

RESEARCH PAPER

Combining citrulline with atorvastatin preserves glucose homeostasis in a murine model of diet-induced obesity

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BACKGROUND AND PURPOSE

NO is a crucial regulator of energy and lipid metabolism, whose homeostasis is compromised during obesity. Combination of citrulline and atorvastatin potentiated NO production *in vitro*. Here we have assessed the effects of this combination in mice with diet-induced obesity (DIO).

EXPERIMENTAL APPROACH

C57BL/6J male mice were given a standard diet (control) or a high fat–high sucrose diet (DIO) for 8 weeks. DIO mice were then treated with DIO alone, DIO with citrulline, DIO with atorvastatin or DIO with citrulline and atorvastatin (DIOcit–stat) for 3 weeks. Thereafter, body composition, glucose tolerance, insulin sensitivity and liver fat metabolism were measured.

KEY RESULTS

DIOcit–stat mice showed lower body weight, fat mass and epididymal fat depots compared with other DIO groups. Unlike other DIO groups, glucose tolerance and insulin sensitivity of DIOcit–stat, along with blood glucose and insulin concentrations in response to feeding, were restored to control values. Refeeding-induced changes in liver lipogenic activity were also reduced in DIOcit–stat mice compared with those of DIO animals. This was associated with decreased gene expression of the transcription factor SREBP-1, liver X receptor α , ChREBP and of target lipogenic enzymes in the liver of DIOcit–stat mice compared with those of other DIO groups.

CONCLUSIONS AND IMPLICATIONS

The citrulline–atorvastatin combination prevented fat mass accumulation and maintained glucose homeostasis in DIO mice. Furthermore, it potentiated inhibition of hepatic *de novo* lipogenesis activity. This combination has potential for preservation of glucose homeostasis in patients receiving statin therapy.

Abbreviations

ACC1 and *Acc1*, acetyl-CoA carboxylase 1; AMPK α , AMP-activated protein kinase α ; ChREBP and *Chrebp*, carbohydrate-responsive element-binding protein; FAS and *Fasn*, fatty acid synthase; GPAM and *Gpam*, glycerol-3-phosphate acyltransferase; LXRA and *Lxr α* , liver X receptor α ; RQ, respiratory quotient; SCD1 and *Scd1*, stearoyl-Coenzyme A desaturase 1; SREBP-1 and *Sreb1*, sterol regulatory element-binding transcription factor 1; VCO₂, carbon dioxide production; VO₂, oxygen consumption

Tables of Links

TARGETS	LIGANDS
<p>Enzymes^a Akt AMPK α, AMP-activated protein kinase α</p> <p>Nuclear hormone receptors^b LXRα, liver X receptor α (NR1H3) PPARα (NR1C1) PPARγ (NR1C3)</p>	<p>Arginine Atorvastatin Citrulline Glucose Insulin L-NAME</p>

These Tables list key protein targets and ligands in this article which are hyperlinked to corresponding entries in <http://www.guidetopharmacology.org>, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Pawson *et al.*, 2014) and are permanently archived in the Concise Guide to PHARMACOLOGY 2013/14 (^aAlexander *et al.*, 2013a, b).

Introduction

Obesity is a major risk factor for metabolic diseases including type 2 diabetes and non-alcoholic steatohepatitis. It has become a global public health priority because its prevalence has begun to rise worldwide. Today, between 15 and 30% of the population is obese, and over 45% of adults are overweight in industrialized countries (World Health Organization, 2014).

NO is a critical regulator of energy and lipid metabolism body composition and insulin sensitivity (Sansbury and Hill, 2014). NO synthesis plays a major role in the peripheral effects of insulin as it directly promotes insulin transport by vascular endothelial cells and insulin delivery to peripheral tissues (Wang *et al.*, 2013). However, NO homeostasis is compromised during obesity. Its bioavailability is decreased in obese and diabetic patients (Higashi *et al.*, 2001) and animal models (Kim *et al.*, 2008). Furthermore, mouse models of partial or total endothelial NO synthase (eNOS) knockout exhibit insulin resistance and glucose intolerance but also visceral obesity and liver steatosis (Duplain *et al.*, 2001; Cook *et al.*, 2004; Schild *et al.*, 2008). At the tissue level, dysregulation of NO production by liver endothelial cells exacerbates hepatic insulin resistance and several associated metabolic disorders, including liver steatosis (Tateya *et al.*, 2011).

To restore NO synthesis, clinical data have shown that nutritional supplementation with arginine is effective in the short term (Bode-Böger *et al.*, 2003) but presents adverse effects in the longer term (Schulman *et al.*, 2006). Furthermore, it is now known that citrulline supplementation is more efficient and more reliable than arginine. Citrulline is characterized by improved gastrointestinal tolerance (Grimble, 2007) and greater bioavailability (Moinard *et al.*, 2008) than arginine. It induces a greater increase in the plasma concentration of arginine than arginine itself (Schwedhelm *et al.*; 2008) and it is a better precursor of NO than arginine (Tangphao *et al.*, 1999). Finally, it is recognized for being safe. Several mechanisms may explain the improvement of NO synthesis with citrulline. First, citrulline inhibits the enzyme arginase, which metabolizes arginine to ornithine and urea and which is induced in diabetes (Kashyap *et al.*, 2008). Second, citrulline increases the arginine/asymmetric dimethylarginine

ratio, thus diminishing the inhibitory pressure of arginine/asymmetric dimethylarginine on eNOS (Schwedhelm *et al.*, 2008). Third, arginine is subject to a splanchnic first pass process whereas citrulline is neither metabolized in the intestine nor taken up by the liver (Curis *et al.*, 2005).

Citrulline is a non-protein amino acid with antioxidant properties and is particularly present in watermelons. It is also synthesized in enterocytes. Several studies describe a modification of its circulating concentration in obesity and diabetes (Sailer *et al.*, 2013; Zhou *et al.*, 2013), but little information is currently available to understand the importance of its endogenous synthesis in the aetiology of metabolic disorders associated with obesity. Although citrulline is known to exert regulatory effects on lipid, energy and protein metabolisms (Faure *et al.*, 2012; Faure *et al.*, 2013; Joffin *et al.*, 2014), few studies have examined the potential effects of citrulline in preventing insulin resistance and diabetes. In Zucker diabetic fatty rats, one study showed a 22% reduction in blood glucose after receiving watermelon juice (Wu *et al.*, 2007). In humans, one 7-day-long study showed no effect of citrulline supplementation on fasting insulin in healthy adults (Thibault *et al.*, 2011). Interestingly, combining citrulline with atorvastatin was recently shown to potentiate NO production in endothelial cells (Berthe *et al.*, 2011). This was mediated through an increased bioavailability of arginine, due to citrulline, and to an enhanced activity of the NO-generating enzyme eNOS because the statins, apart from their ability to decrease cholesterol synthesis, also increase eNOS expression and activity (Berthe *et al.*, 2011).

Our aim was thus to investigate *in vivo* the benefits of combining citrulline to atorvastatin in a mouse model of diet-induced obesity (DIO) known to develop glucose intolerance and insulin resistance within 8 to 10 weeks (Bonnard *et al.*, 2008). The primary objective of our study was to explore the individual and combined effects of citrulline and atorvastatin on body weight and composition and on glucose tolerance and insulin resistance. Our secondary objective was to explore the individual and combined effects of citrulline and atorvastatin on liver fat metabolism, which is closely interrelated with glucose homeostasis (Bechmann *et al.* 2012).

Methods

Animals and experimental protocols

All animal care and experimental procedures complied with the guidelines formulated by the European Community for the use of experimental animals (L358-86/609/EEC) and were approved by the local ethical committee (agreement no. A633-239). INRA animal facilities were approved (C634514) by the French veterinary department. All studies involving animals are reported in accordance with the ARRIVE guidelines for reporting experiments involving animals (Kilkenny *et al.*, 2010; McGrath *et al.*, 2010). A total of 160 animals were used in the experiments described here.

Male C57BL/6J mice (5 weeks old) (Janvier Labs, France) were housed in individual cages, subjected to a 12 h light/dark cycle at a temperature of $22 \pm 2^\circ\text{C}$. All animals had *ad libitum* access to food and water. After 1 week of acclimatization under the control diet (20% protein, 70% carbohydrates and 10% fat), 40 mice were maintained on the control diet for a further 12 weeks (control group). The remaining 120 mice (DIO groups) were fed a high fat–high sucrose diet (20% protein, 18% carbohydrates, 17% sucrose and 45% fat). During the last 3 weeks of the dietary intervention, DIO mice were randomized into four groups. Each group received either DIO alone ($n = 40$), or DIO enriched with citrulline (2.5 g kg^{-1} ; DIOcit, $n = 20$), atorvastatin (10 mg kg^{-1} ; DIOstat, $n = 20$) or the combination of citrulline (2.5 g kg^{-1}) and atorvastatin (10 mg kg^{-1} ; DIOcit–stat, $n = 40$). This dose of 10 mg atorvastatin has been repeatedly shown to be effective on lipid metabolism in mice, especially in mouse models of inflammation (Chen *et al.*, 2014a, b). In addition, our own work has demonstrated that 1 to $5 \text{ g kg}^{-1} \text{ day}^{-1}$ citrulline is required in order to induce metabolic effects in rodents (Moinard *et al.*, 2009; Faure *et al.*, 2012, 2013).

Body weight and food intake were measured every 2 weeks. At the end of the experiment, most animals were examined after an overnight fast. Twenty animals from control, DIO and DIOcit–stat groups were used to explore the effect of refeeding. Briefly after an overnight fast, mice were refed (or not) with their diet during 2 h before killing.

- **Measurement of food intake:** Mice were given *ad libitum* access to a known amount of pellets for 1 week. At the end of the test, leftover food was weighed, and mean daily food intake was calculated in g per day.
- **Indirect calorimetry method:** Oxygen consumption (VO_2) and carbon dioxide production (VCO_2) were measured using a four-chamber TSE systems PhenoMaster/LabMaster (Bad Homburg, Germany). Temperature was maintained at 28°C to reach the animals' thermoneutrality and the light was on from 08:00 to 20:00 h. System settings included a flow-rate of 0.6 L min^{-1} , a sample purge of 5 min and a measurement period of 5 min every 25 min. Sixteen hours prior to data collection, mice were placed in separate calorimetry chambers (each with a volume of 2.5 L), with free access to food and water. Then VO_2 and VCO_2 were monitored during 24 h while food was available *ad libitum*. Energy expenditure was calculated using Brouwer's equation (1965). The respiratory quotient (RQ) was calculated as the ratio of VCO_2 to VO_2 . Daily energy expenditure and RQ were computed over the 24 h period.

- **Body composition determination:** Fat and lean masses (g) were assessed *in vivo* in fasted mice ($n = 5$ per group) using the EchoMRI-100 instrument (Echo Medical Systems LLC, Houston, USA).
- **Glucose homeostasis:** Each group was divided into two subgroups for exploring either insulin sensitivity or glucose tolerance after 4 h of fasting. For insulin sensitivity assessment, basal blood glucose was measured in a drop of blood from the tail using a glucometer (SureStep, Life Line Screening, Independence, USA) and an insulin dose of 0.75 mUI g^{-1} was then injected *i.p.*. Blood glucose was measured after 15, 30, 45, 60 and 120 min. Insulin sensitivity was evaluated from the increase in blood glucose compared with basal glucose, which is the blood glucose area over the curve. To assess glucose tolerance, after basal blood glucose measurement, a dose of glucose (2 mg g^{-1}) was injected *i.p.*, and blood glucose was measured after 15, 30, 60 and 120 min. Glucose tolerance was evaluated from blood glucose area under the curve.
- **Plasma and tissue collection:** Animals were anaesthetized with pentobarbital ($90 \mu\text{g g}^{-1}$, *i.p.*). Blood was collected (EDTA as anticoagulant) and centrifuged, and plasma was aliquoted and frozen at -80°C . Liver, epididymal adipose tissue and leg muscles (quadriceps + gastrocnemius + tibialis) were dissected and weighed. The liver was frozen in liquid nitrogen and kept at -80°C until subsequent analysis.
- **Plasma metabolites and hormones:** Blood was collected in EDTA-coated tubes. Insulin was assessed using an enzyme-linked immunosorbent assay kit (Alpco Diagnostics, Salem, USA). Fasting glucose, triglycerides and free fatty acids were assessed using an automated system (Konelab 20, Kone, Helsinki, Finland). Reagents were obtained from Randox (Crumlin, UK) and ThermoElectron Corporation (Waltham, USA).

To assess plasma concentrations of amino acids, blood was collected into heparinized tubes. Plasma was then deproteinized with sulfosalicylic acid (10%), frozen and stored at -80°C . Plasma amino acids were separated and quantified by ion-exchange chromatography with spectrophotometric detection after ninhydrin derivatization using an AminoTac JLC-500/VAA analyser (Jeol, Croissy-sur-Seine, France).

Cell culture

Human HuH7 hepatocarcinoma cells (from Creative Bioarray, Shirley, NY, USA) were grown at 37°C in a 5% CO_2 atmosphere in a medium composed of DMEM containing 1 g L^{-1} glucose and supplemented with 10% fetal calf serum, 1% glutamine and 1% antibiotics (5 U mL^{-1} penicillin and $50 \mu\text{g mL}^{-1}$ streptomycin). At ~70% confluence, cells were incubated for 24 h with a medium enriched with citrulline (5 mM) and/or atorvastatin (0.02 or 0.05 mM) and/or the NOS inhibitor, L-NAME (0.5 mM); DMSO concentration was the same under all conditions. Thereafter, cells were lysed at 4°C in a lysis buffer (50 mM HEPES pH 7.4, 150 mM NaCl, 10 mM EDTA, 10 mM NaPPI, 25 mM β -glycerophosphate, 100 mM NaF, 2 mM Na_3VO_4 , 10% glycerol, 1% Triton X-100 and 0.5% of a protease-inhibitor cocktail) for further Western blotting

analyses. For the exploration of insulin signalling, DMEM was depleted of fetal calf serum for 3 h; cells were then incubated with 10^{-7} M insulin for 15 min and lysed as described above.

Cell and liver metabolic adaptations to diets

- **Quantification of lipid content:** Lipids from livers were extracted according to Folch and coworkers (Folch *et al.*, 1957). Triglyceride content was assessed colorimetrically using the Triglyceride Colorimetric Assay kit (Cayman, Ann Arbor, USA).
- **Real-time quantitative RT-PCR analysis:** Total RNA was extracted with Tri-Reagent® Solution (Sigma). Target RNA levels were measured by reverse transcriptase followed by real-time PCR using a Rotor-Gene Q system (Qiagen, Courtaboeuf, France). A standard curve was generated from a mixture of all native cDNA and serial dilutions. *Hprt* was used as a reference gene and results are expressed relative to *Hprt* expression. Primer sequences are shown in Table 1.
- **Immunoblotting:** Livers were homogenized in the ice-cold lysis buffer described above. After centrifugation, protein content of supernatants was determined using MicroBCA™ Protein assay kit (Thermo Scientific, Rockford, USA). Lysate proteins were solubilized in 4X Laemmli's buffer, boiled at 95°C for 5 min, resolved by SDS-PAGE and transferred to PVDF membranes (1 h, 100 V) (Merck Millipore, Darmstadt, Germany). Membranes were blocked for 1 h in 5% BSA-TBS 50 mM Tris-Cl, 150 mM NaCl, pH 7.6/Tween 0.01% and then incubated with the specific primary antibodies. Signal was detected with a horseradish peroxidase-conjugated secondary antibody and revealed with an enhanced chemiluminescence system (Pierce, Thermo Scientific). Results are expressed as a ratio of total protein

or α -tubulin expression. Adjustment was made to internal control on each gel and to controls in HuH7 cells and DIOcit-stat in mice.

Data analyses

All data are presented as means \pm SEM. A one-way ANOVA was performed to test the effect of the experimental conditions using the Statview software (SAS Institute, USA). When a significant effect was detected at $p < 0.05$, Fisher's test was used to analyse pairwise differences.

Materials

DMEM was from Sigma (St Louis, USA). Fetal calf serum and heat-inactivated horse serum were purchased from PAN-Biotech GmbH (Aidenbach, Germany). Antibiotics were supplied by Gibco-Invitrogen (Carlsbad, USA). Atorvastatin was a generic product from Pfizer, NY, USA. Insulin Actrapid® was from Novo Nordisk (Bagsvaerd, Denmark). Primary antibodies were against phospho(Ser⁴⁷³)-Akt (Cell Signaling, Danvers, USA, #9271), total-Akt (Cell Signaling, #9272), phospho(Thr¹⁷²)-AMP-activated protein kinase α (AMPK α) (Cell Signaling, #2531), total-AMPK α (Cell Signaling, #2532), sterol regulatory element-binding transcription factor 1 (SREBP-1) (Santa Cruz Biotechnology, Dallas, USA, #sc-8984) and α -tubulin (Sigma, #T5168). Appropriate secondary antibodies were obtained from Cell Signaling and Dako (Glostrup, Denmark). Other chemicals used were of the highest grade commercially available. Citrulline was a gift from Citrage® (Créteil, France).

Results

Otherwise stated, results presented below apply to fasted conditions.

Table 1

Quantitative reverse transcription PCR primers

Gene name	Forward primer	Reverse primer
<i>Acc1</i>	ACCTGGTGGAGTGGCTGGAG	ATGGCGACTTCTGGGTTGGC
<i>Chrebp</i>	GGAGAGCCTGGTACATTCAG	GCTTCCAGTAATTACCCTCCAG
eNOS	GAGAGCGAGCTGGTGTTTG	CTGTGATGGCTGAACGAAGA
<i>Fasn</i>	GGCTGCTGTTGGAAGTCAG	TGCCCTGAACCACTCACAC
<i>Gpam</i>	AGTTCGCGAGTCTGAGTACC	GTCTCTTTGAAAACCCCGATG
<i>Hprt</i>	TTGCTGACCTGCTGGATTAC	AGTTGAGAGATCATCTCCAC
<i>Lxra</i>	GGAGTGTGCGACTTCGCAAAT	GCAGGACTTGAGGAGGTGAG
<i>Pepck</i>	CCCGAAGGCAAGAAGAATA	CGTTTTCTGGGTTGATAGCC
<i>Pgc-1α</i>	GAAGTGGTGTAGCGACCAATC	AATGAGGGCAATCCGTCTTCA
PPAR α	TGTGGCTGCTATAATTTGCTGTGG	CCCTCTGCAACTTCTCAATGTAG
PPAR γ	CAAGAATACCAAAGTGCATCAA	GAGCTGGGCTTTTCAGAATAATAAG
<i>Scd1</i>	CCTCCTGCAAGCTCTACACC	CAGCCGTGCCTTGTAAAGTTC
<i>Srebfl</i>	CACCACCGTACAGCTCAG	GTGGCTGCTGAGTGTTCCT

Hprt, hypoxanthine guanine phosphoribosyl transferase; *Pepck*; phosphoenolpyruvate carboxykinase; *Pgc-1 α* , peroxisome proliferator-activated receptor gamma coactivator 1- α .

Citrulline and atorvastatin combination alters plasma amino acid concentrations

Citrulline supplementation was accompanied by a specific increase in plasma citrulline, arginine, ornithine and proline concentrations. However, these modifications disappeared when treatment with atorvastatin was combined with citrulline (Table 2). In addition, the plasma concentrations of branched-chain amino acids (valine, leucine and isoleucine) and essential amino acids (phenylalanine and threonine) were decreased in the four DIO groups compared with controls (Table 2).

Citrulline and atorvastatin combination prevents weight gain but did not alter daily energy balance

• **BW and body composition:** Before the pharmaco-nutritional intervention started, i.e. after 8 weeks of control of DIO diet, the DIO mice gained 8% more body weight than controls ($p < 0.01$, Figure 1A). After 4 weeks of treatment, the weight of DIOcit-stat mice was significantly lower than those of the other DIO groups whereas it was not significantly different from the controls ($p < 0.05$, Figure 1A). Weights of DIO, DIOcit and DIOstat animals was significantly higher than

controls ($p < 0.001$, Figure 1A). EchoMRI showed that differences in weight were linked to differences in fat mass, lean mass being similar between the five groups (Table 3). Fat mass of DIOcit-stat mice was 22% lower than other DIO groups ($p < 0.05$) but remained 46% higher than controls ($p < 0.05$, Table 3). These results were confirmed by weighing epididymal fat depots at the end of the experiment (Figure 1B, Table 3).

• **Daily energy balance** was similar between DIOcit-stat and other DIO groups. Daily food intake averaged 3 g per day in the five groups, which represents a mean intake of 56 and 78 kJ day⁻¹ for the control and DIO groups respectively. Furthermore, daily energy expenditure of DIOcit-stat mice was similar to other DIO groups but was 13% lower than controls because of a reduced spontaneous physical activity ($p < 0.01$, Table 3). Finally, daily RQ of DIOcit-stat mice was similar to other DIO groups and was significantly lower than controls ($p < 0.0001$, Table 3).

Citrulline and atorvastatin combination improves glucose homeostasis in vivo and in vitro

• **Plasma metabolites and hormones:** Fasting glucose, insulin, triglycerides and free fatty acid plasma concentrations of

Table 2

Plasma amino acid concentration ($\mu\text{mol L}^{-1}$)

	Control	DIO	DIOcit-stat	DIOstat	DIOcit
Citrulline and related amino acids					
Citrulline	49 ± 2	49 ± 3	51 ± 6	46 ± 2	84 ± 19*
Arginine	89 ± 5	94 ± 4	96 ± 11	94 ± 2	134 ± 22*
Ornithine	45 ± 4	35 ± 2*	31 ± 2*	32 ± 1*	52 ± 7 [§]
Glutamine	538 ± 24	521 ± 18	481 ± 17	510 ± 22	472 ± 21
Proline	64 ± 3	62 ± 6	53 ± 3	53 ± 3	87 ± 15*
Branched-chain amino acids					
Valine	242 ± 20	180 ± 8*	145 ± 7*	165 ± 5*	195 ± 11*
Isoleucine	100 ± 6	80 ± 4*	66 ± 4*	79 ± 4*	78 ± 5*
Leucine	185 ± 14	137 ± 7*	111 ± 9*	136 ± 7*	133 ± 10*
Other essential amino acids					
Phenylalanine	78 ± 4	66 ± 3.6*	59 ± 3*	63 ± 2.2*	67 ± 3.1*
Methionine	54 ± 3	53 ± 4	43 ± 2	46 ± 2	56 ± 5
Lysine	241 ± 7	251 ± 14	214 ± 9	230 ± 8	256 ± 18
Threonine	182 ± 10	168 ± 13	144 ± 8*	156 ± 8	191 ± 15*
Histidine	61 ± 3	57 ± 3	50 ± 2	55 ± 2	60 ± 4
Other non-essential amino acids ^A					
Serine	102 ± 4	103 ± 7	92 ± 4	93 ± 3	126 ± 17*
Asparagine	38 ± 2	35 ± 2	31 ± 1*	33 ± 1.4 ^Y	39 ± 2

Results are means ± SEM ($n = 10$ per group). Dietary interventions: standard diet (control), high fat-high sucrose diet (DIO) and DIO enriched in citrulline (2.5 g kg⁻¹) (DIOcit), atorvastatin (10 mg kg⁻¹) (DIOstat) or citrulline and atorvastatin (DIOcit-stat).

^AOnly non-essential amino acids significantly altered by the dietary interventions are shown.

* $P < 0.05$, significantly different from controls;

[§] $P < 0.05$, significantly different from control and other DIO groups.

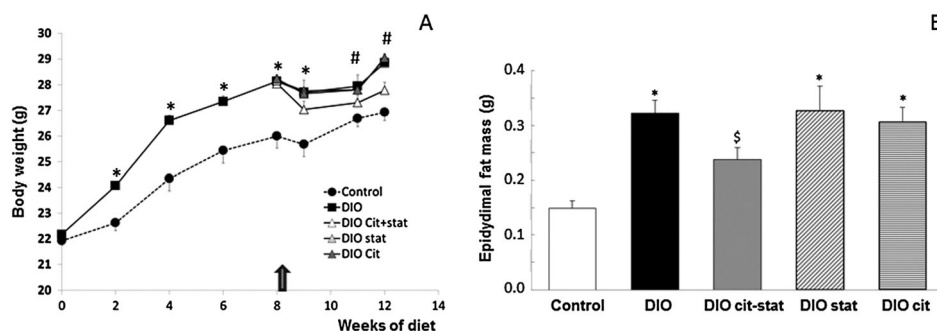


Figure 1

(A) Changes in body weight during the dietary intervention. The arrow indicates the beginning of the pharmaco-nutritional intervention with citrulline (cit) and/or atorvastatin (stat) in the DIO mice. (B) Differences in epididymal fat mass at the end of the protocol. * $P < 0.05$, significantly different from controls; # $P < 0.05$, significantly different from controls and DIOcit–stat group; [§] $P < 0.05$, significantly different from control and other DIO groups.

Table 3

Characterization of the animals according to the dietary intervention

	Control	DIO	DIOcit–stat	DIOstat	DIOcit
Body composition (otherwise stated, $n = 5$ per group)					
Lean mass (g) ^A	22.85 ± 0.85	24.07 ± 0.65	23.58 ± 0.59	22.13 ± 0.47	23.10 ± 0.73
Fat mass (g) ^A	1.17 ± 0.11	2.51 ± 0.17*	1.79 ± 0.12 [§]	2.18 ± 0.19*	2.34 ± 0.24*
Liver (mg) ^B	853 ± 39	972 ± 34	925 ± 12	1000 ± 72	1019 ± 51
Leg muscle (mg) ^B	743 ± 29	752 ± 23	762 ± 4	728 ± 17	759 ± 7
Epididymal fat (mg) ^B ($n = 20$ per group)	149 ± 14	333 ± 97*	254 ± 26 [§]	337 ± 46*	313 ± 28*
Calorimetric chamber measurements ($n = 4$ per group)					
Daily EE (kJ g ⁻¹ LM ⁻¹ day ⁻¹)	1.31 ± 0.03	1.10 ± 0.06*	1.14 ± 0.01*	1.18 ± 0.03*	1.20 ± 0.03*
Daily RQ	0.97 ± 0.01	0.82 ± 0.03*	0.86 ± 0.01*	0.82 ± 0.01*	0.85 ± 0.01*
Plasma concentrations ($n = 10$ per group)					
Triglycerides (g L ⁻¹)	0.40 ± 0.02	0.52 ± 0.04 [§]	0.42 ± 0.02	0.45 ± 0.01	0.44 ± 0.01
Free fatty acids (mM)	0.53 ± 0.02	0.71 ± 0.03*	0.57 ± 0.05	0.63 ± 0.02*	0.64 ± 0.01*
Glucose (g L ⁻¹)	1.31 ± 0.05	1.70 ± 0.10 [§]	1.28 ± 0.05	1.32 ± 0.05	1.47 ± 0.10
Insulin (ng mL ⁻¹)	0.50 ± 0.03	1.03 ± 0.12 [§]	0.63 ± 0.04	0.53 ± 0.03	0.48 ± 0.03

Results are means ± SEM. Dietary interventions: standard diet (control), high fat–high sucrose diet (DIO), and DIO enriched in citrulline (2.5 g kg⁻¹) (DIOcit), atorvastatin (10 mg kg⁻¹) (DIOstat) or citrulline and atorvastatin (DIOcit–stat). Leg muscle mass is the sum of the two *tibialis*, *soleus*, *gastrocnemius* and *quadriceps* muscle mass; Daily EE, energy expenditure measured over 24 h; Daily RQ, respiratory quotient measured over 24 h; LM, lean mass.

^AMeasured using EchoMRI;

^BMeasured at death.

* $P < 0.05$, significantly different from controls;

[§] $P < 0.05$, significantly different from control and other DIO groups.

DIOcit–stat animals were significantly lower than DIO animals ($p < 0.05$, Table 3). In addition, they were not significantly different from controls and from DIOcit and DIOstat groups (Table 3).

- *In vivo*, DIOcit–stat was the only intervention that improved glucose tolerance and insulin sensitivity in comparison with that of DIO and DIOcit ($p < 0.05$) and maintained them at control values (Figure 2). To support the improved glucose homeostasis in response to the combined

supplementation, we explored blood glucose and insulin concentrations 2 h after refeeding. Blood glucose and insulin concentrations in fed DIOcit–stat mice were significantly lower than in DIO ($p < 0.05$) and were similar to controls (Figure 3A and B). These results suggested an improved inhibition of hepatic gluconeogenesis in response to feeding. To support this hypothesis, the hepatic gene expression of phosphoenolpyruvate carboxykinase, a rate-controlling step of gluconeogenesis, was significantly

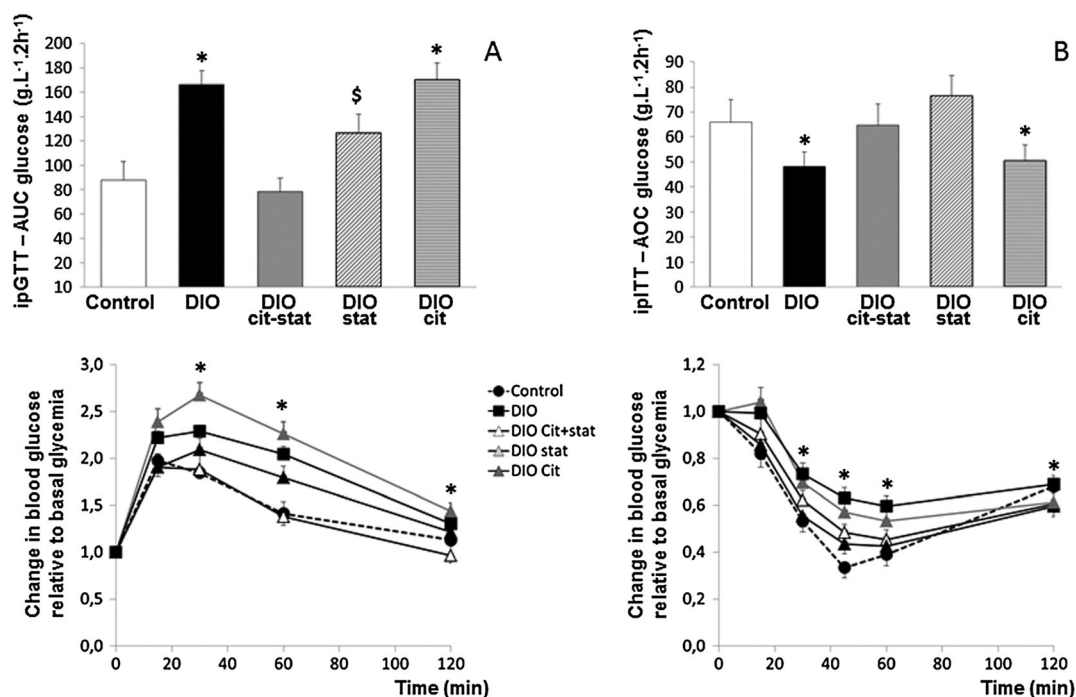


Figure 2

(A) Glucose tolerance [intraperitoneal glucose tolerance test (ipGTT)] and glycemic response to ipGTT, and (B) insulin sensitivity [intraperitoneal insulin sensitivity tolerance test (ipITT)] and glycemic responses to ipITT in diet-induced obese (DIO) mice treated or not with citrulline (cit) and/or atorvastatin (stat) compared with control lean mice. Glycemic responses are expressed relative to basal blood glucose. * $P < 0.05$, significantly different from controls; [§] $P < 0.05$, significantly different from DIOcit-stat group.

down-regulated 2 h after refeeding in DIOcit-stat and in control mice ($p < 0.05$, Figure 3C). This down-regulation was not observed in DIO animals, which exhibited no significant alteration of phosphoenolpyruvate carboxykinase gene expression after refeeding (Figure 3C).

- *In vitro*, the effects of citrulline and atorvastatin on the insulin signalling pathway were examined in HuH7 cells using the ratio tested in the preclinical study, that is, using citrulline 5 mM and atorvastatin 0.02 mM. Whereas citrulline tended to reduce the insulin-induced Akt phosphorylation compared with that of controls ($p > 0.10$), atorvastatin significantly decreased the insulin-induced Akt phosphorylation compared with that of controls ($p < 0.05$, Figure 3D). Interestingly, combining citrulline with atorvastatin significantly increased the insulin-induced Akt phosphorylation compared with that of citrulline and atorvastatin alone ($p < 0.05$), thus restoring Akt phosphorylation to control values (Figure 3D).

Citrulline and atorvastatin combination reduces the lipogenic activity in the liver through AMPK α activation and down-regulation of SREBP-1

- The triglycerides content of liver was not significantly different between groups, particularly between DIOcit-stat and DIO, although DIOcit-stat was close to control values (Figure 4A). However, in response to refeeding, the increase in liver triglyceride content was 27% lower in DIOcit-stat

compared with DIO ($p < 0.05$), but still 41% higher than in controls ($p < 0.05$, Figure 4B).

- AMPK is a cellular energy sensor that, among its many actions, integrates diverse physiological signals to restore energy balance. In particular, it suppresses SREBP-1 cleavage and nuclear translocation and represses SREBP-1 target gene expression leading to reduced lipogenesis and lipid accumulation in hepatocytes (Li *et al.*, 2011).

In vivo, hepatic AMPK α phosphorylation was significantly higher in DIOcit-stat than in DIO ($p < 0.05$, Figure 4D), DIOcit and DIOstat animals being intermediary ($p > 0.10$, Figure 4B). In addition, AMPK α phosphorylation was similar between DIOcit-stat and controls (Figure 4D).

In vitro, studies with HuH7 cells showed that citrulline, like atorvastatin (Sun *et al.*, 2006), activated AMPK α (Figure 5A) and decreased SREBP-1 protein content (Figure 5B). Yet at the ratio used in the preclinical study (citrulline 5 mM/atorvastatin 0.02 mM), citrulline did not further potentiate the atorvastatin-induced activation of AMPK α and decrease in SREBP-1 protein content (*data not shown*). Treatment with L-NAME, a NOS inhibitor, tended to reduce citrulline-induced AMPK α phosphorylation and significantly reduced atorvastatin-induced activation of AMPK α (Figure 5C).

We then analysed the gene expression of SREBP-1 and of major nuclear and transcription factors involved in the regulation of hepatic lipid metabolism. Hepatic *Sreb1* (Figure 4C) and liver X receptor α (*Lxra*) (Table 4) mRNA content was significantly lower in DIOcit-stat compared with all other

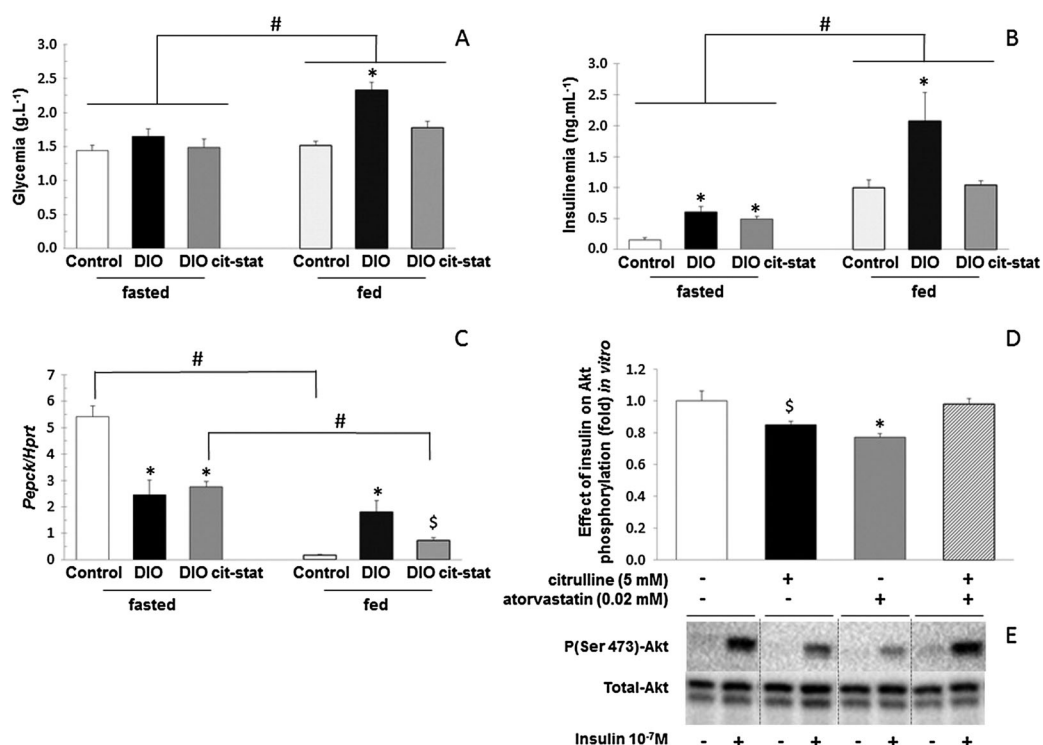


Figure 3

(A) Glycaemia and (B) insulinaemia in fasted and 2 h refed diet-induced obese (DIO) mice treated or not with citrulline (cit) and atorvastatin (stat) and in control lean mice. (C) phosphoenolpyruvate carboxykinase (PEPCK) gene expression in liver of fasted and 2 h refed DIO mice treated or not with citrulline (cit) and/or atorvastatin (stat) and in control lean mice. (D) Impact of citrulline, atorvastatin and combination of the two on insulin-induced changes in phospho(Ser⁴⁷³)-Akt/total Akt in human Huh7 hepatocarcinoma cells and (E) related Western blot images. For A–C: * $P < 0.05$, significantly different from controls; # $P < 0.05$, significantly different fasted conditions; \$ $P < 0.05$, significantly different from control and DIO groups. For D: * $P < 0.05$, significantly different from controls and citrulline–atorvastatin combination; \$ $P < 0.05$, significantly different from citrulline–atorvastatin combination. HPRT, hypoxanthine guanine phosphoribosyl transferase.

groups, including controls ($p < 0.05$). Furthermore, the mRNA for the carbohydrate-responsive element-binding protein (*Chrebp*) was ~42% lower in DIOcit–stat compared with other DIO groups ($p < 0.01$) and reached control values (Table 4). Regarding the gene expression of other regulators of hepatic lipid metabolism, *PPAR α* mRNA content was similar between DIOcit–stat, DIOstat and controls, whereas that it was almost twofold higher in DIO and DIOcit groups compared with that of controls ($p < 0.01$, Table 4). *PPAR γ* mRNA content of DIOcit–stat was 44 to 62% lower than other DIO groups ($p < 0.05$) and was similar to controls (Table 4). Finally, *PPAR γ coactivator1- α* gene expression was significantly reduced in DIOcit–stat, as in DIO and DIOstat, in comparison to controls and DIOcit (–28%, $p < 0.05$, Table 4).

In response to refeeding, *Srebf1* mRNA content was not altered in DIOcit–stat animals contrasting with DIO and control mice (Table 5). By contrast, *Lxr α* , *Chrebp*, *PPAR γ coactivator1- α* , *PPAR α* and *PPAR γ* were similarly altered in DIOcit–stat, DIO and controls (Table 5).

Fatty acid synthase (FAS), acetyl-CoA carboxylase-1 (ACC1), stearoyl-CoA desaturase-1 (SCD1) and glycerol-3-phosphate acyltransferase (GPAM) are key lipogenic enzymes. They are established targets of the LXR α /SREBP-1 pathway but also of ChREBP and PPARs (Poupeau and

Postic, 2011). As expected from alterations in *Srebf1* gene expression, *Acc1*, *Fasn* and *Scd1* mRNA content was significantly lower in DIOcit–stat compared with that of all other groups, including controls ($p < 0.05$, Table 4). By contrast, *Gpam* gene expression was similar between all conditions (Table 4). Furthermore, in response to refeeding, *Fasn* and *Acc1* mRNA contents were enhanced in refed DIOcit–stat as in controls ($p < 0.05$), whereas *Scd1* and *Gpam* gene expression were differently altered between the two groups (Table 5).

Discussion

We aimed at exploring whether the combination of nutrition and pharmacology could have a synergistic effect on major metabolic traits associated with obesity and risk of type 2 diabetes. In a murine model of DIO, the combination of citrulline with atorvastatin synergistically improved glucose homeostasis in comparison with DIO groups receiving each compound alone. Furthermore, the combination also synergistically affected fat metabolism. This was characterized by a lower fat accumulation, notably in epididymal depots, and

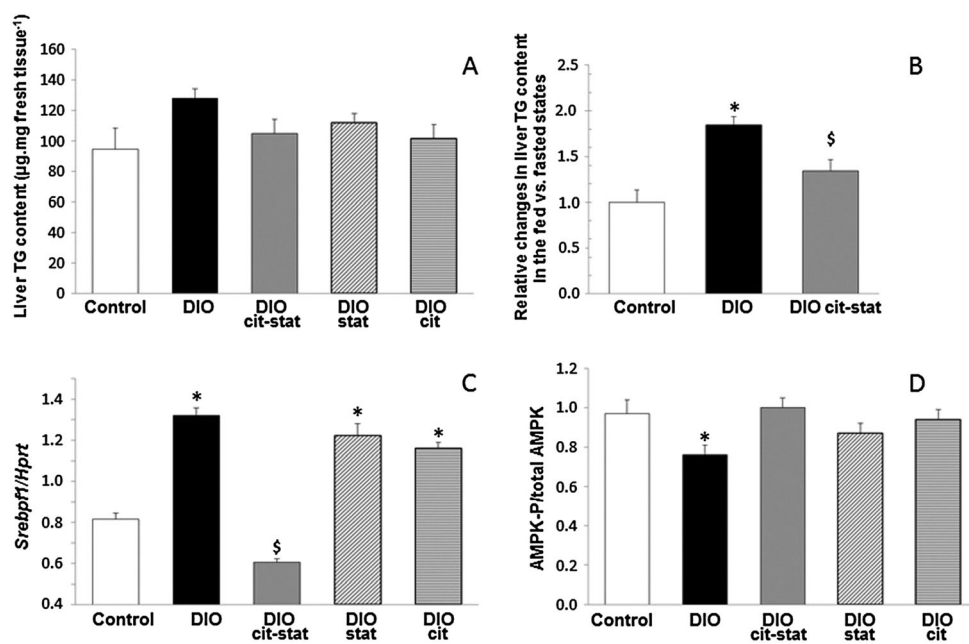


Figure 4

(A) Triglycerides (TG) content, (B) relative changes in liver TG content in 2 h re-fed mice compared with overnight fasted animals, (C) *SREBP1* mRNA content and (D) phospho(Thr172)-AMPK α /total AMPK α , in livers of DIO mice treated or not with citrulline (cit) and/or atorvastatin (stat) and compared with control lean mice. * $P < 0.05$, significantly different from controls; $^{\$}P < 0.05$, significantly different from controls and other DIO groups. *Hprt*, hypoxanthine guanine phosphoribosyl transferase.

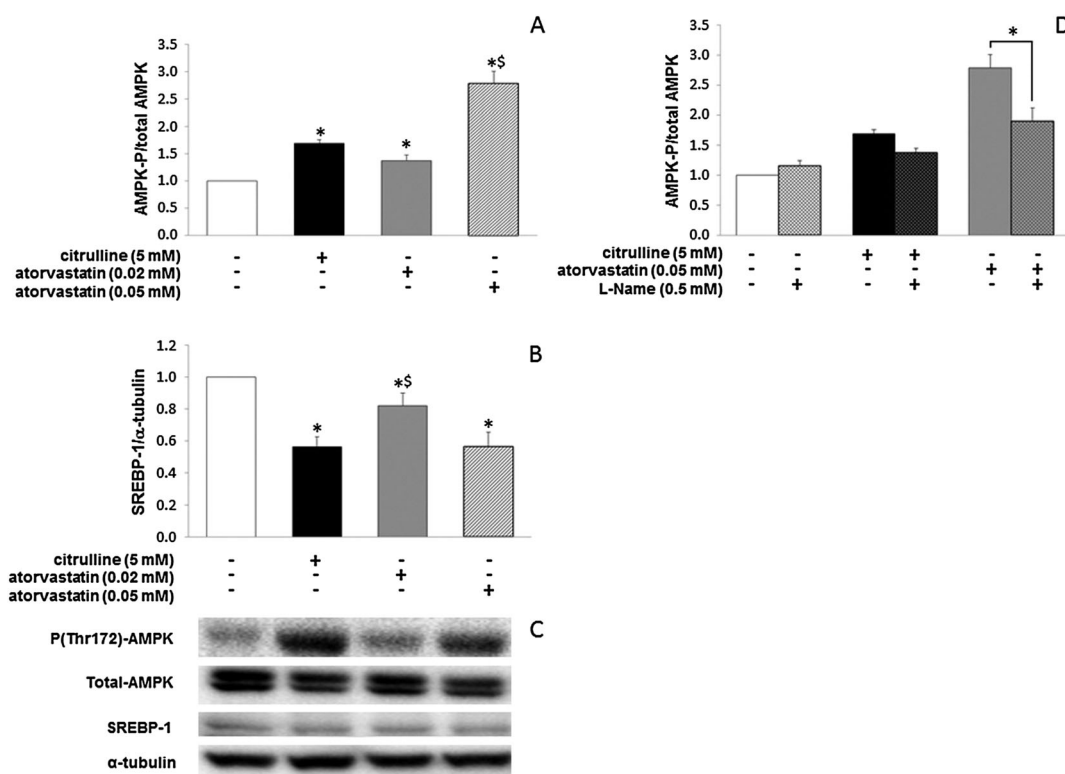
a marked inhibition of hepatic *Srebp1* gene expression and *de novo* lipogenesis activity in DIOcit-stat mice compared with other DIO groups.

Our results could be of particular value for patients treated with statins, because long-term treatment with statins has been associated with several side-effects including alteration of glucose homeostasis and increased risk of developing type 2 diabetes (Food and Drug Administration, 2012). In this respect, the relevancy of our results is strengthened by the fact that they were obtained in a model of DIO. Indeed, the risks for developing diabetes with statins are higher in individuals expressing signs of cardiometabolic disorders compared with those without such risk factors (Waters *et al.*, 2011). Because the ability of statins to decrease major cardiovascular events and mortality outweighs the risk of incident diabetes, the American College of Cardiology/American Heart Association Task Force has recommended not to change clinical practice and not to discontinue statin treatment (Stone *et al.*, 2014). Therefore, it is clearly necessary to develop strategies to minimize the negative metabolic effects of statins and to maintain glucose homeostasis, especially in pre-diabetic patients and in patients at high cardiometabolic risk.

The improved glucose tolerance we observed in response to the pharmaco-nutritional intervention is likely to involve a combination of several beneficial adaptations. *In vivo* and *in vitro* data supported a synergistic effect of the pharmaco-nutritional combination on the improvement of liver insulin sensitivity. These adaptations were further supported by the favourable impact of the product combination on fat mass, especially on the intra-abdominal fat depots, because a high level of visceral adipose tissue is associated in humans with a deterioration of glucose tolerance (Després, 1993). Another

mechanism of citrulline-atorvastatin action may also involve an improved hepatic lipid metabolism, although it is still debated whether the correlation between hepatic *de novo* lipogenesis, lipid accumulation and insulin resistance is due to a causal relationship (Farese *et al.*, 2012). Data showed in the liver that the combination of citrulline and atorvastatin specifically blunted the DIO-induced expression of SREBP-1 and of its downstream targets. In agreement with Li *et al.* (2011), our *in vivo* and *in vitro* data suggested that SREBP-1 down-regulation was mediated by the activation of AMPK α . These synergistic effects of the combination were unexpected. Indeed, the effects of statins on liver lipid metabolism has been little investigated. In rats fed with fructose (Roglans *et al.*, 2002; Rodríguez-Calvo *et al.*, 2009) or high fat (Ji *et al.*, 2011) to induce hepatic *de novo* lipogenesis, a high dose of atorvastatin (30 mg kg⁻¹), but not a low dose (5 mg kg⁻¹), significantly reduced the liver triglyceride content. Because animals from the present study received a low dose of atorvastatin, our data suggest that citrulline is required to increase the hepatic adaptations during statin therapy.

Data are also scarce concerning the effects of statins on adipose tissue mass. Aguirre *et al.* (2013) showed in obese Zucker rats that subcutaneous adipose tissue was significantly increased after treatment with atorvastatin. By contrast, citrulline supplementation was shown to reduce visceral fat depots in old rats (Moinard *et al.*, 2009). Experiments on visceral adipose tissue explants from overweight rats showed that citrulline directly induced phosphorylation of hormone-sensitive lipase and downregulated glyceroneogenesis, allowing an increased fatty acid release from the adipose tissue (Joffin *et al.*, 2014). Surprisingly, these effects were not observed in the present study because mice

**Figure 5**

Citrulline and atorvastatin-induced changes in (A) phospho(Thr¹⁷²)-AMPK α /total AMPK α and (B) SREBP-1 protein expression in human HuH7 hepatocarcinoma cells. (C) Western blot images of phospho(Thr¹⁷²)-AMPK α , total AMPK α and SREBP-1 protein expression from (A) and (B). (D) L-NAME, a NOS inhibitor, reduces citrulline and atorvastatin-induced changes in phospho(Thr¹⁷²)-AMPK α /total AMPK α . * $P < 0.05$, significantly different from controls; [§] $P < 0.05$, significantly different from control and other treated groups.

Table 4

mRNA content of major proteins involved in the regulation of liver *de novo* lipogenesis

	Control	DIO	DIOcit-stat	DIOstat	DIOcit
<i>Lxra</i>	0.95 ± 0.14	1.37 ± 0.13*	0.61 ± 0.08 [§]	1.26 ± 0.11	1.08 ± 0.12
<i>Chrebp</i>	0.98 ± 0.15	1.73 ± 0.26*	0.99 ± 0.15	1.68 ± 0.19*	1.72 ± 0.17*
PPAR α	0.92 ± 0.09	1.32 ± 0.15	0.67 ± 0.10*	0.78 ± 0.10*	1.41 ± 0.18
<i>Pgc-1α</i>	1.37 ± 0.20	1.02 ± 0.12*	0.98 ± 0.08*	0.81 ± 0.07*	1.55 ± 0.25
PPAR γ	0.73 ± 0.09	1.31 ± 0.27*	0.60 ± 0.06 [#]	1.08 ± 0.11	1.16 ± 0.11*
<i>Acc1</i>	1.07 ± 0.10	0.92 ± 0.08*	0.68 ± 0.03 [§]	1.41 ± 0.14	1.29 ± 0.07
<i>Fasn</i>	1.83 ± 0.24	1.21 ± 0.10*	0.71 ± 0.06 [§]	1.18 ± 0.11*	1.14 ± 0.08*
<i>Scd1</i>	1.64 ± 0.21	1.80 ± 0.44	0.36 ± 0.03 [§]	0.86 ± 0.14	1.02 ± 0.13
<i>Gpam</i>	1.13 ± 0.11	1.14 ± 0.12	1.02 ± 0.16	1.15 ± 0.11	1.23 ± 0.07
eNOS	1.60 ± 0.21	1.14 ± 0.11*	0.96 ± 0.09*	0.91 ± 0.12*	0.99 ± 0.10*

Pgc-1 α , peroxisome proliferator-activated receptor gamma coactivator 1- α .

Results are expressed relative to *Hprt* gene expression (mean ± SEM, $n = 10$ per group). Dietary interventions: standard diet (control), high fat-high sucrose diet (DIO) and DIO enriched in citrulline (2.5 g kg⁻¹) (DIOcit), atorvastatin (10 mg kg⁻¹) (DIOstat) or citrulline and atorvastatin (DIOcit-stat).

* $P < 0.05$, significantly different from controls;

[§] $P < 0.05$, significantly different from control and other DIO groups.

[#] $P < 0.05$, significantly different from other DIO groups.

Table 5

Impact of refeeding on mRNA content of major proteins involved in the regulation of liver *de novo* lipogenesis

	Control	DIO	DIOcit-stat
<i>Srebf1</i>	4.54 ± 0.95*	0.60 ± 0.09 ^{#*}	0.90 ± 0.09 [§]
<i>Lxra</i>	1.37 ± 0.17	0.98 ± 0.17	1.08 ± 0.13
<i>Chrebp</i>	0.84 ± 0.12	0.69 ± 0.07*	0.81 ± 0.07
PPAR α	0.33 ± 0.05*	0.53 ± 0.09*	0.46 ± 0.04*
<i>Pgc-1α</i>	0.48 ± 0.05*	0.41 ± 0.07*	0.46 ± 0.10*
PPAR γ	0.96 ± 0.21	0.70 ± 0.14	0.86 ± 0.12
<i>Fasn</i>	2.96 ± 0.99*	1.07 ± 0.08 [#]	2.18 ± 0.32*
<i>Acc1</i>	3.03 ± 0.93*	1.12 ± 0.21	1.61 ± 0.27 *
<i>Scd1</i>	0.48 ± 0.07*	0.81 ± 0.23	1.03 ± 0.34
<i>Gpam</i>	1.30 ± 0.07	0.84 ± 0.08 [#]	0.75 ± 0.03 ^{#*}
eNOS	0.89 ± 0.15	1.10 ± 0.10	1.14 ± 0.22

Hprt, hypoxanthine guanine phosphoribosyl transferase; *Pgc-1 α* , peroxisome proliferator-activated receptor gamma coactivator 1- α .

Results are expressed relative to fasted values (mean ± SEM; $n = 5$ to 7 samples per group). Dietary interventions: standard diet (control), high fat–high sucrose diet (DIO) and DIO enriched in citrulline (2.5 g kg⁻¹ body weight) and atorvastatin (10 mg kg⁻¹ body weight) (DIOcit-stat).

* $P < 0.05$, significantly different from fasted values;

[§] $P < 0.05$, significantly different from control and DIO;

[#] $P < 0.05$, significantly different from control.

receiving citrulline alone did not show any reduction in fat mass in comparison with DIO mice. The direct effect of citrulline on adipose tissue fatty acid release may partly explain why in our model, fat mass was significantly reduced in DIOcit-stat animals whereas daily energy balance was not significantly altered. However, the combination of citrulline and atorvastatin appears necessary to attain a significant reduction in fat mass accretion under DIO conditions.

We hypothesized that the synergistic effects of the combination involved an improved production of NO by eNOS as described by Berthe *et al.* (2011) in endothelial cells. In the present study, the metabolic adaptations induced *in vivo* by the combination of citrulline and atorvastatin were consistent with those observed on glucose homeostasis, visceral obesity and liver steatosis, when NO production was enhanced (Duplain *et al.*, 2001; Cook *et al.*, 2004; Schild *et al.*, 2008). Our observations on adipose tissue mass are also in agreement with those obtained in overweight rats (Joffin *et al.*, 2014). In that study, citrulline stimulated visceral adipose tissue lipolysis, and the effect was abolished by pretreatment of explants with L-NAME (Joffin *et al.*, 2014). Furthermore, our observations on plasma amino acid concentrations showed that atorvastatin interferes with citrulline and arginine metabolism, thus corroborating a possible increase in NO production. Finally, NO is known to induce AMPK α phosphorylation in several tissues (Cardaci *et al.*,

2012; Abudukadier *et al.*, 2013). *In vitro* using HuH7 cells, we confirmed that atorvastatin induced AMPK α phosphorylation (Sun *et al.*, 2006) and showed for the first time that citrulline can also activate this kinase. Experiments using L-NAME also suggest that NO may mediate part of the effect of citrulline and atorvastatin on AMPK phosphorylation. These observations suggest but do not demonstrate the potential involvement of NO in the metabolic adaptations described *in vivo* in the present study. Thus, a limitation of our work lies in the fact that we did not directly look at the involvement of NO in the *in vivo* metabolic adaptations. Additional researches, using NOS inhibitors or through the infusion of citrulline and arginine labelled with stable isotopes (Castillo *et al.* 1996; Fitzgerald *et al.* 2007), are required to gain insight into the effect of the treatment on NO synthesis at the whole-body and tissue levels.

In conclusion, combination treatment with citrulline and atorvastatin preserved glucose homeostasis and prevented fat mass accumulation in a mouse model of DIO. Furthermore, it potentiated the inhibition of SREBP-1 mediated *de novo* lipogenesis activity in the liver, partly by up-regulating phosphorylation of AMPK α . Citrulline–atorvastatin combinations are thus a promising treatment for preventing the alteration of glucose homeostasis and minimizing the risk of developing type 2 diabetes in patients treated with statins and presenting cardiometabolic risk factors.

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Author contributions

F. C., G. C., L. C., C. M. and B. M. contributed to the study conception and design, acquisition of data or analysis and interpretation of data, to drafting the article or revising it critically, and gave final approval of the version to be published. J.-P. B. contributed to the article revision. E. P., J.-P. R., S. L. P., C. D. and C. J. contributed to the study conception, acquisition of data and analysis and interpretation of data.

Conflict of interest

Christophe Moinard, Luc Cynober, Jean-Pascal de Bandt and Servane Le Plenier are shareholders of CITRAGE© (Créteil, France), suppliers of citrulline.

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