Ellagic Acid Reduces Adipogenesis through Inhibition of Differentiation-Prevention of the Induction of Rb Phosphorylation in 3T3-L1 Adipocytes

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1. Introduction

Obesity is a risk factor for numerous metabolic disorders such as type 2 diabetes, hypertension, and coronary heart disease [1]. It is imperative to understand the precise mechanism(s) behind adiposity and weight gain in the pathogenesis of obesity. Excessive weight gain in obesity may be related to, in large part, increased adipogenesis [2]. Adipogenesis is a 2-step process through which clonal expansion of preadipocytes precedes adipocyte differentiation. These sequential events ultimately result in an increase in both number (hyperplasia) and size (hypertrophy) of adipocytes [3]. It is well perceived that overweight and obesity are immediate consequences of excessive adipogenesis [4].
3T3-L1 preadipocytes are a well-established cell line employed to study adipogenesis [5]. Hormonal inducers are capable of facilitating the differentiation process in 3T3-L1 preadipocytes including a synchronous reentry of cells into cell-cycle, mitotic clonal expansion (MCE) and adipocyte phenotypic differentiation [6]. On reaching confluence, Growth of 3T3-L1 preadipocytes is arrested at the G0/G1 cell cycle boundary. The addition of 1-methyl-3-isobutylxanthine (M), dexamethasone (D), and a pharmacological dose of insulin (I) to serum-containing medium (MDI protocol) concurrently activates the cell undergoing G1/S transition. Progression of the cell cycle from G1 to S phase is regulated by the phosphorylation status of the retinoblastoma protein (Rb) [7]. Rb not only binds transcription factors involved in cell cycle progression, but also interacts with transcription factors involved in differentiation. Rb has been shown to bind to members of the CCAAT/enhancer-binding protein (C/EBP) family of transcription factors [8]. MCE is accompanied by induction of CCAAT/enhancer-binding protein (C/EBP)β expression. Following a long lag period (~16 hrs), C/EBPβ acquires DNA-binding activity prior to transcriptional activation of peroxisome proliferator activated receptors γ (PPARγ) and C/EBPα [6]. It is well conceived that C/EBPα and PPARγ work in synchrony to turn on a transcriptional cascade of genes, such as adipocyte protein-2/fatty acid-binding proteins, to generate the adipocyte phenotype in adipogenesis [6].

Ellagic acid (EA) is present in raspberries, strawberries, walnuts, and pomegranate. As with other polyphenols, EA has chemoprotective activities, with growth-inhibiting and apoptosis-promoting properties in cancer cells [9]. Several in vivo and in vitro studies confirmed its antioxidative, anti-inflammatory, and antitumorigenic properties. Several studies showed that exposure of cancer cells to EA arrested cell growth at the G0/G1-phases [10, 11]. Data from a study in humans showed that exposure of cancer cells to EA arrested cell growth at the G0/G1-phases [16, 17]. However, the effect of EA on preadipocyte adipogenesis as well as the underlying mechanism(s) remains elusive.

In the end, the present study was designed to examine the effect of EA on adipogenesis and the underlying mechanism(s) involved. Our data revealed that EA significantly suppressed hormone-induced adipogenesis in dose-dependent manner. And it is plausible that the inhibition was induced through blocking the early stage of adipogenesis, MCE, which is required for adipogenic terminal differentiation. This inhibition may be due to impaired Cyclin A protein expression and retinoblastoma protein (Rb) phosphorylation, an important cell cycle regulator that is critical for adipocyte differentiation.

2. Materials and Methods

2.1. Cell Culture. Low passage 3T3-L1 preadipocytes were kindly provided by Dr. Qiqun Tang from the Institute of Biological Science, Fudan University (Shanghai, China). Cells were cultured in Dulbecco’s Modified Eagle Medium (DMEM) (GIBCO) with 10% fetal bovine serum (FBS) (GIBCO). Two days after reaching confluence (day 0), differentiation was induced by incubation of cells in a differentiation medium containing human recombinant insulin (1μg/mL, Roche), dexamethasone (390ng/mL, Sigma), and isobutyl-methylxanthine (115μg/mL, Sigma). Two days later, media were replaced with DMEM supplemented with 10% FBS and insulin (1μg/mL). Cells were subsequently refed with DMEM supplemented with 10% FBS every two days until day 8 [6]. EA was dissolved in DMSO (stock concentration at 5mg/mL). For cell proliferation, cells were maintained in the differentiation medium with EA (0, 10, 15, and 20μM) (replaced every two days from day 0 to day 8) in culture disks. For determination of Cyclin A, cdk2 protein expression, and retinoblastoma protein (Rb) phosphorylation, EA was added only for 4, 8, 16, 20, and 24 hours with adipogenic inducer. Cell differentiation was observed daily [15].

2.2. Cell Viability. Cell viability was examined using trypan blue exclusion test. Trypan blue stain was prepared fresh as a 0.4% solution in 0.9% sodium chloride. The cells were washed in PBS, trypsinized, and centrifuged. Twenty microlitres of cell suspension were added to 20μL of trypan blue solution in 0.4% solution in 0.9% sodium chloride. The cells were washed in PBS and resuspended in an ice-cold lysis buffer containing 50mM Tris, 150mM sodium chloride, 2mM ethylene diamine tetra acetic acid (EDTA), 1mM phenylmethyl sulfonyl fluoride (PMSF), 1% NP-40, 1% sodium deoxycholate, 0.1% sodium dodecyl sulfate (SDS), 50mM sodium fluoride, 1mM sodium orthovanadate, 15mM sodium pyrophosphate, 10mM β-glycerophosphate, and a protease inhibitor cocktail tablet (Roche). The lysates were centrifuged for 10 min (12,000 rpm at 4°C). Samples were then resolved on 10–12% SDS-polyacrylamide gels for separation of specific protein followed by electrophoretic transfer to a PVDF membrane (Millipore). Primary antibodies (cell signaling) against phospho-Rb, total Rb, and aP2, PPARγ, C/EBPα, β-actin (as loading
concentrations of 0, 10, 15, and 20 M were added to a 6 cm dish and treated for 48 h overnight at 4 °C with a dilution of 1:1000. After incubating with the secondary antibody 1 h at the room temperature, the membrane was detected using the ECL Western blotting analysis system [6, 17, 18].

2.5. Nuclear Protein Extract for C/EBPβ and C/EBPα-Binding Activity Test. For DNA-binding assay, nuclear fractions were prepared (Thermo, Nuclear protein extraction kit) for C/EBPβ and C/EBPα at given time following adipogenic induction. Then 5 μg of nuclear extracts was used to assess levels of C/EBPβ and C/EBPα-specific DNA-binding activity using the TransAM ELISA kit (Active motif, Carlsbad, CA, USA), according to the manufacturer’s instructions [15, 17].

2.6. Assessment of Mitotic Clonal Expansion. For enumeration studies, 3T3-L1 preadipocytes undergoing differentiation were rinsed, trypsinized, and counted on the indicated days using a Neubauer hemacytometer. EdU labeling was performed as previously described in the report [19]. Cells were induced to differentiation with standard protocol. In different time course after induction, cells were labeled for 2 h with 10 mM EdU before fixation with 4% paraformaldehyde. EdU was stained by incubating for 20 min with 100 mM Tris (from 2 M stock, pH 8.5), 0.5 mM CuSO₄, 10 mM fluorescent azide (from 100 mM stocks in DMSO), and 50 mM ascorbic acid (added last to the mix from a 0.5 M stock in water). The staining mix was prepared fresh each time and used for staining cells immediately after the addition of ascorbate. Cells were then washed and counterstained with Hoechst. The incorporation of EdU was visualized by fluorescence microscopy. At each step described above, the cells were washed with PBS three times after treatment.

For FACS analysis, cells were trypsinized, fixed with 2% (wt/vol) paraformaldehyde in 1× PBS. Then they were treated with 0.5 mg/mL RNase A for 1 h at room temperature and incubated with 0.1 mg/mL PI (Sigma) for 45 min at 37°C. DNA content was determined by flow cytometry analysis [19].

2.7. Cell Apoptosis Analysis by Annexin V-FITC/PI Double Staining. Annexin V-FITC/PI double staining of the cells was determined using the Annexin V-FITC kit (ANNEX100F, SEROTEC, UK). To detect early apoptosis, late apoptosis, and necrosis induced by EA, 3T3-L1 preadipocytes (1 × 10⁶ cells/dish) were added to a 6 cm dish and treated for 48 h at 37°C in 5 mL of culture medium containing EA at final concentrations of 0, 10, 15, and 20 μM. 3T3-L1 preadipocytes (1 × 10⁶) were then stained for 30 min at room temperature with FITC-conjugated Annexin V-FITC and PI in a Ca²⁺-enriched binding buffer (Annexin V-FITC kit) and analyzed by a FACScan flow cytometer. Annexin V-FITC and PI emissions were detected in the FL1 and FL2 channels of a FACScan flow cytometer, using emission filters of 525 and 575 nm, respectively. The Annexin V-FITC−/PI− population was regarded as normal healthy cells, while the Annexin V-FITC+/PI− cells were taken as a measure of early apoptosis, Annexin V-FITC+/PI+ as late apoptosis, and Annexin V-FITC−/PI+ as necrosis. Approximately 1 × 10⁴ counts were made for each sample. The percentage of apoptotic cells was calculated using CELL Quest software [20].

2.8. Statistical Analysis. Statistical significance was determined using Student’s t-test (for comparison between groups), or ANOVA followed by a Tukey analysis test (for multiple comparison), and differences with P values <0.05 were considered significant.

3. Results

3.1. Effect of EA on 3T3-L1 Cell Viability and Survival. The Trypan blue dye exclusion test revealed subtle change in 3T3-L1 cell viability in response to EA (0, 10, 15, and 20 μM) challenge for 24, 48, and 96 hrs. Cells treated with 0.05% DMSO were used as a solvent control. EA obviously had no significant cytotoxicity against 3T3-L1 preadipocytes (Figure 1).

3.2. EA Inhibits 3T3-L1 Adipocyte Differentiation. Using the 3T3-L1 preadipocyte model, the effect of EA (0, 10, 15, and 20 μM) on adipogenic differentiation was examined using Oil Red O staining (indicative of lipid droplet formation). With hormone cocktail administration, 3T3-L1 preadipocytes were exposed to EA and were refed every two days. Adipocyte differentiation assessed on day 8 under light microscopy revealed a robust reduction in neutral lipid content in cells treated with EA throughout the entire differentiation period. EA treatment inhibited conversion of preadipocytes to mature lipid-storing adipocytes at the concentrations of 15 μM and the above, the result shown by Oil Red O staining (indicative of lipid droplet formation) (Figure 2(a)). Roscovitine, a cyclin-dependent kinase 2 inhibitor, was used
Figure 2: Inhibition of 3T3-L1 adipocyte differentiation by EA. (a) Representative photographs showing the cell monolayers stained with Oil Red O (×100). 3T3-L1 preadipocytes were seeded, cultured until 2 day after confluence, and then differentiated by incubating in the growth medium containing 10% fetal bovine serum in the presence of IBMX, dexamethasone, and insulin (MDI). Either EA or roscovitine was added to the incubation medium at the time inducing cell differentiation with MDI. At day 9, the cells were stained with Oil Red O. (b) Immunoblottings for adipogenic markers. Cell lysates were subjected to immunoblottings for PPARγ or aP2. Immunoblotting for β-actin verified equal loading of proteins. (c and d) Densitometric data are expressed as mean ± SD of 4 independent experiments. * P < 0.05 versus control group (without EA treatment).
3.3. EA Inhibition of Preadipocyte Clonal Expansion. We continued to examine the effect of EA on 3T3-L1 adipocyte differentiation with the purpose of identifying the inhibitory time period of EA on hormone-induced adipogenesis. The different protocol of EA challenge was used (Figure 3(a), left panel). In the experiment, the inhibition of EA on adipogenesis was caused by EA treatment during the early stage (day 0–4) of differentiation period of time. In contrast, preadipocytes exposed to EA from day 4 to day 8 were mostly differentiated to adipocytes, suggesting that EA might have an inhibitory effect at an initial stage (Figure 3(a)). The inhibition of EA on adipogenesis in the experiments was confirmed by alterations in the expression of PPARγ and aP2 (Figures 3(b), 3(c), and 3(d)), the specific adipogenic markers.

When induced to differentiate, growth-arrested 3T3-L1 preadipocytes synchronously reenter the cell cycle and undergo mitotic clonal expansion (MCE) followed by expression of genes that produce the adipocyte phenotype. And the cell number will increase correspondently. We next assessed the effect of EA on proliferation of 3T3-L1 preadipocytes triggered by MDI. When 2-day postconfluent cells were incubated in a medium containing MDI, the cell number increased significantly during the clonal expansion phase as a function of time at least up to 4 days. In contrast, simultaneous EA (20 μM) treatment resulted in a greater than 50% decrease in cell number at day 4, compared with MDI alone. Accordingly, 3T3-L1 cell number increased from 2.02 × 10^6 to 6.07 × 10^6 cells/dish within 4 days of induction of adipogenesis in control medium, a 3-fold increase (n = 3; P < 0.05 compared to day 0), and this increase was inhibited by 69% (day 4; n = 3; P < 0.05) when 3T3-L1 preadipocytes were induced to differentiate in the presence of EA (n = 3; P < 0.05) (Figure 4(a)).

As for the cell number reduced, there are two possible situations leading to the results, cell apoptosis increasing, or cell cycle restraining. The percentage of apoptotic cells, assessed by PI and Annexin V-FITC double stainings, was minimal in all culture conditions, indicating that changes in apoptosis were not the explanation for the lower number of cells observed when adipogenesis was reduced in the presence of EA (Figure 4(b)).

For the cell cycle progression, PI staining was used and flow cytomtery analysis revealed that EA administration resulted in a significant reduction in the cell population in S phase (from 38.1 to 19%) with a concomitant increase in the cell population in G1 phase (from 58.71 to 70.9%) at 24 h after induction (Figures 4(c) and 4(d)). These data suggest that EA prevents MCE through inhibiting G1/S transition. Consistently, cell proliferation was significantly inhibited within 4 days after induction (G1/S transition takes place probably within 4 days after MDI [6]).

To confirm the effect of EA on refraining 3T3-L1 preadipocytes from entering the S phase, the effect EA on DNA synthesis was evaluated by using EdU labeling. As shown in Figure 4(e), EA administration caused less EdU incorporation. These data consolidate the antiproliferation property of EA in 3T3-L1 preadipocyte during MCE.

3.4. EA Inhibits Differentiation-Induced Cyclin A Protein Expression and Phosphorylation of Rb. The expression of key regulators of cell cycle progression is tightly governed during the clonal expansion phase of adipogenesis. Retinoblastoma protein (Rb) is a critical regulator of cell cycle progression at the G1/S transition [8]. Previous studies have shown that Rb and C/EBP family members are able to interact in in vitro experiments [8].

In the present study, Cyclin A protein levels changed dynamically during this time and were clearly upregulated at 16 h after adipogenic induction, from initially undetectable levels. The large increase in Cyclin A at these time points was strongly inhibited by 46.9% (16 h), 30.6% (20 h), and 38.7% (24 h), respectively (n = 3; P < 0.05; Figures 5(a) and 5(b)). Rb phosphorylation, assessed using a phospho-specific antibody, was strongly stimulated in 3T3-L1 preadipocytes 16 hrs after adipogenic induction. However, a 29.3%, 41.7%, and 33.4% inhibition in Rb phosphorylation was observed in cells exposed to EA and adipogenic inducers with 16 hrs, 20 hrs, and 24 hrs, respectively (n = 3; P < 0.05; Figures 5(c) and 5(d)). But our data fail to show that EA exerts any effect on cdk2 protein expression (data not shown). These results suggest that regulation of Rb downstream targets may be impaired by EA.

3.5. EA Inhibit Differentiation-Induced C/EBPα DNA-Binding Activity and Its following Protein Expression. C/EBPβ function might be altered, as it has been reported to be influenced by the state of Rb phosphorylation [21]. Upon the addition of adipogenic inducers under control conditions, DNA-binding activity of C/EBPβ was observed at 12, 24, and 36 hrs after induction (Figure 6(a)). Data shows that EA resulted in a trend of lower levels of DNA-binding activity of C/EBPβ, but this did not reach significance. Furthermore, quantitative assessment of DNA binding demonstrates that EA does inhibit C/EBPα DNA-binding activity compared to cells differentiated in the presence of control medium (n = 5; P < 0.05; Figure 6(b)). For the effect of EA on C/EBPα protein expression, data showed that EA can significantly inhibit its protein expression (n = 5; P < 0.05; Figures 6(c) and 6(d)).

4. Discussion

Obesity is characterized by increased adipose tissue mass that results from both increased fat cell number and increased fat cell size [2]. The amount of adipose tissue mass can be regulated by the inhibition of adipogenesis from fibroblastic preadipocyte to mature adipocyte. Blocking adipocyte differentiation is an important antiobesity strategy. Therefore,
Figure 3: EA was added at different day of differentiation to check time window of EA action. Data shows that the inhibitory effect on lipid accumulation was diminished when the addition of EA was delayed to day 2 of differentiation. (a) Morphology data from Oil Red O staining data show that EA inhibition of adipogenesis takes place predominantly within the first 2 days. (b) The first 2-day inhibitory effect exerted by EA was further confirmed by changes in the expression of specific adipocyte marker expression: PPARγ and aP2. β-actin loading as control. (c and d) Densitometric data are expressed of PPARγ and aP2 as mean ± SD of 3 independent experiments. *P < 0.05 compared to control (without EA treatment).

adipogenesis by modulation of adipocyte differentiation has been a target of antiobesity strategies [2]. The utilization of antiadipogenic compounds from natural sources could be helpful in reducing obesity and its consequent side effects [22].

3T3-L1 preadipocytes are a well-established cell model to investigate adipogenesis [5]. Confluent preadipocytes, growth arrest at the G0/G1 phase, differentiate to adipocytes through the one or two round(s) of clonal expansion and expression of adipocyte markers by hormonal induction. C/EBPβ is expressed early in the adipocyte differentiation program. After a long lag period (10–12 h) after adipogenic induction, C/EBPβ does acquire DNA-binding activity [6]. Acquisition of binding activity occurs as the cells synchronously reenter the cell cycle, traverse the G1/S checkpoint, and begin MCE. Later then expression of C/EBPα and
Figure 4: EA inhibits the cell number increase at early clonal expansion phase of adipogenesis. 3T3-L1 preadipocytes were induced to differentiate in the presence of control medium or EA for the indicated times. (a) On indicated days, cells were trypsinized and enumerated. (b) Cells were treated for 48 h at 37°C in 5 mL culture medium containing EA at final concentrations of 0 and 20 μM. Then cells were stained with PI and Annexin V-FITC and assessed by FACScan flow cytometer, the apoptotic rate of cell compared with control group (without EA). (c and d) DNA content was analyzed by PI staining and flow cytometry at given time points and analyzed. (e) At 16, 20, and 24 hrs after induction, cells were labeled with EdU for 2 hrs and then stained with Hoechst. The fluorescence of EdU (red) and Hoechst (blue) was detected with a fluorescence microscope. Results are the mean ± SD of 3 independent experiments, each performed in triplicate.
PPARγ was activated [6, 18]. Rb is a critical regulator of cell cycle progression at the G1/S transition. Previous studies have shown that Rb and C/EBP family members are able to interact in vitro experiments [7].

Fruits and nuts may prevent or reverse common human health conditions such as obesity, diabetes, and hypertension; together, these conditions are referred to as metabolic syndrome, an increasing problem [23]. Ellagic acid is one such polyphenol that is present in raspberries, strawberries, walnuts, and pomegranate. It attenuated diet-induced symptoms of metabolic syndrome with normalization of protein levels of Nrf2, NF-κB, and CPT1 [13]. Other than metabolic-regulating effect, EA has been reported to induce apoptosis in pancreatic and leukemia cancer cells [24, 25]. Several in vivo and in vitro studies confirmed its antiproliferative, anti-inflammatory, and antitumorigenic properties [26–28]. Aortic smooth muscle cell proliferation stimulated by ox-LDL was also inhibited by ellagic acid according to G0/G1 cell-cycle arrest [29]. Previous researchers have reported that ellagic acid was able to induce, in a dose-dependent manner, G0/G1 arrest and apoptotic cell death in human bladder cancer T24 cells, as determined by flow cytometric analysis of hypodiploid nuclei [30]. Panchal et al. found that EA induces fat redistribution in rats of high fat diet-induced symptoms of metabolic syndrome [13]. However, its biological and pharmacological effects on adipogenesis are still poorly defined. And data are still lacking to depict a direct correlation between EA and early differentiation of preadipocytes.

Data shown in present study demonstrated that, with adipogenic induction, 3T3-L1 preadipocytes were exposed to EA and were reinfused every two days. Our Oil Red O staining data revealed that EA lowered the neutral lipid content in 3T3-L1 preadipocyte with concentration-dependent manner. The expression of specific adipogenic marker, PPAR and αP2, confirmed this result. Using the Trypan blue dye exclusion test, we confirmed that the inhibitory effect of EA on adipogenesis was independent of potential cytotoxicity. Our data are consistent with the results shown by Losso et al. that EA at concentration in the range 10–100 μM/L did not affect the viability of normal fibroblast cells during a 24-hour incubation, where EA manifests strong antiproliferative activity against the colon, breast, and prostatic cancer cell lines [31]. Ellagic acid is a phenolic compound found in fruits including grape juice (10.2mg/100g), grape wine (5.6mg/100g), blueberries (0.9mg/100g), blackberries (42.4mg/100g), raspberries (17.9mg/100g), and strawberries (19.8mg/100g) [32]. The typical dietary intake of ellagic acid in humans is approximately 40–80 mg/d if 200 g strawberries or blackberries are eaten [32]. Previous studies also indicated that other polyphenols, such as resveratrol (20–100 μM) also inhibited adipogenesis in 3T3-L1 preadipocytes [33, 34]. In the present study, we found that 20 μM-EA treatment profoundly reduced adipogenesis in 3T3-L1 preadipocytes.
This dose (20 μM) of ellagic acid is equivalent to the dietary intake of approximately 80 g of berries subject to absorption [29].

During the normal course of adipogenesis, 3T3-L1 preadipocytes run through two distinct phases, early clonal expansion and terminal adipogenic differentiation. In order to check the time window of EA inhibition on 3T3-L1 preadipocytes adipogenesis, EA was administered during different periods of adipogenesis. The results revealed that EA is required at the onset of differentiation for maximal inhibition of adipogenesis. Enumeration test shows that EA can significantly reduce the cell number within 2–4 days after induction. It indicated that EA overtly inhibits the early mitotic clonal expansion, which is required for terminal differentiation. Our FACS data on cell-cycle analysis supported the blocking effect of EA on MDI-induced cell-cycle progression and preadipocytes proliferation. It revealed that EA blocked the cell cycle at the G1/S transition and caused cells to remain in the preadipocyte state. Our data were consistent with the work done by Yarmo et al.. The results there revealed that clonal expansion, an early event required for 3T3-L1 adipogenesis, was reduced in the presence of medium conditioned by murine J774 macrophages (MacCM). And then the central lipid accumulation was reduced in 3T3-L1 adipocytes [15].

Propidium iodide and Annexin V-FITC double staining shows no alteration of apoptotic effect of EA on 3T3-L1 preadipocyte, which indicated that the inhibitory effect of EA on cell number reduction and preadipocyte proliferation was independent of potential proapoptosis of EA. The results are
similar to the previous data that EA-mediated G0/G1 cell-cycle arrest in Aortic smooth muscle cell and cancer cell to induce antiproliferation [29, 30], but without effect on 3T3-L1 adipocytes apoptosis.

CCAAT element-binding protein β/δ plays an important role in the clonal expansion of preadipocytes through facilitating PPARγ and C/EBPα expression by their interactions with the DNA element(s) located in the genes sequence. The retinoblastoma protein (Rb) is a nuclear protein that regulates proliferation, differentiation, and apoptosis. As the cells synchronously transition through the G1/S checkpoint (12–16 h), formation of active Cyclin A/cdk2 complexes fully phosphorylate Rb, leading to the release of E2F1 and initiating the limited proliferation that defines this early required phase of adipogenesis [6, 35]. The upregulation of Cyclin A expression is a critical component of the 3T3-L1 preadipocyte MCE [6]. Rb can be phosphorylated by cdk5, which are activated upon association with cyclins [36]. The data in the present study showed that EA can reduce the Cyclin A protein expression and Rb phosphorylation significantly but not completely. However, EA fails to exert any effect on cdk2 protein expression, which does not influence the role of EA on Rb phosphorylation. Vanella et al. also reported that EA treatment represents a new approach and highly effective strategy in reducing carcinogenesis by decreasing the level of Rb phosphorylation [12]. Rb is also a critical regulator of C/EBPβ DNA-binding activity during the early stages of 3T3-L1 adipogenesis [8]. Here, EA has the trend to inhibit C/EBPβ DNA-binding activity but not to reach significant differentiation, assessed by centromeric localization and ELISA-based specific DNA binding. It does raise the possibility that transcriptional activity of C/EBPβ in our model could potentially still be affected.

However, EA triggered decrease of C/EBPα DNA binding activity at 36 and 48 hrs following MDI induction, probably through inhibition of C/EBPα protein expression. The data here did show that EA inhibits the following C/EBPα protein expression. C/EBPβ, which is coexpressed with C/EBPβ to initiate the expression of C/EBPα and PPARγ, [6, 37], may also be the candidate for EA to impact the C/EBPα DNA-binding activity and numerous adipocyte gene expression, which need further investigation.

In summary, this report suggests that EA inhibits 3T3-L1 preadipocyte adipogenesis. The mechanisms by which EA regulates adipogenesis possibly include the inhibition of mitotic clonal expansion by reducing Cyclin A protein expression and Rb phosphorylation to block 3T3-L1 preadipocyte entering into S phase at the early stage of adipocyte differentiation. Also, the expression of the adipogenic transcription factors such as C/EBPα and PPARγ was involved. Therefore, these results may be of help in understanding the basic biology of the antiadipogenic process, and developing the potent antiadipogenic modulators that are crucial to reducing obesity.

**Authors’ Contribution**

Xinmin Mao and Jun Ren contributed equally to the work.

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