro and microvesicular aspects, according to the NAFLD Activity Score (NAS) parameter (Liang et al., 2014). Then, ten fields were randomly selected (20x objective), which were evaluated in each section, and the following scores were considered: 0 (<5% of the microscopic field), 1 (5-33%), 2 (33-66%), or 3 (>66%). Other liver samples were frozen and included in an Optimal Cutting Temperature medium (Tissue-Trek, USA), and 10 μm sections were obtained and stained with Oil red (Sigma Aldrich, USA) to obtain representative photos. To assess the presence and deposition of collagen sections samples were stained with picosirius red with subsequent detection by refringent images on optical microscope. The METAVIR score system was used to evaluate the level of fibrosis: F0: no fibrosis; F1: portal fibrosis without septa; F2: portal fibrosis with septa; F3: septa without cirrhosis; F4: cirrhosis.

2.6.2. Immunohistochemistry for CD68

Other 5 μm thick liver sections were submitted to antigen retrieval in 0.01M citrate buffer (pH 6.0, 120°C, five min.) in a Pascal pressure chamber (Dako Cytomation, Denmark). After blocking endogenous peroxidase with 10% H₂O₂ in phosphate-buffered saline (PBS) (15 minutes), the slides were treated with skimmed milk for 60 minutes to block nonspecific binding. Then, the histological sections were incubated in a humid chamber (4°C overnight) with a primary anti-CD68 antibody solution (macrophages, ab125212, 1:500 dilution, Abcam, UK). Then, the slides were incubated with HRP polymer (Horseradish peroxidase) in a single step (EasyPath - Erviegas, Brazil) (20 minutes at room temperature). The reactions were revealed with the chromogen 3'-diaminobenzidine (DAB) (Sigma Aldrich, USA), and the slides were counterstained with Harris hematoxylin. CD68+ macrophages were counted in 10 randomly selected fields (20x) and divided by field area (mm²) using Image J software (NIH, USA) (Romualdo et al., 2022).

2.6.3. Oxidative Stress markers

Malondialdehyde (MDA) level was used to assess lipid peroxidation. Briefly, 250 μL of supernatant from hepatic tissue was added to 750 μL of 10% trichloroacetic acid for protein precipitation. The samples were centrifuged (3000 rpm, for 5 minutes; Eppendorf® Centrifuge 5804-R, Hamburg, Germany), and the supernatant was removed. Thiobarbituric acid (ATB) was added at a ratio of 0.67% (1:1), and the samples were heated for 15 minutes at 100°C. After cooling, the reading was performed at 535 nm in the Spectra Max 190 microplate reader (Molecular Devices®, Sunnyvale, CA, USA). The MDA concentration was obtained by the molar extinction coefficient (1.56 x

105 M⁻¹cm⁻¹) and the absorption of the samples, and the final result was reported in nmol/g of protein (Siqueira et al., 2023).

To quantify protein carbonylation (CBO), 100 µL of liver tissue homogenate was used for 100 µL of 2,4-dinitrophenylhydrazine (DNPH, 10mM in 2M HCl). The samples were incubated for 10 minutes at room temperature, and then 50 µL of sodium hydroxide (6M NaOH) was added and incubated again for 10 minutes at room temperature. Reading at 450 nm was performed on a Spectramax 190 microplate reader (Molecular Devices®, Sunnyvale, CA, USA). The results were obtained by the molar extinction coefficient (22,000 M⁻¹ cm⁻¹) of DNPH and expressed in nmol/g of protein. Protein carbonylation levels were expressed in nmol of DNPH/mg of protein (Garcia et al., 2022).

Advanced oxidation protein products (AOPP) were detected in liver tissue. Tissue samples were homogenized in PBS buffer at 1:10 (sample:PBS). AOPP quantification was performed by diluting 1:5 of 200 µl of the sample (homogenate). 200 µl of Chloramine-T (0-100 umol/L) was used for the calibration curve, and 200 µl of PBS as a blank. 10 µl of 1.16M KI (potassium iodide) and 20 µl of acetic acid were added to the samples. Absorbance was read immediately at 340 nm (Multiskan Ascent spectrophotometer, Labsystems, Vantaa, Finland). The final concentration of AOPP is expressed in chloramin units (umol/L), and the result in the liver tissue is corrected for the amount of protein (Ferron et al., 2019).

2.6.4. Hepatic Levels of Triglycerides and Cholesterol

Intracellular levels of hepatic triglycerides and cholesterol were determined by a colorimetric method (Colorimetric Triglycerides and Cholesterol Assay Kit; Cayman Chemical, Ann Arbor, MI, USA). The results were corrected for the amount of tissue (Francisqueti-Ferron et al., 2022).

2.7. Gene Expressions

Gene expression of peroxisome proliferator-activated receptor α (PPAR-α; gene *Ppara*, primers F-AGATGGGCTGGTTCTTCCTC and R-CAGTGCATTTGCCAGG), very long-chain acyl-CoA dehydrogenase (VLCAD, gene *Acadvl*, primers F-ATGGAATTCATGGCCAGTGAGGCCACTC and R-TTTACGAATTCTCAGACTCTAAGGGGGT) and β-actin (gene *Actb*, primers F-GGGAAATCGTGCGTGACAT and R-CAGGAGGAGCAATGATCTT) were performed using real time PCR (polymerase chain reaction). Total RNA (ribonucleic acid) was extracted from liver tissue using the TRIzol® reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's specifications. The RNA was subjected to reverse transcription for conversion to cDNA (complementary deoxyribonucleic acid) (High-Capacity cDNA Reverse Transcription Kit, Applied Biosystems, CA, USA) using the

provided primers (Exxtend Biotecnologia Ltda, Campinas-SP, Brazil). The real-time PCR kit used the cDNA (TagMan® Gene Expression Assays, Applied Biosystems, Foster City, CA, USA) to evaluate expressions. The β -actin was used to normalize expressions of PPAR- α and VLCAD.

2.8. Statistical Analysis

Results are expressed as mean $\pm$ standard deviation. Two-way ANOVA determined differences between groups, and data were subjected to Tukey's post hoc test to compare all groups. Histological parameters are expressed as medians, and Kruskal-Wallis was used with Tukey's post hoc test. Statistical analyzes were performed using Sigma Plot 12.0 software for Windows (Systat Software Inc., San Jose, CA, USA). A p-value <0.05 was considered statistically significant and all performed statistical analyzes were tested to ensure the power to detect differences among groups and the sample size.

3. Results

3.1. Nutritional characterization and obesity

The nutritional characterization and obesity is presented in Table 2. It is possible to observe that the MASLD and MASLD+CAR groups had lower food intake, higher water, caloric intake, and more significant weight gain when compared to the Co and Co+CAR groups, respectively. There was no difference in the initial body weight between groups. The MASLD group had a higher final weight when compared to the Co group. The adiposity index determined the presence of obesity, and it can be seen that the MASLD induction groups (MASLD and MASLD+CAR) had higher adiposity indices when compared to the control groups (Co and Co+CAR, respectively).

Table 2. Nutritional characterization and obesity.

	Groups				Effects		
	Co	Co+CAR	MASLD	MASLD+CAR	Diet	CAR	Interactions
Food intake (g/day)	37.0 $\pm$ 2.31	36.3 $\pm$ 2.07	25.6 $\pm$ 0.904*	25.3 $\pm$ 1.72†	<0.001	0.528	0.858
Water intake (ml/day)	53.2 $\pm$ 3.79	52.0 $\pm$ 7.17	68.0 $\pm$ 4.02*	64.9 $\pm$ 3.48†	<0.001	0.320	0.564
Caloric intake (kcal/day)	133 $\pm$ 8.28	130 $\pm$ 7.42	166 $\pm$ 7.84*	168 $\pm$ 5.44†	<0.001	0.849	0.584
Initial body weight (g)	324 $\pm$ 25.0	324 $\pm$ 26.8	320 $\pm$ 16.4	314 $\pm$ 27.1	0.450	0.775	0.753
Final body weight (g)	502 $\pm$ 57.2	477 $\pm$ 25.8	569 $\pm$ 38.0*	530 $\pm$ 68.5	0.008	0.133	0.743
Weight gain (g)	145 $\pm$ 48.9	137 $\pm$ 23.2	250 $\pm$ 34.6*	198 $\pm$ 48.6†	<0.001	0.085	0.201
Adiposity index (%)	4.15 $\pm$ 1.60	3.12 $\pm$ 1.11	8.51 $\pm$ 1.03*	7.68 $\pm$ 2.56†	<0.001	0.19	0.883

Data are presented as mean $\pm$ standard deviation. Effects are the comparison by two-way ANOVA with Tukey's post-hoc test, with $p < 0.05$ as statistically significance. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine. * vs Co; † vs Co+CAR.

3.2. Systolic Blood Pressure

Figure 1 shows the Systolic Blood Pressure (SBP) levels. The MASLD group had a higher SBP compared to the Co group.

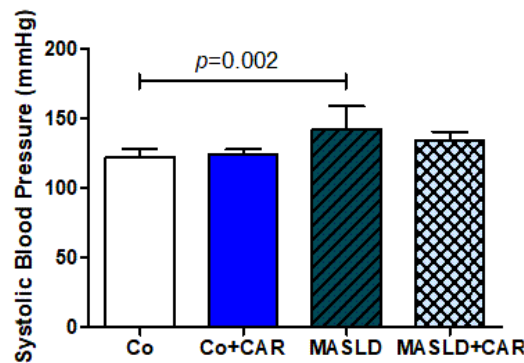


Figure 1. Systolic Blood Pressure Levels. Data are presented as mean $\pm$ standard deviation. Comparison by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.3. Biochemical and hormonal analysis

Table 3 shows the biochemical and hormonal markers. The MASLD and MASLD+CAR groups had lower urea levels when compared to the Co and Co+CAR groups, respectively. There was no significant difference uric acid, and creatinine. ALT levels were significantly lower in control treated with CAR and MASLD induction groups (MASLD and MASLD+CAR) had higher blood glucose levels when compared to the control groups (Co and Co+CAR, respectively). Co+CAR showed lower concentrations of total cholesterol compared to the Co group, while MASLD+CAR showed higher concentrations of total cholesterol compared to the Co+CAR group. Regarding triglycerides, the MASLD and Co+CAR groups had higher and lower concentrations when compared to the Co group. In contrast, the MASLD+CAR group had lower and higher levels when compared to the MASLD and Co+CAR groups, respectively. In addition, the MASLD+CAR group had higher HDL cholesterol (high-density lipoprotein) levels than the MASLD group. There was no statistical difference for non-HDL cholesterol levels. The MASLD and MASLD+CAR groups had higher insulin and HOMA-IR values when compared to the Co and Co+CAR groups, respectively.

Table 3. Biochemical and hormonal markers.

	Groups				Effects		
	Co	Co+CAR	MASLD	MASLD+CAR	Diet	CAR	Interactions
AST (U/l)	155 $\pm$ 34.3	125 $\pm$ 18.6	114 $\pm$ 21.0	130 $\pm$ 32.4	0.128	0.532	0.058
ALT (U/l)	104 $\pm$ 14.4	69.5 $\pm$ 16.1*	83.0 $\pm$ 26.5	87.5 $\pm$ 21.5	0.858	0.084	0.028
Uric acid (mg/dl)	0.967 $\pm$ 0.175	0.800 $\pm$ 0.420	1.27 $\pm$ 0.761	1.67 $\pm$ 1.69	0.150	0.768	0.476

Urea (mg/dl)	58.8 ± 8.66	61.0 ± 9.63	35.0 ± 15.5*	44.2 ± 4.58†	<0.001	0.195	0.417
Creatinine (mg/dl)	4.48 ± 3.08	4.59 ± 5.01	4.48 ± 1.68	4.36 ± 0.985	0.931	0.998	0.930
Blood Glucose (mg/dl)	78.0 ± 8.29	86.3 ± 11.4	88.8 ± 6.74*	97.3 ± 4.50†	0.004	0.020	0.980
Total Cholesterol (mg/dl)	62.3 ± 7.12	46.5 ± 10.1*	52.7 ± 9.63	62.0 ± 8.32†	0.430	0.380	0.002
Triglycerides (mg/dl)	70.4 ± 16.9	41.0 ± 8.32*	205 ± 34.6*	112 ± 27.4#†	<0.001	<0.001	0.004
HDL (mg/dl)	14.5 ± 3.02	14.0 ± 1.79	11.4 ± 2.33	15.5 ± 3.27#	0.471	0.114	0.047
n-HDL (mg/dl)	47.8 ± 4.67	40.5 ± 10.9	40.2 ± 6.59	46.5 ± 5.79	0.785	0.870	0.035
Insulin (ng/ml)	0.224 ± 0.031	0.175 ± 0.039	0.306 ± 0.039*	0.293 ± 0.073†	<0.001	0.132	0.379
HOMA - IR	1.24 ± 0.199	1.07 ± 0.227	1.93 ± 0.304*	2.03 ± 0.535†	<0.001	0.800	0.342

Data are presented as mean ± standard deviation. Effects are the comparison by two-way ANOVA with Tukey's post-hoc test, with p<0.05 as statistically significance. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine; AST: aspartate aminotransferase; ALT: alanine aminotransferase; HDL: high density lipoprotein; n-HDL: non-high-density lipoprotein; HOMA-IR: homeostasis model assessment – insulin resistance. * vs Co; # vs MASLD; † vs Co+CAR.

3.4. Liver Tissue Analysis

3.4.1. Histological Parameters

Figure 2 shows the determination of hepatic steatosis by histological parameters. The MASLD induction group supplemented with carnosine (MASLD+CAR) attenuated the steatosis with macro and microvesicular aspects developed by the untreated MASLD induction group (MASLD). Also, picosirius red staining demonstrate septal fibrosis deposition (black arrow) in MASLD that is significantly reduced in MASLD treated with CAR.

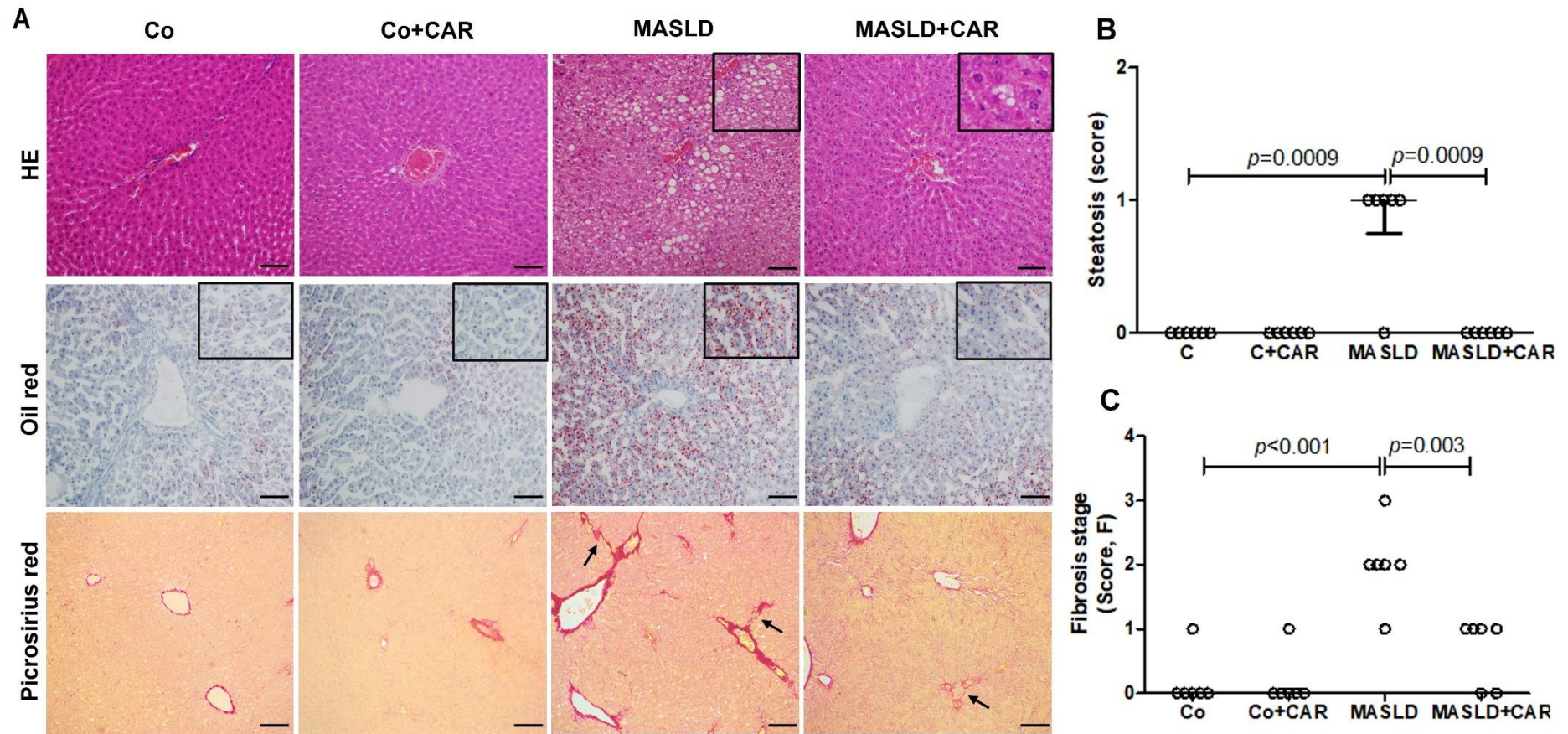


Figure 2. Determination of hepatic steatosis by histological parameters. A) Representative photomicrographs of liver sections stained with HE, Oil red and Picrosirius red (scale bar: 50 μ m); B) Steatosis Score; C) Fibrosis Stage. Data are presented as median (max – min) and individual data points, and were analyzed by Kruskal-Wallis with Tukey's post-hoc test. $p < 0.05$ statistically significant; $n = 6$ animals/group. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.4.2. Immunohistochemistry for CD68+

Figure 3 shows the effects of CAR supplementation on the density of CD68+ macrophages. The density of macrophages is lower in the MASLD+CAR group compared to the MASLD group, pointing to an attenuation of the inflammation generated by CAR.

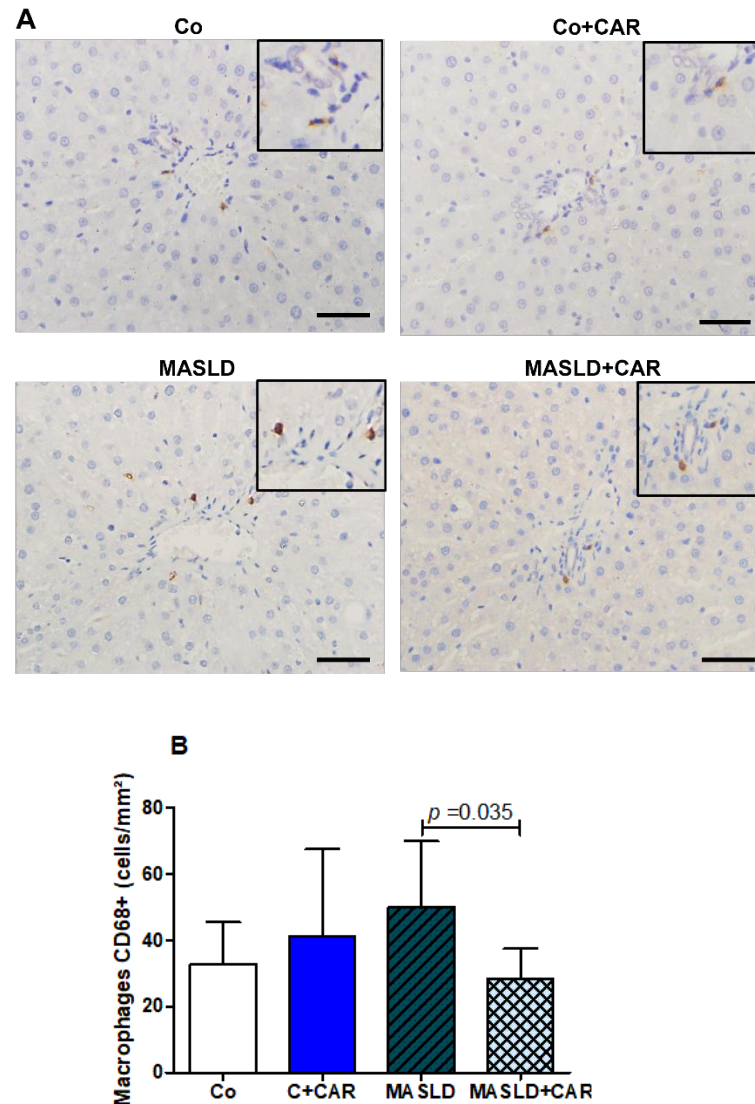


Figure 3. Effects of CAR treatment on CD68+ expression density. A) Representative photomicrographs of immunostained liver sections (details, scale bar: 100 μ m). B) Semi-quantitative analysis. Data are presented as mean $\pm$ standard deviation and were analyzed by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant; $n = 6$ animals/group. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.4.3. Oxidative Stress Markers

Figure 4 shows levels of MDA (Figure 4A), CBO (Figure 4B) and AOPP (Figure 4C) in liver tissue. The MASLD group showed higher levels of MDA, CBO, and AOPP compared to the Co group, and in contrast, the MASLD+CAR group attenuated MDA,

CBO, and AOPP levels compared to the MASLD group. Levels of CBO and AOPP of the MASLD+CAR group are elevated compared to the Co+CAR group.

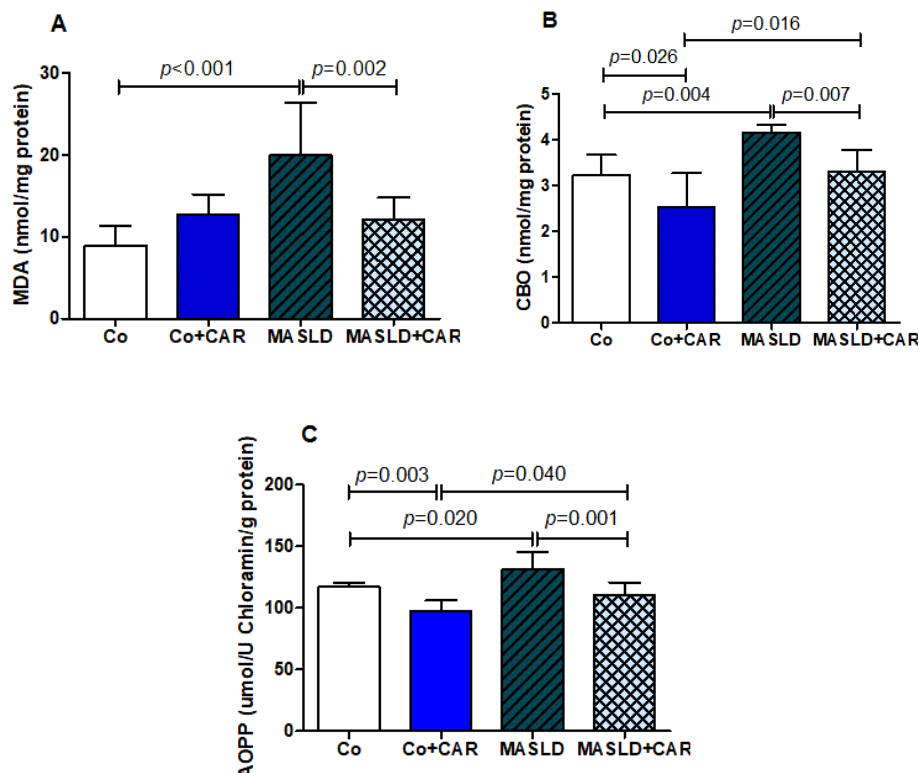


Figure 4. Oxidative Stress Markers. A) Liver levels of MDA; B) Liver levels of CBO; C) Liver levels of AOPP. Data are presented as mean $\pm$ standard deviation. Comparison by two-way ANOVA with Tukey's post-hoc test. $p < 0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine; MDA: Malondialdehyde; CBO: Protein carbonylation; AOPP: Advanced protein oxidation products.

3.4.4. Liver levels of triglycerides and cholesterol

Figure 5 shows the hepatic levels of triglycerides and cholesterol. The MASLD group had higher triglyceride and cholesterol concentrations in the liver than the Co group. The CAR-treated MASLD induction group (MASLD+CAR) had elevated triglyceride concentrations relative to the Co+CAR group but attenuated hepatic triglyceride and cholesterol levels compared to the untreated MASLD induction group.

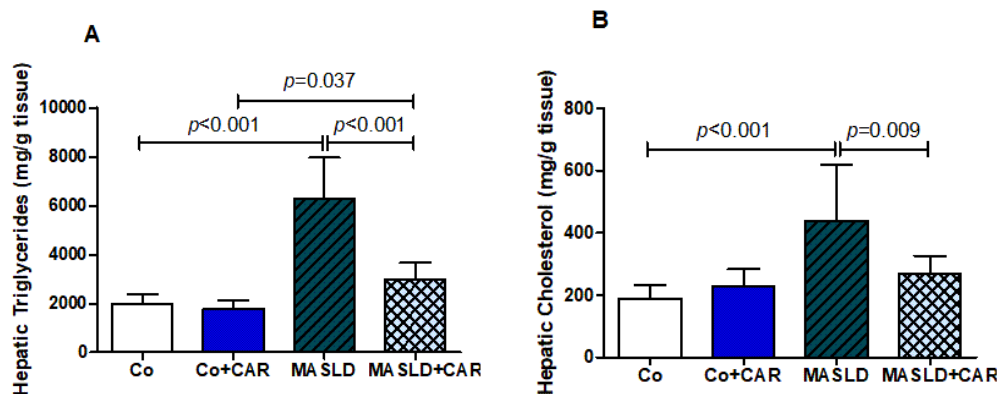


Figure 5. Liver levels of triglycerides and cholesterol. A) Hepatic levels of triglycerides; B) Hepatic cholesterol levels. Data are presented as mean $\pm$ standard deviation. Comparison by two-way ANOVA with Tukey's post-hoc test. $p<0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

3.5. PPAR- α and VLCAD Gene Expressions

Figure 6 shows gene expression by real-time PCR in liver tissue. It can be seen that the MASLD group had lower PPAR- α and VLCAD mRNA levels compared to the Co group. In contrast, the MASLD+CAR group had higher levels of PPAR- α than MASLD group. MASLD+CAR also showed higher VLCAD mRNA levels than MASLD but not enough to get statistical significance.

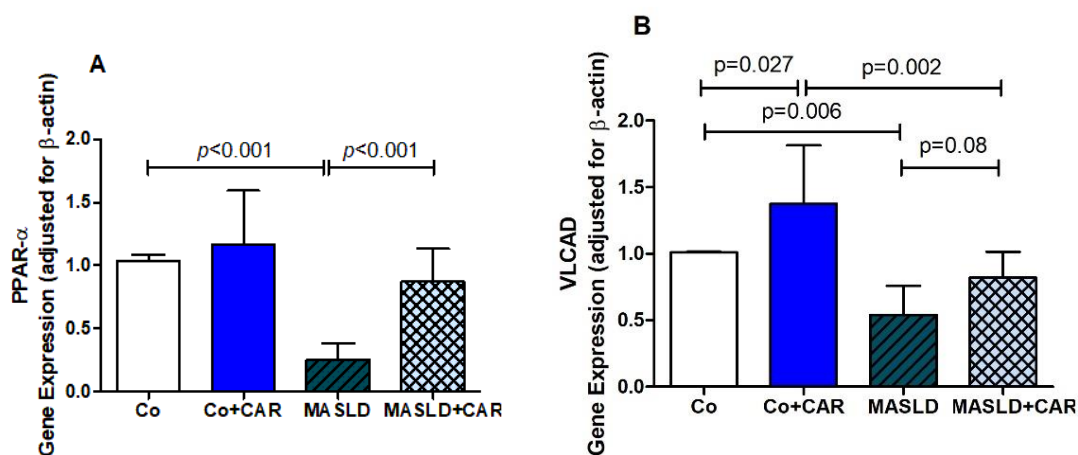


Figure 6. Gene expressions in liver tissue. A) PPAR- α Gene Expression; B) VLCAD Gene Expression. Comparison by two-way ANOVA with Tukey's post-hoc test. $p<0.05$ statistically significant. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine.

4. Discussion

The present study aimed to verify the effects of CAR supplementation on diet-induced MASLD in rats. Studies have shown that the high consumption of diets rich in

sugars and fats, as well as sugary drinks, favors the development of MASLD, a condition associated with several morbidities, such as metabolic syndrome, insulin resistance, inflammation and oxidative stress (Carneros et al., 2020). In this condition, the present study demonstrated beneficial effects of CAR in the treatment of MASLD by influencing activity of key modulators of lipid metabolism, reflecting in lower lipid deposition and lower inflammatory and oxidative injuries in liver tissue.

MASLD was recently defined by the presence of hepatic steatosis associated with one of the five cardiometabolic risk factors: overweight or obesity, type 2 diabetes mellitus or hyperglycemia, hypertension or antihypertensive drug treatment, and dyslipidemia (Rinella et al., 2023). In this study, the MASLD group presented obesity (higher adiposity index), hypertension, insulin resistance and dyslipidemia, evidencing the presence of the condition. Higher water and caloric intakes were observed in the MASLD group, with a consequent increase in final body weight and weight gain, as observed in other similar studies (Garcia et al., 2022; Siqueira et al., 2022). Although the animals in the MASLD group had lower food intake than the Co group, they had higher sugar-sweetened-water (25% sucrose) consumption, justifying the weight gain and higher final weight of these animals. This dietary habit in association with physical inactivity generates a positive energy balance, which results in hypertrophy and hyperplasia of adipocytes. This change in adipose tissue causes accumulation of fatty acids as well as changes in the production of inflammatory cytokines (Garcia et al., 2019; White and Ravussin, 2019). When hepatocytes reach their lipid-storing capacity, lipotoxicity and hepatocellular stress lead to apoptosis, and then, chemokines are released to activate resident macrophages, kupffer cells, and circulating monocyte-derived macrophages (Li et al., 2020). Macrophages are essential factors in the progression of MASLD. Their activation and proliferation promote several stimuli for inflammatory perpetuation with consequent progression to MASH (Ju and Tacke, 2016). Oxidative stress is also a critical alteration that favors the development of MASLD since the hepatic accumulation of lipids favors the overproduction of ROS, which at high concentrations leads to a decrease in the activity of antioxidant systems, causing several structural and oxidative changes in the metabolism of cellular macromolecules (Bayarsaikhan et al., 2022; Gabbia et al., 2021; Hong et al., 2021). Hepatic storage of triglycerides and cholesterol is an initial mechanism to prevent toxicity; however, excess steatosis can induce lipotoxicity and glucotoxicity in the liver, producing ROS in various organelles (Sabir et al., 2022; Yang and Jo, 2018). Once established the pathophysiology of MASLD, the present study aimed to verify the effects of CAR treatment on markers for the development of this condition.

The treatment with CAR did not influence food, water and caloric intake, weight gain, and final weight of the animals in the treated MASLD induction group (MASLD+CAR), corroborating data found in the literature, where CAR supplementation did not influence food and water intake, as well as animal weight parameters (Everaert et al., 2021; Giriş et al., 2014). Treatment with CAR did not influence levels of adiposity index, hypertension, and total cholesterol in the MASLD group. Instead, it was observed attenuated plasma triglyceride levels, as well as improved HDL cholesterol concentrations. Al-Sawalha et al. (2019) conducted a study in rats with metabolic syndrome and CAR supplementation for 16 weeks, where changes similar to those found in our work on the lipid profile were also observed. The positive effects of CAR on lipid parameters were also observed in humans with type 2 diabetes (Houjehani et al., 2018) who received CAR for 12 weeks, demonstrating that CAR can act as an anti-dyslipidemic agent and may have protective effects in models of obesity-related complications.

The mechanism by which CAR intervenes on lipid profile is not well understood. In the present work, along with a better plasma lipid profile and lower deposition of lipids in liver tissue it was observed important modulatory activity on PPAR- α gene expression. PPAR- α is a key transcription factor that regulates transcription of genes involved in fatty acid β -oxidation and thermogenesis (Todisco et al., 2022). Moreover, in response of PPAR- α activation or not, it was observed a modulatory effect on mRNA expression of VLCAD, an enzyme responsible for the catalysis of the first reaction of the mitochondrial very long chain fatty acids β -oxidation (Koo, 2013). The oxidation of fatty acids, mainly regulated by PPAR- α , reduces the levels of fatty acids available in the liver, using them as an energy source. However, in the face of mitochondrial dysfunction, the β -oxidation process is impaired, thus generating fat accumulation, as well as the production of ROS (Arroyave-ospina et al., 2021; Chen et al., 2020; Mong et al., 2011). In the present study, the MASLD group showed lower PPAR- α gene expression than the Co group. CAR supplementation in the MASLD+CAR increased mRNA levels compared to the MASLD group, pointing to a possible role on this component. Additionally, mRNA levels of VLCAD showed similar behavior, demonstrating high response in Co+CAR and a restoring effect in the MASLD+CAR, even still not statistically significant. There are no data in the literature reporting the role of CAR on PPAR- α , but there are studies where β -alanine, a precursor of CAR, was able to stimulate the expression of PPAR- β/δ , a key regulator of muscle lipid balance, suggesting that histidyl dipeptides can be natural ligands of PPARs (Schnuck et al., 2016). In line with this, these findings suggest a possible role of CAR on dysfunctions of lipid metabolism, in view of the reduction of

plasma and hepatic triglycerides, hepatic cholesterol, steatosis and oxidative stress, as well as an increase in PPAR- α gene expression in MASLD.

Although no significant difference was observed between the MASLD group and the Co group for CD68 expression, CAR significantly attenuated their activation compared to the MASLD group. The reduction of macrophage infiltration in MASLD+CAR is in line with the reduction of collagen deposition marked by Sirius red staining and fibrosis score in the same group, suggesting a possible role of CAR in the attenuation of MASLD progression to MASH. Similar data were found in the literature, where CAR attenuated inflammatory infiltrates (monocytes and macrophages) (Chen et al., 2017). Another study reported a significant decrease in CD68 macrophages in mice with atherosclerotic lesions treated with CAR (Barski et al., 2013). CAR's action on macrophages may be related to its potential to reduce oxidative stress, which is directly related to inflammation. However, CAR may also be acting on MIF, generating a decrease in the activity of CD68 macrophages in the obesity model. Aiello et al., 2022 observed that CAR increased MIF in a 3D skin model, associating this effect with the antioxidant action of CAR and its power to reduce oxidative stress and its positive effect on aging.

In line with this hypothesis, the present work showed that the MASLD group had higher hepatic levels of MDA, carbonyl proteins (CBO), and AOPP in comparison to the control group, but, the treatment with CAR (MASLD+CAR group) attenuated the hepatic concentrations of MDA, CBO, and AOPP in comparison to the MASLD group. It is also possible to notice lower concentrations of CBO and AOPP in the Co+CAR group compared to the Co group, which may be related to the health and quality of the animals used in the study. These data corroborate those described in the literature, where CAR reduced the plasmatic and hepatic levels of AOPP, as well as the hepatic levels of MDA and CBO in rats with diabetes induced by a hypercaloric diet and streptozotocin, possibly due to its protective effect against stress oxidative, detoxifying reactive components and producing compounds less harmful to the oxidant system, as is the case of carnosinyl proteins, which inhibit the oxidative conversion of glycated proteins into advanced glycation end-products (AGEs) (Fatih Aydın et al., 2017; Hipkiss et al., 2001). Yılmaz et al., 2017 reported that CAR decreased plasma and hepatic levels of CBO and AOPP in rats with oxidative stress induced by methylglyoxal, associating this activity with its potential antioxidant effect, as an ion chelator and ROS scavenger, preventing modifications in lipids, proteins and nucleic acids caused by metal ions to generate free radicals or by ROS themselves. Studies have described that the imidazole ring present in the structure of CAR has an essential role in inhibiting the formation of reactive

species, in addition to the anti-glycation potential of CAR, preventing the formation of abnormal proteins that can generate AGEs (Boldyrev et al., 2013; Chmielewska et al., 2020).

5. Conclusion

The administration of CAR promoted changes in markers for the MASLD pathophysiology. The supplementation of CAR in MASLD was involved with the attenuation of plasma and hepatic lipid deposition, improvement of plasma HDL cholesterol and attenuation of oxidative damage. It was also observed modulation of the PPAR- α mRNA expressions, a key component in the control of lipid metabolism and oxidative functions in liver tissue. These findings suggest that the PPAR- α transcription factor can be a potential target for the CAR effects in MASLD. More studies are still needed to better elucidate and determine the exact way by means CAR induces PPAR- α modulation. For instance, we conclude that supplementation with CAR is effective for treating lipid metabolism dysfunctions and oxidative stress in MASLD.

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CRedit authorship contribution statement

Núbia Alves Grandini: Conceptualization, Data curation, Methodology, Project administration, Writing – original draft. **Mariane Róvero Costa:** Conceptualization, Data curation, Methodology. **Cristina Schmitt Gregolin:** Methodology. **Juliana Silva Siqueira:** Data curation, Methodology. **Taynara Aparecida Vieira:** Methodology. **Artur Junio Togneri Ferron:** Methodology. **Fabiane Valentini Francisqueti-Ferron:** Data curation, Methodology. **Guilherme Ribeiro Romualdo:** Data curation, Methodology. **Ana Lúcia dos Anjos Ferreira:** Methodology. **Giancarlo Aldini:** Conceptualization. **Camila Renata Corrêa:** Data curation, Supervision, Project administration. **Fernando Moreto:** Conceptualization, Data curation, Supervision, Project administration, Writing – original draft.

Conflict of interest: none.

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Highlights

- Consumption of high sugar-fat diets is related with MASLD,
- Carnosine improved lipid metabolism dysfunctions and oxidative stress in MASLD,
- Carnosine promoted modulation of PPAR- α mRNA expression,
- Carnosine can be effective for treating MASLD.

Table 1. Diet composition and nutritional values.		
Components	Normocaloric	Hypercaloric
Soybean meal (g/kg)	335	340
Sorghum (g/kg)	278	80
Soy hulls (g/kg)	188,5	116,7
Dextrin (g/kg)	146,5	20
Maize starch (g/kg)	278	80
Sucrose (g/kg)	-	80
Fructose (g/kg)	-	180
Soybean oil (g/kg)	23	-
Lard (g/kg)	-	154,3
Minerals (g/kg)	25	25
Salt (g/kg)	4	4

Table 2. Nutritional characterization and obesity.							
	Groups				Effects		
	Co	Co+CAR	MAFLD	MAFLD+CAR	Diet	CAR	Interactions
Food intake (g/day)	37.0 ± 2.31	36.3 ± 2.07	25.6 ± 0.904*	25.3 ± 1.72†	<0.001	0.528	0.858
Water intake (ml/day)	53.2 ± 3.79	52.0 ± 7.17	68.0 ± 4.02*	64.9 ± 3.48†	<0.001	0.320	0.564
Caloric intake (kcal/day)	133 ± 8.28	130 ± 7.42	166 ± 7.84*	168 ± 5.44†	<0.001	0.849	0.584
Initial body weight (g)	324 ± 25.0	324 ± 26.8	320 ± 16.4	314 ± 27.1	0.450	0.775	0.753
Final body weight (g)	502 ± 57.2	477 ± 25.8	569 ± 38.0*	530 ± 68.5	0.008	0.133	0.743
Weight gain (g)	145 ± 48.9	137 ± 23.2	250 ± 34.6*	198 ± 48.6†	<0.001	0.085	0.201
Adiposity index (%)	4.15 ± 1.60	3.12 ± 1.11	8.51 ± 1.03*	7.68 ± 2.56†	<0.001	0.19	0.883

Data are presented as mean ± standard deviation. Effects are the comparison by two-way ANOVA with Tukey's post-hoc test, with p<0.05 as statistically significance. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine. * vs Co; † vs Co+CAR.

Table 3. Biochemical and hormonal markers.							
	Groups				Effects		
	Co	Co+CAR	MASLD	MASLD+CAR	Diet	CAR	Interactions
AST (U/l)	155 ± 34.3	125 ± 18.6	114 ± 21.0	130 ± 32.4	0.128	0.532	0.058
ALT (U/l)	104 ± 14.4	69.5 ± 16.1*	83.0 ± 26.5	87.5 ± 21.5	0.858	0.084	0.028
Uric acid (mg/dl)	0.967 ± 0.175	0.800 ± 0.420	1.27 ± 0.761	1.67 ± 1.69	0.150	0.768	0.476
Urea (mg/dl)	58.8 ± 8.66	61.0 ± 9.63	35.0 ± 15.5*	44.2 ± 4.58†	<0.001	0.195	0.417
Creatinine (mg/dl)	4.48 ± 3.08	4.59 ± 5.01	4.48 ± 1.68	4.36 ± 0.985	0.931	0.998	0.930
Blood Glucose (mg/dl)	78.0 ± 8.29	86.3 ± 11.4	88.8 ± 6.74*	97.3 ± 4.50†	0.004	0.020	0.980
Total Cholesterol (mg/dl)	62.3 ± 7.12	46.5 ± 10.1*	52.7 ± 9.63	62.0 ± 8.32†	0.430	0.380	0.002
Triglycerides (mg/dl)	70.4 ± 16.9	41.0 ± 8.32*	205 ± 34.6*	112 ± 27.4#†	<0.001	<0.001	0.004
HDL (mg/dl)	14.5 ± 3.02	14.0 ± 1.79	11.4 ± 2.33	15.5 ± 3.27#	0.471	0.114	0.047
n-HDL (mg/dl)	47.8 ± 4.67	40.5 ± 10.9	40.2 ± 6.59	46.5 ± 5.79	0.785	0.870	0.035
Insulin (ng/ml)	0.224 ± 0.031	0.175 ± 0.039	0.306 ± 0.039*	0.293 ± 0.073†	<0.001	0.132	0.379
HOMA - IR	1.24 ± 0.199	1.07 ± 0.227	1.93 ± 0.304*	2.03 ± 0.535†	<0.001	0.800	0.342

Data are presented as mean ± standard deviation. Effects are the comparison by two-way ANOVA with Tukey's post-hoc test, with p<0.05 as statistically significance. Co: control group; Co+CAR: control group treated with carnosine; MASLD: MASLD induction group; MASLD+CAR: MASLD induction group treated with carnosine; AST: aspartate aminotransferase; ALT: alanine aminotransferase; HDL: high density lipoprotein; n-HDL: non-high-density lipoprotein; HOMA-IR: homeostasis model assessment – insulin resistance. * vs Co; # vs MASLD; † vs Co+CAR.

