

Anti-inflammatory and antiaging properties of chlorogenic acid on UV-induced fibroblast cell

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Background: Skin aging is the most common dermatological problem caused by intrinsic and extrinsic factor, such as exposure to (ultraviolet) UV rays. Chlorogenic acid (CA) is a phenolic compound which is known for its antioxidant properties against oxidative stress. **Objectives:** This study investigates the antiaging and anti-inflammatory properties of CA on UV-induced skin fibroblast cells. **Methods:** Anti-inflammatory properties of CA were assessed by measuring inflammatory-related proteins IL-1 β and TNF- α , while antiaging properties of CA were assessed by measuring reactive oxygen species (ROS), apoptosis, live and necrotic cells, and COL-3 gene expression level. **Results:** Treating UV-induced skin fibroblast cells with CA decreased the level of ROS, IL-1 β , TNF- α , apoptotic cells, and necrotic cells and increased live cells and COL-3 gene expression. **Conclusion:** CA has the potential as the protective compound against inflammation and aging by decreasing the level ROS, pro-inflammatory cytokines IL-1 β and TNF- α , apoptotic cells, and necrotic cells and by increasing live cells and COL-3 gene expression.

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Running Title: Anti-inflammatory and Antiaging of Chlorogenic Acid

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Abstract

Background: Skin aging is the most common dermatological problem caused by intrinsic and extrinsic factor, such as exposure to ultraviolet (UV) rays. Chlorogenic acid (CA) is a phenolic compound which is known for its antioxidant properties against oxidative stress.

Objectives: This study investigates the antiaging and anti-inflammatory properties of CA on UV-induced skin fibroblast cells.

Methods: Anti-inflammatory properties of CA were assessed by measuring inflammatory-related proteins IL-1 β and TNF- α , while antiaging properties of CA were assessed by measuring reactive oxygen species (ROS), apoptosis, live and necrotic cells, and COL-3 gene expression level.

Results: Treating UV-induced skin fibroblast cells with CA decreased the level of ROS, IL-1 β , TNF- α , apoptotic cells, and necrotic cells and increased live cells and COL-3 gene expression.

Conclusion: CA has the potential as the protective compound against inflammation and aging by decreasing the level ROS, pro-inflammatory cytokines IL-1 β and TNF- α , apoptotic cells, and necrotic cells and by increasing live cells and COL-3 gene expression.

Keywords: Antiaging, anti-inflammatory, apoptosis, chlorogenic acid, ultraviolet

Introduction

Skin aging is the most common dermatological problem, especially for women. While aging could happen as a result of intrinsic mechanisms, the external causes also play a big role in causing skin aging (Bauman, 2007; Farage, 2008). The external causes in aging, mostly from environmental factor i.e., sun (ultraviolet), infrared and heat, smoking, air pollution, and low

antioxidant diet (Vierkötter & Krutmann, 2012, Addor, 2018). Ultraviolet (UV) exposure is one of the factors that play an important role in affecting all body systems, especially for the skin as the outer barrier against the external environment. Direct exposure to UV rays could cause progressive loss of integrity and physiology manifested in visible rough, sagging, and wrinkles in the skin (Bauman, 2007). Infrared (IR) and heat induce premature skin aging with the mechanism: (1) IR stimulates the expression of Matrix Metalloproteinase-1 (MMP-1) and decreases type 1 procollagen expression *in vivo*. Acute IR irradiation also increases new, leaky vessel formation, and induces inflammatory cellular infiltration. (2) Heat energy increase skin temperature which increases MMP-1, -3, and -12 expression, and modulates elastin and fibrillin synthesis, resulting in the development of solar elastosis. Acute heat shock in human skin stimulates new vessel formation, recruit inflammatory cells, and causes oxidative DNA damage (Cho et al., 2009). Smoking induce human skin aging by the additive induction of MMP-1 expression (Yin et al., 2001). Air pollution can induce skin aging by generating Reactive Oxygen Species (ROS) by the particulate matter contained within the pollution (Donaldson et al., 2005). Low antioxidant diet correlate with the antioxidant that is micronutrient in various vegetable to prevent the skin aging. If people didn't consume enough antioxidant the case mostly there are wrinkles on the skin (Addor, 2018).

At the cellular level, UV exposure affects the production of free radicals known as ROS. An imbalance in ROS production causes oxidative stress, which leads to cellular or tissue changes (Davalli et al., 2016). While ROS was naturally formed by cells as a normal condition, UV exposure is known to increase the intracellular ROS level, causing oxidative stress. Oxidative stress leads to the activation of a series of signaling pathways to the apoptosis of skin fibroblast (Tanigawa, 2014). ROS production is also closely related with inflammation. In contrast to young individuals, aged individuals have more numbers of inflammatory cytokines, such as IL-6, IL-1 β , and TNF- α (Sanada et al., 2018). The existence of these substances plays an important role in the progression of inflammatory diseases. The inhibition of oxidative stress now becomes an important approach for treating skin damage due to aging (Rinnerthaler et al., 2015, Liguori et al., 2018).

Chlorogenic acid (CA) is one of the phenolic compounds from the hydroxycinnamic family, which is known for its antioxidant properties against free radicals (Santana-Gálvez et al., 2017). This compound is also found in beverages prepared from herbs, fruits, and vegetables (Clifford, 1999). *In vitro* and *in vivo* analyses reported the roles of CA in mitigating oxidative and inflammatory stresses (Liang & Kitts, 2015).

CA is naturally found in snake fruit peel (Girsang et al., 2019a; Fitri, 2015). Recent *in silico* studies about snake fruit peel compound i.e. CA reported its antiaging and antioxidant properties (Girsang et al., 2019b). CA is also reported to have a protective effect by suppressing ROS level and to have anti-inflammatory properties against to lead (Pb) poisoning (Girsang et al., 2019c). Therefore, this study aimed to determine the antiaging and anti-inflammatory properties of CA on UV-induced fibroblast cells, being reported for its antioxidant activities.

Materials and Methods

Materials

Skin fibroblast cells (BJ cell line, ATCC no CRL-2522) were obtained from Biomolecular and Biomedical Research Center, Aretha Medika Utama, Bandung, Indonesia. Cells were grown in MEM (Biowest, L0416-500) with 10% fetal bovine serum (FBS, Biowest, S1810-500), 1% antibiotic/antimycotic (ABAM, Biowest, L0010-100), 1% Nanomycopolitine (Biowest, L-X16-100), 1% amphotericin B (Biowest, L0009-050), and 0.1% gentamicin (Gibco 15750060) respectively. Cells were incubated at 37°C in a humidified atmosphere with 5% CO₂. The cells with 80% confluency (n = 10⁶ in six-well plate) were incubated with UV exposure for 75 min (37°C, 5% CO₂). The cells induced from UV exposure were treated for 4 days with CA (Chengdu Biopurify Phytochemical, BP0345) and were harvested with 0.25% trypsin-EDTA (Gibco, 25200072) after 4 days treatment (Widowati et al., 2016; Girsang et al., 2019c).

Measurement of IL-1β and TNF-α

The conditioned medium of the treated cells was used to measure the concentration of IL-1β and TNF-α. The measurement was conducted according to Human IL-1β ELISA Kit (BioLegend, Elisa Max Deluxe, 437004) and Human TNF-α ELISA Kit (BioLegend, Elisa Max Deluxe, 430204) protocol. The treatments used in this experiments were as follows: (1) normal control; (2) BJ cells + DMSO 1% (vehicle control); (3) UV-induced BJ cells (positive control); (4) UV-induced BJ cells + CA 6.25 μg/ml; and (5) UV-induced BJ cells + CA 25 μg/ml (Laksmitawati et al., 2017; Widowati et al., 2016; Widowati et al., 2019a; Lister et al., 2020).

Total Protein Measurements

The total protein measurement was conducted to assess the IL-1β and TNF-α level by mg protein. The BSA stock was obtained by dissolving 2 mg of BSA in 1000 μL ddH₂O (Sigma, A9576, Lot. SLB2412). The BSA solution was obtained from diluting the BSA stock. The standard solution with 20 μL in number and 200 μL of Quick Start Dye Reagent 1× (Biorad, 5000205) were added to each well. The plate was incubated for 5 min at room temperature. The result was read at 595 nm (Widowati et al., 2019a; Lister et al., 2020).

Measurement of ROS Level

The measurement of ROS levels was assessed using DCFDA Cellular Reactive Oxygen Species Detection Assay Kit (Ab113851) according to the protocol, with slight modification by flow cytometry (MACSQuant Analyzer 10, Miltenyi). The cells were harvested, counted, and added with 500 μl DCFDA working buffer. They were then stained with 25 μM DCFDA and incubated for 30 min in 37°C, with 5% CO₂. The result was analyzed using MACSQuant Analyzer 10 Flow Cytometer (Miltenyi) (Widowati et al., 2014; Niocel et al., 2016; Girsang et al., 2019c; Lister et al., 2020).

Measurement of Apoptotic, Live, Necrotic Cells

The measurement of apoptotic, live, and necrotic cells was assessed by flow cytometry analysis. The cells were placed in a 2 × 6-well plate (n = 500.000) and were incubated for 2 h. The growth medium was discarded in 4 days and harvested and centrifuged for 5 min at 1600 rpm. The

pellet was added with 500 μ L FACS buffer and centrifuged. The pellet was added with 100 μ L FACS buffer and stained by annexin and propidium iodide (PI). The stained cells were incubated for 1 h and analyzed using flow cytometry (Lister et al., 2020; Widowati et al., 2019b).

Measurement of COL-3 Gene Expression

Measurement of COL-3 gene expression was performed using real-time quantitative reverse transcription polymerase chain reaction (qRT-PCR) (Thermo Fisher Scientific PikoReal 96). The reaction mix was set for 40 \times reaction consisting of 8 μ L nuclease-free water, 20 μ L SsoFast EvaGreen supermix, 8 μ L mix primer, and 4 μ L cDNA. The reaction mix was then run with the cycle for pre-denaturation (95°C for 5 min), denaturation (95°C for 1 min), annealing (58°C for 40 s), pre-elongation (72°C for 1 min), and elongation (72°C for 5 min). All the reaction was set for 55–90°C melting curve and 4°C infinite hold, in a total of 40 reaction cycles (Widowati et al., 2019b; Afifah et al., 2019).

Statistical Analysis

All the data are presented as the mean \pm standard deviation. The data were analyzed using Shapiro–Wilk test followed by Mann–Whitney–Wilcoxon test and independent t-test. P-values < 0.05 were considered significant.

Results

IL-1 β Level

The conditioned medium of the treated fibroblast cell with CA was measured and compared to the normal control and UV-induced fibroblast cell (positive control). Treatment with DMSO 1% (vehicle control) was also included to compare with the normal control. The result shows a significant decreasing level of IL-1 β compared to the positive control (**Fig. 1a**). Total protein assay was present to measure the IL-1 β level by mg protein. The result showed a significant decrease in the IL-1 β level compared to the positive control (**Fig. 1b**).

TNF- α Level

Measurement of tumor necrosis factor- α (TNF- α) was performed using Human TNF- α ELISA Kit (BioLegend, Elisa Max Deluxe cat no. 430204). The result was compared to the normal and positive controls. Treatment with 2.5 μ g/ml CA shows a significant difference with the normal control, and both treatments show a significant difference of TNF- α level compared with the negative control (**Fig. 1c**). Total protein assay was present to measure the TNF- α level by mg protein. The result showed a significant decrease in the TNF- α level compared to both controls (**Fig. 1d**).

ROS Level

The level of reactive oxygen species (ROS) of UV-induced fibroblast cells was measured using flow cytometry with DCFDA single staining. The dot blots show the population of the analyzed cells, and the peak shows the positive ROS cells. The normal control, vehicle control,

positive control, treatment with CA 6.25 µg/ml, and treatment with CA 25 µg/ml show 0.02%, 0.05%, 16.01%, 4.95%, and 6.86% positive ROS levels, respectively (**Fig. 2**). Both cells treated with CA show a significant result related to controls (**Fig. 3**).

Apoptosis Level

The apoptosis level was analyzed using flow cytometry. The surface markers PI and annexin were used as stains. **Figure 4** shows the representative of dot blots effect CA towards apoptosis cells. Flow cytometry analysis reveals the percentage of live cells, necrotic cells, early apoptosis, and late apoptosis of cells. **Figure 5** shows the comparison between each treatment on live of cells (**a**), necrosis of cells (**b**), death of cells (early apoptosis) (**c**), and apoptosis (late apoptosis) (**d**).

COL-3 Gene Expression

The COL-3 gene expression was examined using real-time quantitative reverse transcription polymerase chain reaction (qRT-PCR). **Table 1** presents the primer and annealing temperature used for performing COL-3 gene expression, while **Table 2** shows the concentration and RNA purity. The COL-3 gene expression shown in **Figure 6** revealed a significant difference among treatments.

Discussion

UV-induced aging mechanism triggered free radicals in the form of increased intracellular ROS level, leading to oxidative stress, which then triggered inflammation leading to cell death (apoptosis). BJ fibroblast cells are the model of aging cells induced by UV (Girsang et al., 2019c). Fibroblast cells play a role in tissue granulation and scar formation during the inflammation process. The result of the anti-inflammatory test shows treating UV-induced BJ cells with CA decreased the IL-1β and TNF-α levels related to the aging model (**Fig. 1**). The inflammation triggered by UV rays resulted in the increase of the pro-inflammatory cytokine group IL-1β and TNF-α, which are used as inflammatory markers for anti-inflammatory assessment (Chung et al., 2009).

The presence of ROS as one of the inflammation trigger also being one of the direct UV-exposures effect in cells (Tanigawa, 2014). The accumulation of ROS leads to cell death known as apoptosis, which results in the decreasing level of healthy cells. The result of the assessment with fluorescence intensity as an indicator by DCFDA staining showed that CA decreased the level of ROS compared with the positive control (**Fig. 2 and Fig. 3**). The previous study about CA also stating the reducing ROS level induced by lead (Pb) and hydrogen peroxide effect by this substances (Girsang et al., 2019c; Hoelzl et al., 2010), CA has antiaging property by inhibiting enzyme related skin aging i.e. MMP-1 and human skin fibroblast elastase (HSFE) according *in silico* analysis (Girsang et al., 2019b).

UV exposure of BJ-cells also affects the level of apoptosis. The oxidative stress caused by UV rays promotes apoptosis in healthy cells (Kanan & Jain, 2000). Flow cytometry analysis revealed the lowest number of live cells in UV-induced BJ cells. Treatment with CA on UV-exposed cells resulted in increased number of live cells and decreased cell death, necrosis, and

apoptosis (**Fig. 4 and Fig. 5**). The protective effect of CA toward apoptosis is also reported in rat liver with oxidative stress triggered by methotrexate (Ali et al., 2017).

While CA could potentially decrease the levels of ROS, IL-1 β , and TNF- α , the measurement of COL-3 gene expression can also be conducted to determine the repairing effect of CA toward aging. COL-3 gene is related to the pattern of tendon fibril thickness and is more abundant in younger individuals than aged individuals of the animal model (Ribitsch et al., 2020). UV exposure could also decrease the COL-3 gene expression level (Chen et al., 2019). The result of this research showed the effect of UV exposure and the treatment of CA toward COL-3 gene expression level. Treatment with CA increased the COL-3 gene expression level compared to the aging model (**Fig. 6**). According to these results, we proposed the mechanism of anti-inflammatory and antiaging properties of CA against UV-induced BJ cells (**Fig. 7**).

Conclusion

This study shows that CA has the potential as the protective compound against inflammation and aging by reducing ROS, pro-inflammatory cytokines IL-1 β and TNF- α , apoptosis, and necrotic cells and by increasing live cells and COL-3 gene expression.

Acknowledgements

We are gratefully acknowledging Universitas Prima Indonesia, Medan, North Sumatera, Indonesia for the financial support. This research also funded, facilitated and supported by Aretha Medika Utama, Biomolecular and Biomedical Research Center, Bandung, West Java, Indonesia. We also thank to Hanna Sari Widya Kusuma, Cintani Dewi Wahyuni from Aretha Medika Utama Biomolecular and Biomedical Research Center for the their valuable assistance.

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Table and Figure Legends

Table 1. RT-qPCR detail of COL-3 gene

Table 2. Concentration and purity of RNA

Figure 1. Effect CA towards IL-1 β and TNF- α level on UV-induced fibroblast cells

(a) : IL-1 β level (pg/ml), (b) : IL-1 β level (pg/mg protein)

(I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) UV-induced BJ cells (positive control), (IV) UV-induced BJ cells +CA 6.25 μ g/ml, (V) UV-induced BJ cells+CA 25 μ g/ml.

Data is presented as mean \pm standard deviation. Single asterisk symbol () marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

(c) : TNF- α level (pg/ml), (d) : TNF- α level (pg/mg protein)

(I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) UV-induced BJ cells (positive control), (IV) UV-induced BJ cells +CA 6.25 μ g/ml, (V) UV-induced BJ cells+CA 25 μ g/ml.

Data is presented as mean \pm standard deviation. Single asterisk symbol () marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

Figure 2. The representative of dot blots of various concentration CA on UV-induced fibroblast cell toward ROS level by flow cytometry

(a) : BJ cells (negative control); 0.02%, (b) : BJ cells+DMSO 1% (vehicle control) : 0.05%, (c) : BJ cells+UV (positive control) : 16.01%,

(d) : UV-induced BJ cells+CA 6.25 μ g/ml: 4.95%, (e) : UV-induced BJ cells +CA 25 μ g/ml: 6.86%.

Figure 3. Effect CA towards ROS level on UV-induced fibroblast cells

(I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) UV-induced BJ cells (positive control), (IV) UV-induced BJ cells +CA 6.25 μ g/ml, (V) UV-induced BJ cells+CA 25 μ g/ml.

Data is presented as mean \pm standard deviation. The single asterisk symbol () marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test

Figure 4. The representative of dot blots of various concentration CA on UV-induced skin fibroblast cell toward apoptosis level by flow cytometry

- (a) BJ cells (negative control): live cells: 100%, necrotic: 0.0%, early apoptosis: 0.0%, late apoptosis: 0.0%
- (b) BJ cells+DMSO 1% (vehicle control) : live cells: 95.14%, necrotic: 0.38%, early apoptosis: 7.86%, late apoptosis: 2.61%
- (c) BJ cells+UV (positive control): live cells: 38.60%, necrotic: 18.0%, early apoptosis: 16.98%, late apoptosis: 26.43%
- (d) UV-induced BJ cells+CA 6.25 µg/ml: live cells: 80.35%, necrotic: 0.78%, early apoptosis: 10.48%, late apoptosis: 8.39%
- (e) UV-induced BJ cells+CA 25 µg/ml: live cells: 85.50%, necrotic: 0.92%, early apoptosis: 6.51%, late apoptosis: 6.97%

Figure 5. Effect CA towards apoptosis cells on UV-induced fibroblast cells

- (a) : live of cells, (b) : necrosis of cells, (c) : death of cells (late apoptosis of cells), (d): apoptosis of cells (early apoptosis of cells)
- (I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) BJ cells+UV (positive control), (IV) UV-induced BJ cells+CA cells +CA 6.25 µg/ml, (V) UV-induced BJ cells+CA 25 µg/ml.
- *Data is presented as mean ± standard deviation. The single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

Figure 6. Effect CA towards COL-3 gene expression on UV-induced fibroblast cells

- (I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) BJ cells+UV (positive control), (IV) UV-induced BJ cells+CA cells +CA 6.25 µg/ml, (V) UV-induced BJ cells+CA 25 µg/ml.
- *Data is presented as mean ± standard deviation. The single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

Figure 7. Proposed mechanism of CA protective effect towards UV-induced skin fibroblast cells.

UV exposures causing an increasing level of ROS leading to an increase of pro-inflammatory cytokines IL-1 β and TNF- α leading to inflammation. The presence of ROS triggered the production of caspase 9 (Casp-9) along with the production of Casp-8 by the presence of TNF- α leading to apoptosis. These two condition of inflammation and apoptosis leading to aging in cells. On another side, UV-exposures decreasing the COL-3 gene expression causing the loss of fibril pattern leading to aging as well. The presence of CA inhibits the production of ROS level thus suppressing the pro-inflammatory cytokines IL-1 β and TNF- α . The presence of CA also increasing the level of COL-3 gene expression.

Table 1 (on next page)

RT-qPCR detail of COL-3 gene

1

Gene Symbol s	Primer Sequence (5' to 3')		Product Size (bp)	Annealing (°C)	Cycle	References
	Upper strand : sense	Lower strand : antisense				
COL-3	5'- CCAGGAGCTAACGGTCTCAG -3'	5'- CACGGTTTCCATCTCTTCCA -3'	103	54	40	Ye et al., 2012
β-Actin	5'-AGACCTGTACGCCAACACAG-3'	5'-TTCTGCATCCTGTCGGCAAT-3'	24	60	40	Palumbo et al., 2018

2

Table 2(on next page)

Concentration and purity of RNA

1

2

	RNA concentration (ng/μl)	RNA purity (λ 260/ λ 280 nm)
BJ cells (negative control)	190.2	2.3468
BJ cells+DMSO 1% (vehicle control)	179.2	2.3271
UV-induced BJ cells (positive control)	177.70	2.3179
UV-induced BJ cells + CA 6.25 μg/ml	118.80	2.2235
UV-induced BJ cells+CA 25 μg/ml	105.64	2.2848

3

Figure 1

Effect CA towards IL-1 β and TNF- α level on UV-induced fibroblast cells

(a) : IL-1 β level (pg/ml), (b) : IL-1 β level (pg/mg protein) (I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) UV-induced BJ cells (positive control), (IV) UV-induced BJ cells +CA 6.25 μ g/ml, (V) UV-induced BJ cells+CA 25 μ g/ml. *Data is presented as mean \pm standard deviation. Single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test. (c) : TNF- α level (pg/ml), (d) : TNF- α level (pg/mg protein) (I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) UV-induced BJ cells (positive control), (IV) UV-induced BJ cells +CA 6.25 μ g/ml, (V) UV-induced BJ cells+CA 25 μ g/ml. *Data is presented as mean \pm standard deviation. Single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

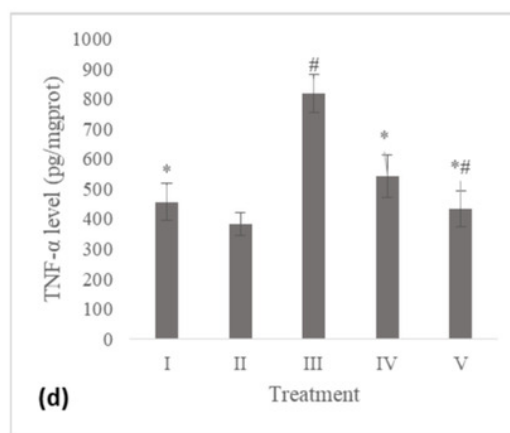
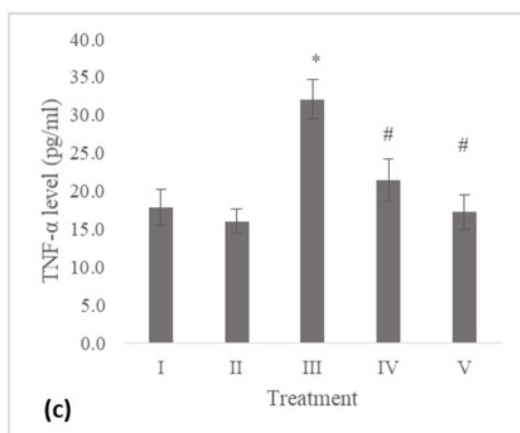
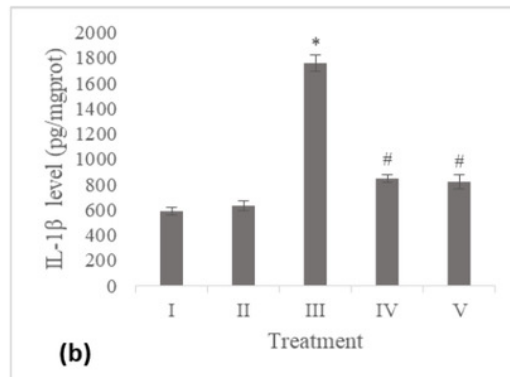
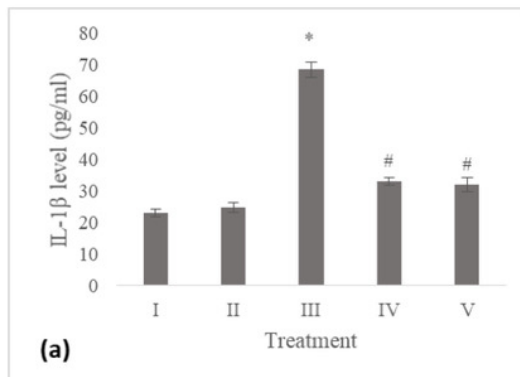


Figure 2

The representative of dot blots of various concentration CA on UV-induced fibroblast cell toward ROS level by flow cytometry

(a) : BJ cells (negative control); 0.02%, (b) : BJ cells+DMSO 1% (vehicle control) : 0.05%, (c) : BJ cells+UV (positive control) : 16.01%, (d) : UV-induced BJ cells+CA 6.25 μ g/ml: 4.95%, (e) : UV-induced BJ cells +CA 25 μ g/ml: 6.86%.

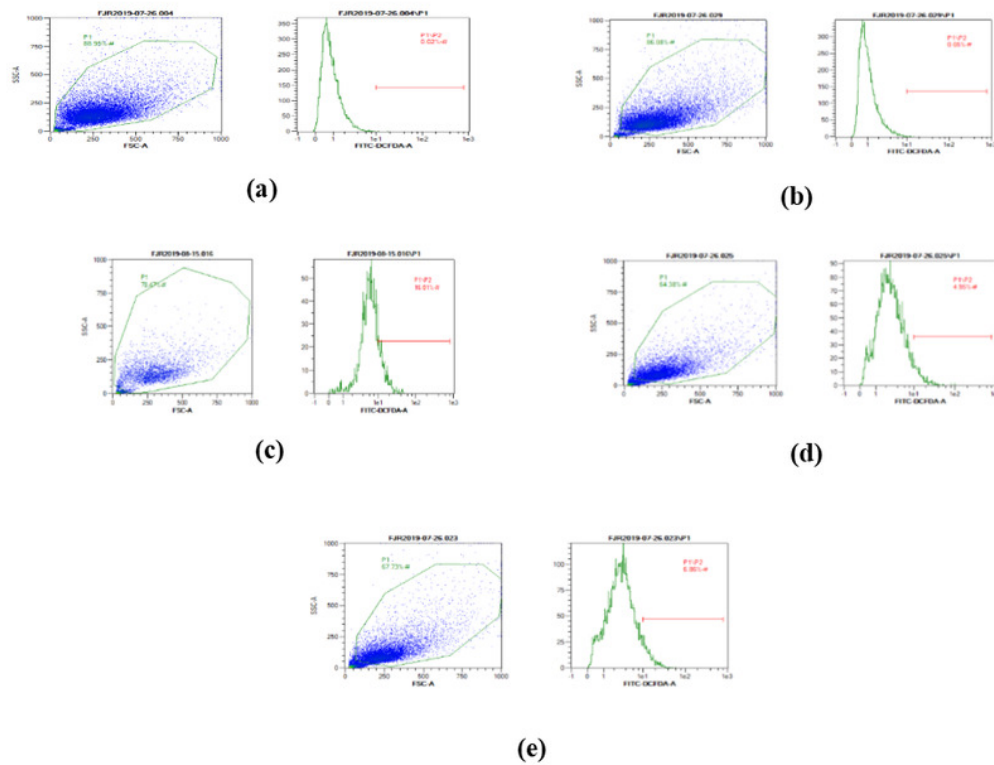


Figure 3

Effect CA towards ROS level on UV-induced fibroblast cells

(I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) UV-induced BJ cells (positive control), (IV) UV-induced BJ cells +CA 6.25 µg/ml, (V) UV-induced BJ cells+CA 25 µg/ml. *Data is presented as mean ± standard deviation. Single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test

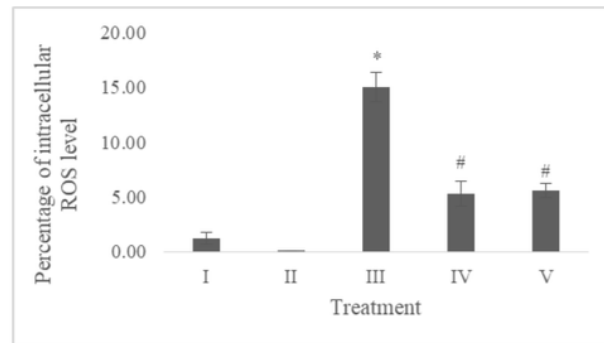


Figure 4

The representative of dot blots of various concentration CA on UV-induced skin fibroblast cell toward apoptosis level by flow cytometry

(a) BJ cells (negative control): live cells: 100%, necrotic: 0.0%, early apoptosis: 0.0%, late apoptosis: 0.0% (b) BJ cells+DMSO 1% (vehicle control) : live cells: 95.14%, necrotic: 0.38%, early apoptosis: 7.86%, late apoptosis: 2.61% (c) BJ cells+UV (positive control): live cells: 38.60%, necrotic: 18.0%, early apoptosis: 16.98%, late apoptosis: 26.43% (d) UV-induced BJ cells+CA 6.25 µg/ml: live cells: 80.35%, necrotic: 0.78%, early apoptosis: 10.48%, late apoptosis: 8.39% (e) UV-induced BJ cells+CA 25 µg/ml: live cells: 85.50%, necrotic: 0.92%, early apoptosis: 6.51%, late apoptosis: 6.97%

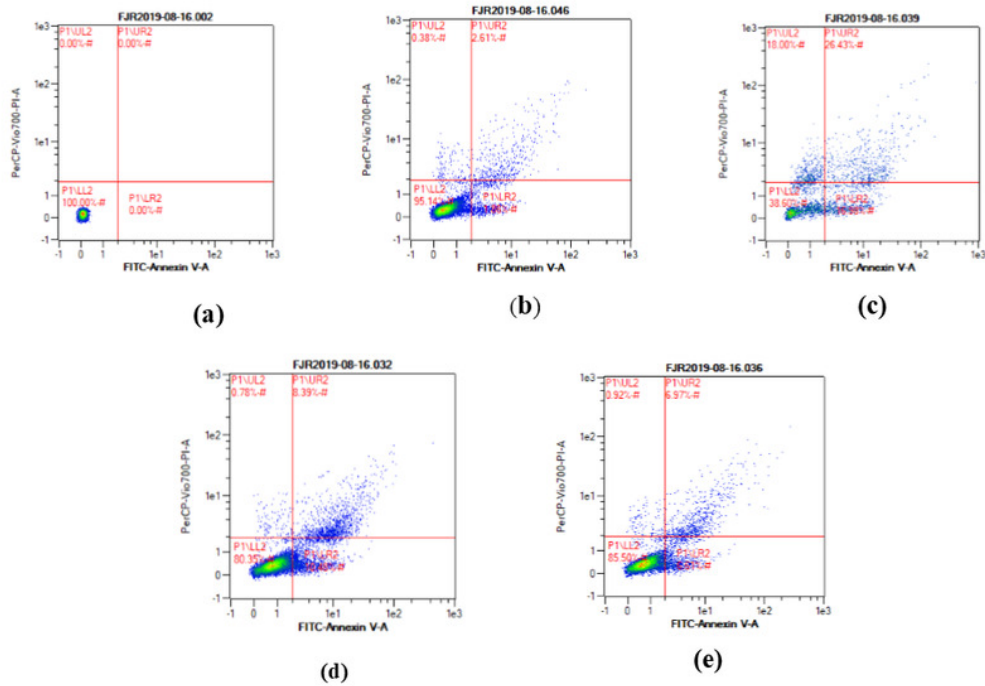


Figure 5

Effect CA towards apoptosis cells on UV-induced fibroblast cells

(a): live of cells, (b) : necrosis of cells, (c) : death of cells (late apoptosis of cells), (d): apoptosis of cells (early apoptosis of cells) (I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) BJ cells+UV (positive control), (IV) UV-induced BJ cells+CA cells +CA 6.25 µg/ml, (V) UV-induced BJ cells+CA 25 µg/ml. *Data is presented as mean \pm standard deviation. The single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

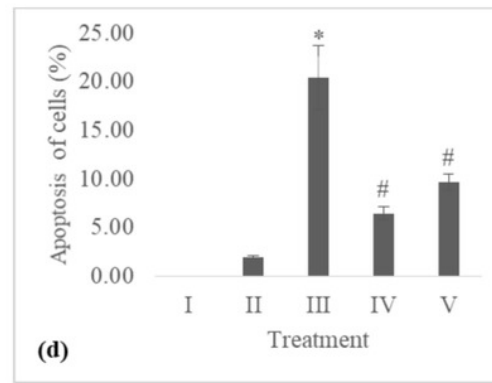
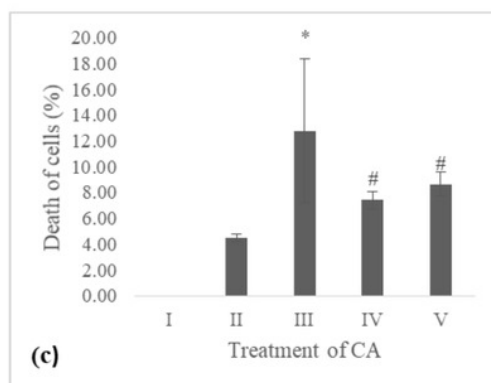
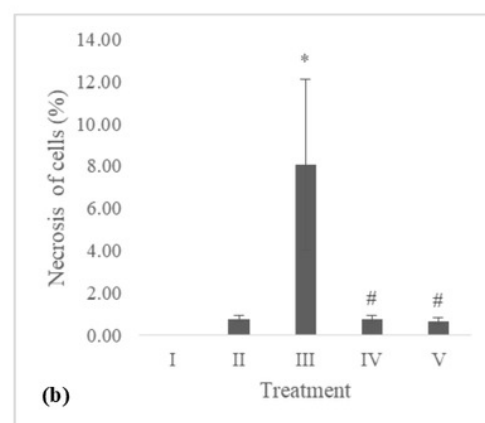
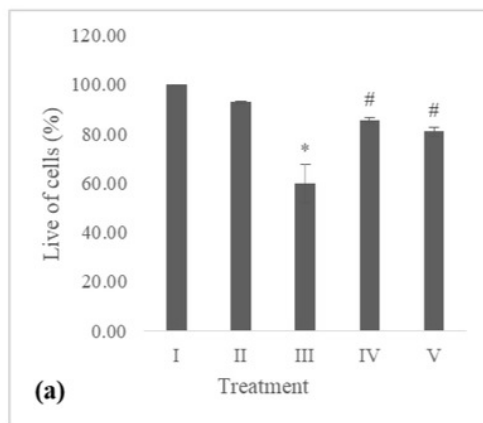


Figure 6

Effect CA towards COL-3 gene expression on UV-induced fibroblast cells

(I) BJ cells (negative control), (II) BJ cells+DMSO 1% (vehicle control), (III) BJ cells+UV (positive control), (IV) UV-induced BJ cells+CA cells +CA 6.25 µg/ml, (V) UV-induced BJ cells+CA 25 µg/ml. *Data is presented as mean ± standard deviation. The single asterisk symbol (*) marks statistical difference compared to negative control group at 0.05 significance level, single hashtag (#) marks statistical difference compared positive control at 0.05 significance level based on Tukey HSD post hoc test.

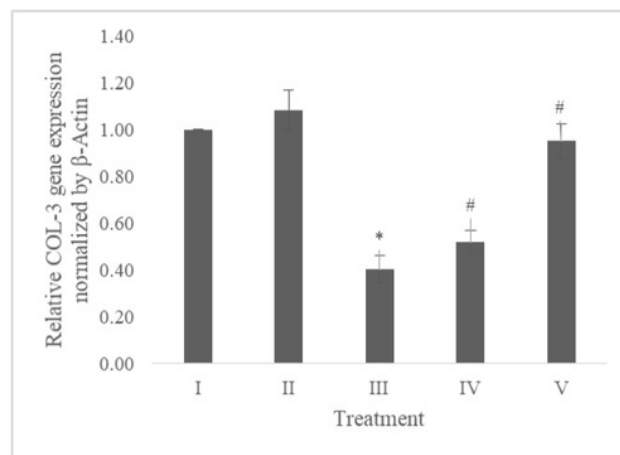


Figure 7

Proposed mechanism of CA protective effect towards UV-induced skin fibroblast cells.

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