



Role of angiotensin type 2 receptor in improving lipid metabolism and preventing adiposity

Sourashish Nag¹ · Sanket Patel¹ · Shailaja Mani^{2,3} · Tahir Hussain¹

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Abstract

Recent studies on mice with null mutation of the angiotensin type 2 receptor (AT₂R) gene have implicated the involvement of AT₂R in regulating adipocyte size and obesity, a major risk factor for metabolic syndrome. However, the outcome from these studies remains inconclusive. Therefore, current study was designed to test whether pharmacological activation of AT₂R regulates adiposity and lipid metabolism. Male mice (5-weeks old) were pre-treated with vehicle or AT₂R agonist (C21, 0.3 mg/kg, i.p., daily, for 4 days) and fed normal diet (ND). Then these animals were subdivided into ND and high-fat diet (HFD) regimen and concomitantly treated with vehicle or C21 through day 14. Vehicle-treated HFD-fed mice demonstrated an increase in epididymal white adipose tissue (eWAT) weight and adipocyte size, which were associated with increased eWAT expression of the lipogenic regulators, fatty acid binding protein and fatty acid synthase, decreased expression of adipose triglyceride lipase and increased expression of hormone-sensitive lipase. Interestingly, C21 pre-treatment altered HFD-induced changes in lipogenic and lipolytic regulators. C21 pre-treatment prevented decrease in expression of uncoupler protein-1 in brown adipose in HFD-fed mice, which was associated with increased core temperature. In addition, C21 pre-treatment ameliorated plasma-free fatty acids, triglycerides, insulin and tumor necrosis factor- α in HFD-fed mice. *Ex-vivo* study in isolated primary epididymal adipocytes revealed that C21 inhibits long chain fatty acid transporter, via a nitric oxide synthase/guanylate cyclase/protein kinase G-dependent pathway. Collectively, we propose pharmacological activation of AT₂R regulates fatty acid metabolism and thermogenesis and prevents HFD-induced adiposity in mice.

Keywords Adipocyte · AT₂R · Fatty acid transporter (FATP) · Lipid metabolism · Adiposity

Introduction

Obesity, characterized by increase in body weight and adiposity, is a major risk factor for the development of dyslipidemia, atherosclerosis, insulin resistance, diabetes and hypertension [1–5]. The incidence of obesity has been

increasing worldwide and is now considered a global epidemic. According to the international guidelines, the lifestyle modification (decreased caloric intake and increased physical exercise) is a primary modality of body weight management [6]. However, the long-term effects of diet and exercise on weight have been mostly disappointing [7]. Further strategies for body weight management include pharmacological intervention and bariatric surgery [8]. But most anti-obesity drugs are associated with serious side effects [9]. Thus, there has been a pressing need for the discovery of newer anti-obesity drugs.

Lipogenesis, lipolysis and thermogenesis are some determinants of body weight gain and adiposity. Under fed condition, the surplus or extra energy from lipid is stored in adipocytes as triglycerides (TAG) for future use [10]. As and when the energy demand increases, the stored TAGs are broken down to free fatty acids. Fatty acid binding protein-4 (FABP4) and fatty acid synthase (FASN) (lipogenic), adipocyte triglyceride lipase (ATGL) and hormone-sensitive

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✉ Tahir Hussain
thussain@central.uh.edu

¹ Department of Pharmacological & Pharmaceutical Sciences, College of Pharmacy, University of Houston, 4849 Calhoun Rd., Health 2, Houston, TX 77204, USA

² Department of Molecular and Cellular Biology, Baylor College of Medicine, Houston, TX 77030, USA

³ Department of Neuroscience, Baylor College of Medicine, Houston, TX 77030, USA

lipase (HSL) (lipolytic) and uncoupling protein-1 (UCP1) (thermogenic) are some of the enzymic regulators of these processes. Fatty acid transporter protein (FATP) is a family of plasma membrane transporters regulating long chain fatty acids (LCFA) uptake into adipocytes [11]. Knockdown of FATP1, which is an insulin-sensitive LCFA transporter, prevents HFD-induced resistance in skeletal muscle [12]. Also, the FATP4 expression in human adipose tissue positively correlates with obesity and insulin resistance [13].

Although renin angiotensin system (RAS) is mostly known for its role in cardiovascular function and blood pressure regulation, it has been recently implicated to have a role in lipid metabolism and adiposity [14–16]. Adipocytes are known to have all the components of the RAS. In high-fat diet (HFD)-induced obesity, an increased formation of angiotensin-II (ang-II) in adipose tissue has been reported. Ang-II is thus considered as a trophic factor for the adipose tissue growth. It acts via angiotensin type 1 and type 2 receptors (AT₁R and AT₂R). The physiologic effects of these receptors generally are antagonistic [17, 18]. While the blocking of AT₁R decreases adipocyte size in mice [19], the role of AT₂R in adiposity is not yet established. Recent studies demonstrated that the genetic deletion of AT₂R resulted in a decrease in adiposity in mice [20, 21]. On the contrary, deficiency of AT₂R in Apolipoprotein E-knockout (KO) mice (a model of atherosclerosis) increased adipose tissue weight in mice [22]. Thus, the role of AT₂R on adiposity in knockout studies has remained inconclusive. Therefore, we designed the current study to investigate the effects of the AT₂R agonist C21, a novel non-peptide and orally active compound, on lipid metabolism and adiposity.

Materials and methods

Animals and experimental protocols

Twenty-four male C57BL/6 mice (5-week-old) were obtained from Harlan (Indianapolis, IN). The mice were housed at the University of Houston's animal care facility with free access to food and water and maintained under a 12-h light/dark cycle. The animal experimental protocols were approved by Institutional Animal Care and Use Committee and performed in accordance with the National Institutes of Health Guidelines.

Mice were randomized and pre-treated with 'Compound 21' (C21) at a dose of 0.3 mg/kg [23, 24] or equivolume of normal saline (i.p., daily, days 1–4 at 10 am) and fed normal diet (29% protein, 56% carbohydrate and 15% fat) (referred as ND, 7022). Thereafter the mice on C21 and vehicle were further sub-divided into two groups: ND or high-fat diet (18.4% protein, 21.3% carbohydrate and 60.3% fat) (referred as HFD, TD.06414) for the next 10 days along

with concurrent treatment with saline or C21. Sodium salt of C21 (PubChem CID: 9804984) was provided as a gift from Vicore Pharma, Sweden. Food intake and body weight of the mice were measured as reported previously [21]. Core temperature was measured by mice rectal probe (RET-3) and thermocouple meter (WD-35627-00) (Kent Scientific corp., Torrington, CT) on terminal days 12–14 in morning immediately after drug administration. At the end of day 14, mice were euthanized in non-fasting state by cervical dislocation under isoflurane anesthesia. Plasma and brown adipose tissue (BAT) were collected and stored at -80°C until further use. Epididymal white adipose tissue (eWAT) pads were removed, patted dry, weighed and a part of the tissue was preserved in buffered formalin for histology. The remaining eWAT was stored frozen at -80°C for biochemical measurements. Details on various assay methods are available in the online-only Data Supplement.

Statistical analysis

The data were analyzed using GraphPad Prism (GraphPad software, San Diego, CA). In vitro results are subjected to one-way ANOVA with Newman–Keuls post hoc test and unpaired Student's *t* test. To determine diet–treatment interaction, in vivo results are subjected to two-way ANOVA followed by Fisher's LSD test at $p < 0.05$ to be considered statistically significant. Data are presented as mean \pm SEM.

Results

Calorie intake and body weight

Food intake was calculated in terms of total kilo calorie (kcal) consumed during 2 weeks of treatment period. As shown in Table 1, HFD-fed mice had significantly higher kcal intake in 2 weeks (ND 64 ± 4.0 vs. HFD 130 ± 5.7 kcal) compared with ND group. The C21 pre-treatment did not affect the kcal intake under ND or HFD conditions in these mice. Body weight gain during the treatment period remained same in all the groups (Table 1). We did not observe adverse events related to animals' health during the study.

Epididymal WAT weight/body weight ratio and epididymal adipocyte size

The hematoxylin and eosin staining of the eWAT sections from control and C21-treated mice on ND and HFD is shown in Fig. 1a. Compared with ND, HFD caused an increase in epididymal adipocyte size (ND 1118 ± 54 vs HFD $2636 \pm 79 \mu\text{m}^2$), which was reduced by C21-treatment ($1407 \pm 102 \mu\text{m}^2$) (Fig. 1b). C21 treatment of ND-fed mice

Table 1 Metabolic and hormonal parameters and rectal body temperature of control and C21-treated 5-week-old male C57BL/6 mice fed with ND or HFD

Parameters/mice groups	ND	ND-C21	HFD	HFD-C21	p value		
					Diet	Treatment	Interaction
Total kcal intake (2 weeks)	64 ± 4	55 ± 5	130 ± 6*	133 ± 7	0.0001	n.s.	n.s.
Body weight gain (g) (terminal – initial)	3.43 ± 0.95	3.38 ± 0.73	4.13 ± 0.95	3.38 ± 0.53	n.s.	n.s.	n.s.
Initial body weight (g)	17.00 ± 1.33	16.88 ± 1.19	16.00 ± 1.30	16.75 ± 0.56	n.s.	n.s.	n.s.
Terminal body weight (g)	20.43 ± 0.57	20.29 ± 0.81	20.13 ± 0.44	20.13 ± 0.44	n.s.	n.s.	n.s.
Plasma FFA (nmol/μL)	2.30 ± 0.20	2.30 ± 0.18	5.24 ± 0.36*	3.27 ± 0.19 [#]	0.0001	0.0002	0.0002
Plasma TAG (nmol/μL)	1.96 ± 0.23	2.20 ± 0.34	4.03 ± 0.25*	3.30 ± 0.18 [#]	0.0001	n.s.	0.0408
Plasma insulin (ng/mL)	0.78 ± 0.06	0.73 ± 0.06	2.34 ± 0.09*	1.33 ± 0.19 [#]	0.0001	0.0001	0.0004
Plasma TNF-α (pg/mL)	35 ± 3	34 ± 4	72 ± 16*	43 ± 4 [#]	0.0147	n.s.	n.s.
Plasma adiponectin (ng/mL)	4.41 ± 0.19	3.60 ± 0.55	2.68 ± 0.43*	5.24 ± 0.68 [#]	n.s.	n.s.	0.0029
Rectal body temperature days 12–14 (°C)	37.01 ± 0.19	37.29 ± 0.18	37.39 ± 0.13	38.04 ± 0.13 [#]	0.0021	0.0087	n.s.

ND normal diet, ND-C21 normal diet on C21, HFD high-fat diet, HFD-C21 high-fat diet on C21

*Significantly different from ND

[#]Significantly different from HFD; data were analyzed using two-way ANOVA with Fisher's LSD test at $p < 0.05$

had no effect on the weight ratio or the adipocyte size. High-fat diet caused significant increase only in the eWAT weight/body weight ratio (Fig. 1c), while other fat depots weight remained unchanged among treatment groups (SI Table 1). C21-treatment reduced the HFD-induced increase in eWAT and eWAT weight/body weight ratio (by 34%).

Plasma FFA and TAG

The HFD-fed mice had significantly increased levels of plasma FFA (ND 2.30 ± 0.20 vs. HFD 5.24 ± 0.36 nmol/μL), which were reduced by C21-treatment (HFD-C21 3.27 ± 0.19 nmol/μL) (Table 1). Similarly, HFD-feeding increased the plasma TAG levels, which were attenuated by C21 treatment (HFD 4.03 ± 0.25 vs. HFD-C21 3.3 ± 0.18 nmol/μL). Both TAG and FFA in ND-fed mice were not affected by C21 treatment (Table 1).

Plasma insulin, TNF-α and adiponectin

The plasma insulin level (non-fasting) was nearly three-fold higher in HFD-fed mice (ND 0.78 ± 0.06 vs. HFD 2.34 ± 0.09 ng/mL). This increase in plasma insulin in HFD-fed mice was attenuated (HFD-C21 1.33 ± 0.19 ng/mL) by C21 treatment, which had no effect in ND-fed mice (Table 1). The HFD feeding significantly increased plasma TNF-α levels (ND 35 ± 2.6 vs. HFD 72 ± 15.8 pg/mL), which were significantly reduced (HFD-C21 43 ± 4.5 pg/mL) by C21 treatment (Table 1). On the other hand, plasma adiponectin was significantly decreased by HFD feeding compared with ND (ND 4.41 ± 0.19 vs. HFD 2.68 ± 0.43 ng/mL). The C21 treatment significantly increased the adiponectin levels (HFD-C21

5.24 ± 0.68 ng/mL) under HFD condition but had no effect in the ND-fed mice (Table 1).

Lipogenic regulators' protein expression in eWAT

Western blotting revealed the presence of the lipogenic regulators fatty acid-binding protein-4 (FABP4) (15 kDa) and fatty acid synthase (FASN) (273 kDa) protein expression in the eWAT. Densitometric analysis of the bands suggested that FABP4 (99%) and FASN (68%) expression in eWAT in HFD was significantly greater than ND (Fig. 2a and b, respectively). C21-treatment under HFD condition reduced the expression of both the proteins—FABP4 (by 26%) and FASN (by 62%).

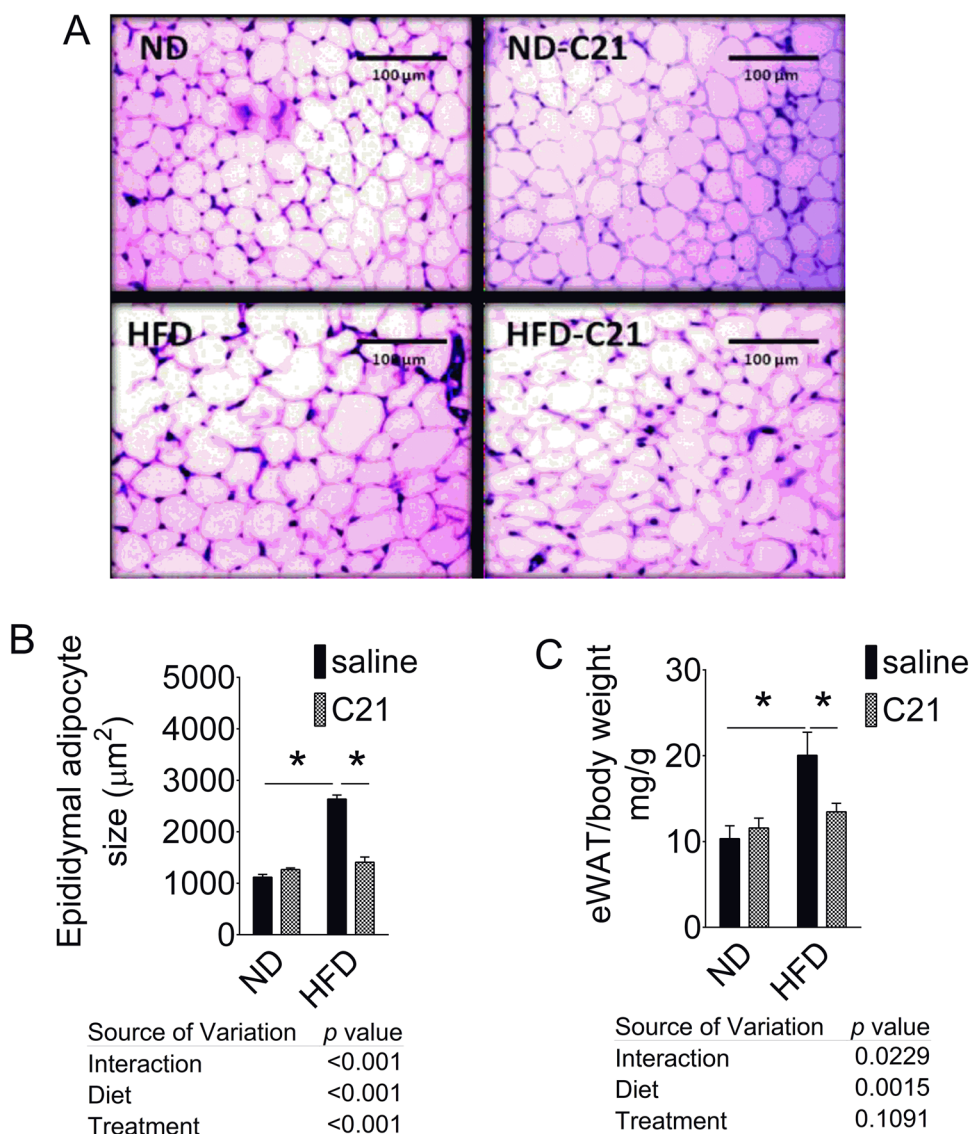
Lipolytic regulators' protein expression in eWAT

Distinct bands for adipose triglyceride lipase (ATGL) and hormone-sensitive lipase (HSL) were detected approximately at 56 kDa and 84 kDa, respectively, in the eWAT by western blotting. Densitometric analysis of the bands revealed that compared with ND, HFD-fed mice expressed reduced ATGL (16%) and increased HSL (62%) proteins (Fig. 2c and d, respectively). C21 treatment of HFD-fed mice caused an increase in the ATGL expression (47%) and decreased the HSL expression (52%). Surprisingly, C21 treatment under ND condition significantly increased the protein expression of HSL (50%) (Fig. 2d).

Fatty acid uptake by isolated primary epididymal adipocytes

With the increasing number of epididymal adipocytes in the assay tube, there was an increase in the FA uptake, which

Fig. 1 Epididymal WAT weight-to-body weight ratio and adipocyte size of control and C21-treated mice fed with ND or HFD. **a** Epididymal WAT weight-to-body weight ratio; $N=6-8$ in each group. **b** Representative micrograph of epididymal adipose tissue (magnification, $\times 10$). **c** Epididymal adipocyte size was expressed as cell area (μm^2); $N=3$ (3 sections per slide from 3 mice). Results are mean \pm SEM; *significantly different compared with ND, #significantly different compared with HFD. Data were analyzed using two-way ANOVA followed by Fisher's LSD test at $p < 0.05$ (ND normal diet, ND-C21 normal diet on C21, HFD high-fat diet, HFD-C21 high-fat diet on C21)



plateaued off at ~ 3600 s after initiating the assay. After conducting preliminary experiments, we chose 40,000 cells/assay for further experiments and terminated the assay at 4200 s. The FA uptake was inhibited by the FATP inhibitor, triacsin C (Fig. 3a) in a concentration (1–20 μM)-dependent manner suggesting the involvement of FATP. Further, FA uptake was stimulated by insulin in a dose-dependent manner (Fig. 3b), with a significant increase at ≥ 10 nM concentrations of insulin.

Adipocyte FA uptake

Figure 4 depicts FA uptake was significantly decreased by C21 at 1 μM concentration. A lower dose (0.1 μM) of C21 was not effective and a higher dose (5 μM) did not cause any further reduction in FA uptake (Fig. 4a). In subsequent experiments, we used 1 μM of C21. The C21-induced

decrease in FA uptake was blocked by the AT_2R antagonist PD123319 (10 μM), but not by the AT_1R antagonist losartan (1 μM) (Fig. 4b) suggesting the involvement of AT_2R . To understand the potential role of NO/cGMP signaling pathway in AT_2R -mediated inhibition of FA uptake in adipocytes, we utilized specific inhibitors. The C21 inhibition of FA uptake was blocked by the NO synthase (NOS) inhibitor L-NAME (1 mM), the guanylyl cyclase (GC) inhibitor ODQ (10 mM) or the cGMP-dependent protein kinase-G (PKG)-inhibitor Rp-8-Br-PET-cGMPS (1 μM) (Fig. 4c).

Core temperature and expression of UCP1 in BAT

The treatment of AT_2R agonist C21 significantly increased body temperature in HFD-fed mice, but not in ND-fed mice (Table 1). HFD feeding did not affect body temperature as compared to ND-fed mice. The expression of UCP1 was

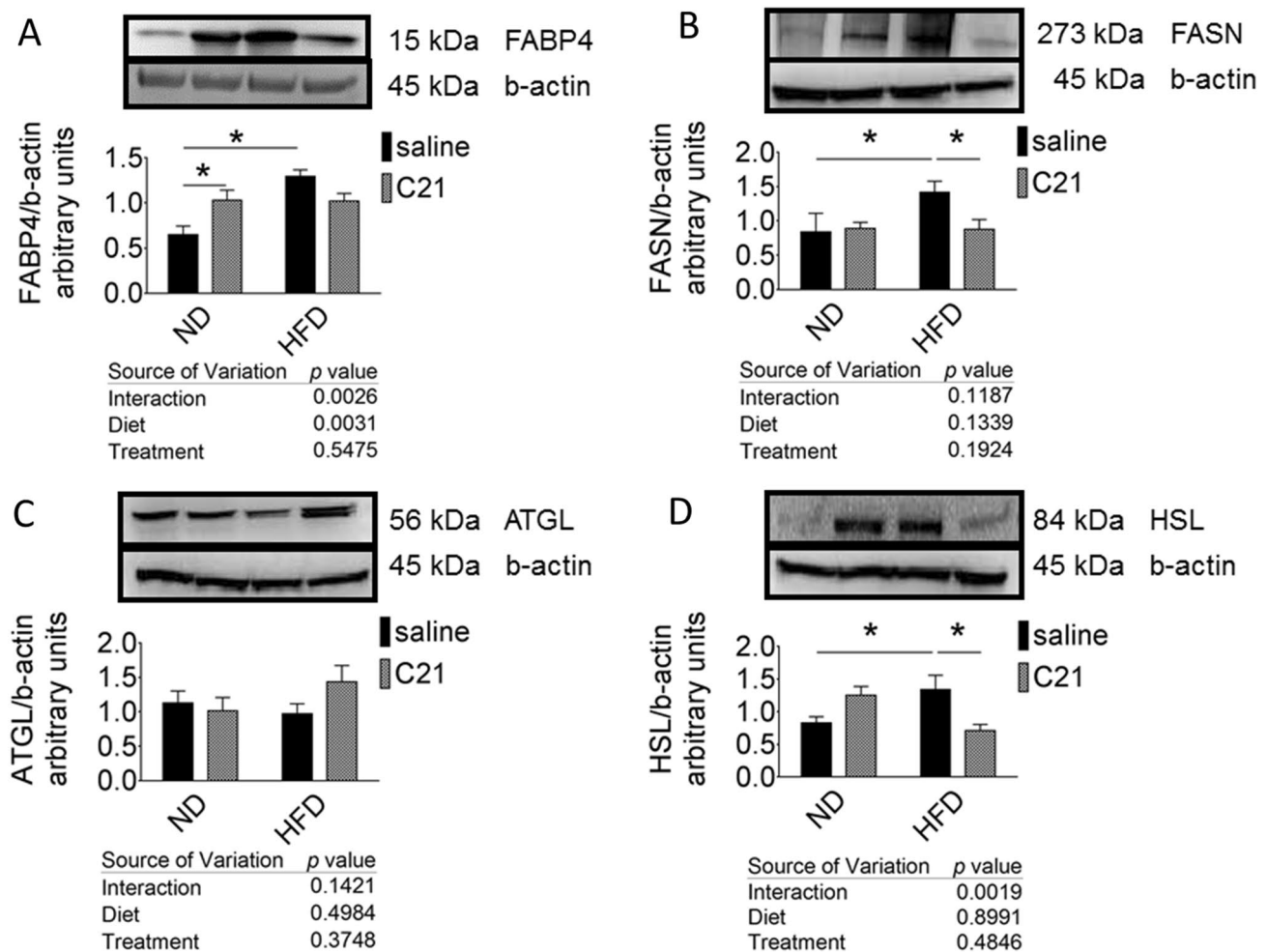


Fig. 2 Protein expression of lipogenic and lipolytic regulators in the epididymal WAT of control and C21-treated mice fed with ND or HFD. Representative western blots for **a** FABP4, **b** FASN, **c** ATGL and **d** HSL with loading control β -actin. The bar graphs represent the ratios of densities of FABP4, FASN, ATGL, and HSL normalized with β -actin protein bands. *Significantly different compared

with ND, #significantly different compared with HFD. Results are mean \pm SEM; data were analyzed using two-way ANOVA followed by Fisher's LSD test at $p < 0.05$; $N = 4-6$ in each group (ND normal diet, ND-C21 normal diet on C21, HFD high-fat diet, HFD-C21 high-fat diet on C21)

detected at 30 kDa in BAT (Fig. 5). The BAT expression of UCP1 was lower in HFD-fed mice (~30%) and C21 treatment of these mice attenuated the decrease in UCP-1 expression.

Discussion

This is a short-term study where 2-weeks HFD regimen, as expected, had no effect on body weight gain yet could initiate metabolic changes [25]. Numerous studies including ours suggest that 6–8 weeks HFD regimen is needed to cause a significant shift in body weight [21]. However, this regimen clearly shows various metabolic parameters were affected by HFD and the AT₂R agonist C21 treatment significantly prevented those changes. Particularly findings revealed that

AT₂R agonist treatment of HFD-fed mice prevented increase in adipocyte size, improved plasma indices of dyslipidemia, which were associated with the changes in expression of lipogenic (FABP4 and FASN), lipolytic (ATGL and HSL) and thermogenic (UCP1) enzymes. Additionally, the AT₂R agonist C21 inhibits primary adipocyte FA uptake via NO/cGMP pathway, suggesting adipose tissue as a potential site of AT₂R action affecting FA transport and metabolism.

The size of the adipocyte is built-up by the process of lipogenesis. In the postprandial phase, the FAs are first transported inside the adipocytes by the FATPs [26]. Once inside the aqueous cytosol, FFAs are then bound to the FABP4 for cytosolic transportation [27]. These FFAs along with the de-novo synthesized FFAs from the Krebs's cycle (by FASN) are packed and stored as TAG in the lipid droplets of adipocytes [28]. An earlier study has reported that AT₂R-KO

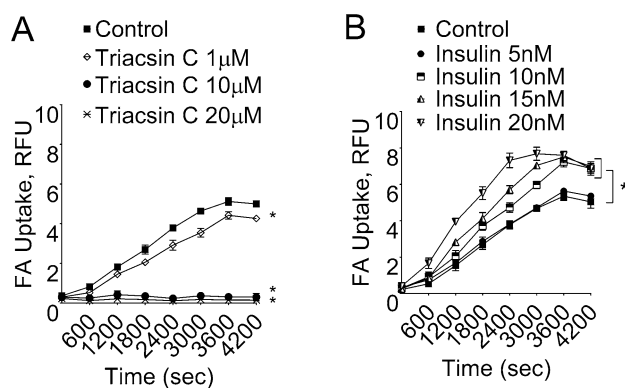


Fig. 3 Fatty acid uptake in isolated primary epididymal adipocytes assessed using fluorescence assay. **a** In absence or presence of different concentrations (1, 10 and 20 μM) of triacsin C, the FATP inhibitor and **b** in absence or presence of different concentrations (5, 10, 15 and 20 nM) of insulin, the FATP-activator. *Significantly different control adipocytes (without triacsin C treatment) (**a**), *significantly different control adipocytes (without insulin treatment) (**b**). Results are mean \pm SEM; data were analyzed using one-way ANOVA followed by Neuman–Keuls post hoc test and Student's *t* test, $p < 0.05$; $N = 3$ in each group

mice fed HFD had smaller adipocytes and decreased expression of FABP4 and FASN gene [20]. In the present study we observed, HFD significantly increased the protein expression of the eWAT lipogenic regulators (FABP4 and FASN) explaining the basis of adipocyte size increase under HFD condition.

The adipocyte size is also regulated by lipolytic enzymes (ATGL and HSL). Recently, it has been reported that the ATGL-KO mice show increased adipose mass and adipocyte size [29]. In contrast to ATGL-KO mice, the HSL-KO mice do not have increased fat deposition, body weight and adipocyte size [30, 31]. A non-coordinated regulation of ATGL and HSL mRNA as well as protein expressions have been reported during inflammatory conditions, exercise, weight loss [32] or during acute lipolytic phase, i.e., in the lipid mobilization phase and the protein breakdown phase such as fasting [33]. In concurrence with these observations, in the present study we observed that along with the increased eWAT weight, HFD dysregulated lipolysis by decreasing the ATGL expression which might have led to increased HSL expression in adipocytes, suggesting a differential regulation of these enzymes by the HFD and plausibly accounts for the increased levels of plasma insulin and TNF- α and reduced adiponectin [34]. However, the AT₂R agonist treatment significantly altered the protein expression of both the epididymal WAT lipolytic regulators. The increased level of ATGL by C21 under HFD condition might be critical in regularizing the imbalanced energy homeostasis state in obesity. However, the significant increase in protein expression of HSL by C21 treatment under ND condition could plausibly be due to the body's physiological increase of the protein levels to compensate the reduction of FA uptake and restore the lipid metabolism under ND condition. Since this study measures only the protein expression, which is not a direct

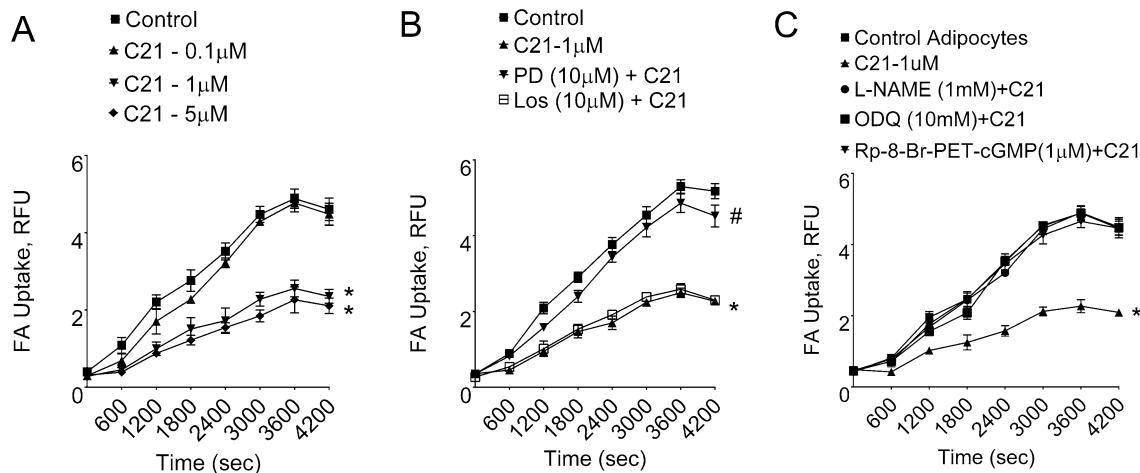


Fig. 4 Fatty acid uptake in isolated primary epididymal adipocytes assessed using fluorescence assay. **a** In absence or presence of different concentrations (0.1, 1.0 and 5.0 μM) of C21, the AT₂R agonist, **b** in absence or presence of C21 (AT₂R agonist), presence of PD (PD123319-AT₂R antagonist)+C21 and presence of losartan (AT₁R antagonist)+C21, and **c** in absence and presence of C21 (AT₂R agonist), presence of L-NAME (NO synthase inhibitor)+C21, presence of ODQ (guanylyl cyclase inhibitor)+C21 and presence of Rp-8-Br-

PET-cGMPs (PKG-inhibitor)+C21. *Significantly different control adipocytes (without C21 treatment), #significantly different from C21 treatment group. Results are mean \pm SEM; data were analyzed using one-way ANOVA followed by Neuman–Keuls post hoc test and Student's *t* test, $p < 0.05$; $N = 3$ in each group. L-NAME-L-NG-Nitroarginine Methyl Ester; ODQ-1*H*-[1,2,4] Oxadiazolo[4,3-*a*]quinoxalin-1-one; PKG-I-PKG-inhibitor/Rp-8-Br-PET-cGMPs

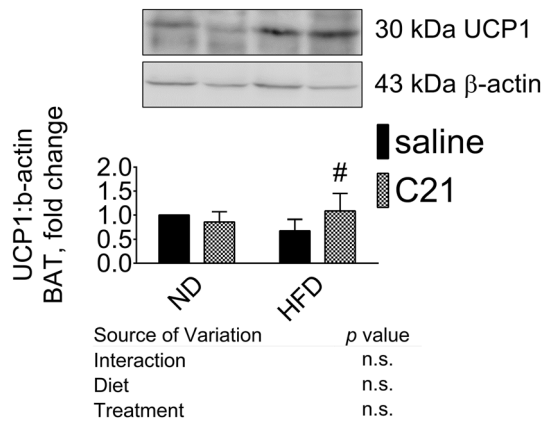


Fig. 5 Protein expression of uncoupling protein 1 (UCP1) in brown adipose tissue of control and C21-treated mice fed with ND or HFD. Representative western blots of UCP1 with loading control β -actin. The bar graphs represent the ratios of densities of UCP1 normalized with β -actin protein bands. *Significantly different compared with ND; #significantly different compared with HFD at $n=13$, 10% alpha error level and 50% beta error level. Results are mean+SEM; data were analyzed using two-way ANOVA followed by Fisher's LSD test at $p<0.05$; $N=6$ in each group (ND normal diet, ND-C21 normal diet on C21, HFD high-fat diet, HFD-C21 high-fat diet on C21)

reflection of the enzyme activities, this can be considered a limitation of these measurements.

The lipogenic and lipolytic regulators are governed by the function of the FA transport into the adipocytes [35]. In adipocytes, plasma membrane FATP facilitates the LCFA uptake [36]. Modulation of FATP function results in alteration of energy homeostasis and insulin sensitivity [37]. Treatment with angiotensin-II alone and with PD123319 (AT_2R blocker) has been shown to increase the uptake of palmitic acid (LCFA) in immortalized cardiomyocytes [38]; this indicates that the AT_2R is involved in the FA uptake. To test whether the AT_2R regulates FA transport, we performed an ex vivo study using isolated mouse primary epididymal adipocytes. Our data demonstrate that the FA uptake was reduced by the AT_2R agonist C21 in the primary epididymal adipocytes. Moreover, the C21-induced decrease in FA uptake was blocked by PD123319, indicating involvement of AT_2R . Furthermore at the molecular level, studies have shown AT_2R increases NOS activity [39]. Also it has been shown that the overexpression of NOS prevents adiposity [39]. So, we further explored whether the pharmacological activation of AT_2R reduces fatty acid uptake via NOS-dependent pathway. We observed that the pre-treatment of primary epididymal adipocytes with inhibitors of NOS/GC/PKG blocked the AT_2R -mediated decrease in FA uptake indicating the involvement of NOS/GC/PKG pathway. Whether these enzyme inhibitors alone would have their own effects (increase) on FA uptake and alter the interpretation of C21 effects are not clear thus limiting the study.

Moreover, further studies are needed to explore whether NOS/GC/PKG pathway is involved in AT_2R -mediated anti-obesity effects. Given that insulin is a potent stimulator of FA uptake (Fig. 3), these inhibitors alone would have a modest, if any, increase in FA uptake, thus unlikely to affect the net outcome on their ability to markedly attenuate AT_2R -mediated inhibition of FA uptake inhibition. This reduction in FA transport possibly as a consequence prevents the HFD-induced changes in lipogenic and lipolytic regulators and thus prevents epididymal adipocyte size and eWAT weight increase under HFD. FATP1 is a major insulin-sensitive LCFA transporter in adipose and skeletal muscle [11, 12, 40]. Inactivation of FATP1 prevents HFD-induced resistance in skeletal muscle [12]. In addition, another report indicates that the FATP4 expressions in human adipose tissue positively correlate with obesity and insulin resistance [13]. In accordance, we observed that the AT_2R agonist treatment inhibits FA uptake in adipocytes, and reduces plasma FA and TAG levels under HFD.

Thermogenesis could increase energy expenditure and reduce adiposity. Tsukuda et al. [41] recently reported that depletion of AT_1R , a counter-regulator of AT_2R , resulted in increase in thermogenic regulators in adipose tissue. Conversely, current results show that pharmacological activation of AT_2R also increased core temperature and prevented a decrease in expression of thermogenic regulator UCP1 in BAT. Similarly, chronic administration of AT_2R agonist C21 resulted in upregulation of adipose UCP1 [42]. Our findings are in accordance with other study [43] showing that activation of AT_2R by C21 increases browning of adipose tissue and increases body surface temperature.

Increased adiposity is associated with hyperlipidemia, hyperinsulinemia and increased inflammation. The larger adipocytes under HFD not only have impaired lipid metabolism but also triggered the pro-inflammatory state of obesity leading to insulin resistance [44–46]. In postprandial phase, the presence of diet stimulates the increase in plasma insulin to facilitate the uptake of glucose in cells. However, in obesity as adipocytes become resistant to insulin, the plasma FFA and TAG levels increase. The larger adipocytes also trigger the inflammation resulting in increased plasma TNF- α , a pro-inflammatory cytokine and decreased plasma adiponectin, an anti-inflammatory and insulin-sensitizing cytokine. The AT_2R agonist C21 treatment improved insulin signaling and sensitivity that was associated with increase in adipose adiponectin [42]. The AT_2R -KO mice developed insulin resistance with decreased adiponectin [20] while the AT_2R -KO atherosclerotic mice have increased plasma FFA [22]. Pharmacological stimulation of AT_2R with CGP42112A (a peptide AT_2R agonist) has also been reported to increase plasma FFA and TAG in rats [47], albeit these experiments were conducted in ND-fed rats. Another study described opposite role of AT_2R in

regulation of thermogenic capacity of adipose tissue; however, these researchers employed the mice model exhibiting transgenic activation of brain RAS and partial AT₂R agonist CGP42112a [48]. There is some evidence showing that AT₂R activation may have opposite effects in lean vs obese rats. For example, we have reported opposing effects of AT₂R agonist on inflammation in lean and obese rats, i.e., anti-inflammatory effects in obese vs inflammatory in lean rats [49]. In the present study we observed that C21 treatment in HFD condition prevented all the HFD-induced biochemical changes in young (5-week age) as well as old (12-week age) male mice (SI Table 2). Thus, C21 treatment not only prevents adiposity but also protects against hyperlipidemia, hyperinsulinemia and increased inflammation under HFD condition regardless of age. However, a direct role of AT₂R in reducing inflammation and improving insulin resistance cannot be ruled out. We and others have reported that the AT₂R exerts anti-inflammatory effects and also protects against insulin resistance [50–52].

Overall, there is a diet–treatment interaction regarding biochemical indices of metabolism (insulin, adiponectin, FFA and triglyceride), expression of lipogenic (FABP4) and lipolytic (HSL) enzymes, eWAT cell size and body weight gain indicating a dependency of the effects of AT₂R agonist on these indices upon intake of type of diet. Our *in vivo* finding showing increase in expression of FABP4 in ND-fed mice does not correlate with finding showing acute inhibition of FA uptake *in vitro*. However, this discrepancy could be related to AT₂R's facilitation of initial steps of differentiation of pre-adipocytes [51]. Literature suggest that AT₂R stimulation reduces the size of large lipid droplets once adipocytes mature and gain full capacity to store lipids [51] and that ND-fed mice treated with AT₂R agonist C21 for 12 weeks did not affect body weight gain [16, 51]. Therefore, we speculate that increase in expression of FABP4 in WAT upon C21 treatment of ND-fed mice may help maintain basal fat mass but would not promote weight gain. Our earlier study reported that AT₂R activation produces anti-adiposity effects and that beneficial effect is estrogen-independent in female mice [16], whereas the current study shows that anti-adiposity effects of AT₂R may be related to improvement in fat metabolism in terms of transport, synthesis and break-down. Additionally, we have reported that liver steatosis, not measured in the present study, caused by HFD in females is attenuated by AT₂R agonist treatment [16]. The effects of AT₂R agonist were independent of estrogen, i.e., ovary-intact and ovariectomized females responded to C21 treatment in regard to the plasma metabolic markers and adipocyte size [16]. Since the effects of C21 are estrogen-independent [16], it can be argued that the soy-based diet which might contain traces of phytoestrogens may not have any bearing on the outcome of this study. Since obesity is connected to cardiovascular disease, affects kidney function

in terms of fluid retention and is a major cause of hypertension and diabetes, long-term study may shed light as to the beneficial effects of AT₂R agonist treatment in obesity. We have shown that AT₂R agonist treatment for 2-weeks reduces high salt-induced blood pressure only in obese Zucker and not in obese rats *per se* [53]. Moreover, 2-weeks HFD regimen does not increase blood pressure and it is unlikely C21 treatment will have any effect on blood pressure in HFD-fed animals.

Conclusions

Adiposity and obesity are metabolic disorders which are intricately connected to various cardiovascular diseases, hypertension and diabetes. Present study is an attempt to identify AT₂R as new target to combat adiposity and obesity. Our results provide evidence that pharmacological activation of AT₂R prevents HFD-induced adiposity, dyslipidemia and inflammation and insulin resistance. Inhibition of fatty acid uptake by adipose and thermogenesis could be possible mechanisms responsible for these beneficial effects of the AT₂R agonist. Collectively, studies indicate AT₂R activation as a potential therapeutic approach for controlling obesity and obesity-associated disorders such as insulin resistance and inflammation.

Author contributions SN and SP designed and performed the study and acquired, analyzed and interpreted data; SN, SP, SM and TH drafted the article and revised critically for important intellectual content; SN, SP, SM and TH approved the final version of the manuscript.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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