

Antiglycation and antioxidant activities of mogroside extract from *Siraitia grosvenorii* (Swingle) fruits

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Abstract *Siraitia grosvenorii* (Swingle) is one kind of medical and edible plants with various health-promoting properties. Recently, its hypoglycemic and antidiabetic activities have been reported, but the underlying mechanism remains to be explored. The current study was aimed to investigate the antioxidant and antiglycation activities of mogroside extract (MGE) from *Siraitia grosvenorii* (Swingle). The results showed that compared to glycated BSA, MGE at middle (125 µg/mL) and high dose (500 µg/mL) significantly inhibited BSA glycation evidenced by decreased fluorescent AGEs formation, protein carbonyls and N^ε-(carboxymethyl) lysine (CML) level at 500 µg/mL by 58.5, 26.7 and 71.2%, respectively. Additionally, the antiglycative activity of MGE (500 µg/mL) was comparable to aminoguanidine (AG) at the equal concentration. However, the inhibitory effect of MGE on glycation-induced increase of fructosamine level and decrease of thiol level was not remarkable. MGE was a potent peroxide

radicals scavenger (851.8 µmol TE/g), moderate DPPH and ABTS radicals scavenger with IC₅₀ 1118.1 and 1473.2 µg/mL, respectively, corresponding to positive controls ascorbic acid of IC₅₀ 9.6 µg/mL, and trolox of IC₅₀ 47.9 µg/mL, respectively, and mild reducing power. These findings suggest that MGE may serve as a new promising antiglycative agent against diabetic complications by inhibiting protein glycation and glycooxidation.

Keywords *Siraitia grosvenorii* (Swingle) · Cucurbitane triterpene glycosides · Antiglycation · Antioxidant

Abbreviations

1-DMF	1-Deoxy-1-morpholino-fructose
3-DG	3-Deoxyglucosone
AAPH	2,2'-Azobis (2-amidinopropane) dihydrochloride
ABTS	2,2'-Azinobis-(3-ethylbenzothiazoline-6-sulfonic acid)
AG	Aminoguanidine
AGEs	Advanced glycation end products
ANOVA	Analysis of variance
AUC	Area under curve
BSA	Bovine serum albumin
CML	N ^ε -(carboxymethyl)lysine
DNPH	2,4-Dinitrophenylhydrazine
DPPH	2,2-Diphenyl-1-picrylhydrazyl
DPTC	4,5-Dimethyl-3-phenacylthiazolium chloride
DTNB	5,5'-Dithiobis-(2-nitrobenzoic acid)
ELISA	Enzyme-linked immunosorbent assay
FI	Fluorescence intensity
GO	Glyoxal
GRAS	Generally recognized as safe
MGE	Mogroside extract
MGO	Methylglyoxal

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NBT	4-Nitro blue tetrazolium
ORAC	Oxygen radical absorbance capacity
PBS	Phosphate buffered saline
ROS	Reactive oxygen species
SD	Standard deviation
TCA	Trichloroacetic acid
Trolox	6-Hydroxy-2,5,7,8-tetramethylchromane-2-carboxylic acid

Introduction

Nonenzymatic glycation reaction, also known as Maillard reaction, occurs between carbonyl groups of monosaccharide and nucleophilic compounds mainly from amino acids, proteins, lipids and nucleic acids (Singh et al. 2001). A complex group of end products, termed as advanced glycation end products (AGEs) are formed through condensation, enolization, dehydration, rearrangement, cyclization, fragmentation, and oxidation reactions during glycation in vitro and in vivo (Fig. 1). Aging and diabetes related hyperglycemia induces an accelerated progression on AGEs formation (Levine and Stadtman 2001). A growing body of evidence indicates that AGEs formation and accumulation is greatly related to microvascular and macrovascular diseases, such as diabetes complications, atherosclerosis and Alzheimer's disease (Ahmed 2005; Takeda et al. 1996). Thus, AGEs inhibitors are promising therapeutic agents to prevent or alleviate these diseases.

Apart from glycation, a large number of studies revealed that reactive oxygen species (ROS) is not only generated by glycooxidation, but also induces and enhances glycation (Hunt et al. 1993; Nagai et al. 1997). Although ROS is not the exclusive mechanism which mediates glycation and AGEs formation, many compounds exhibit powerful antiglycative activity which is mainly ascribed to their ROS-scavenging or antioxidant activities (Aljohi et al. 2016; Jariyapamornkoon et al. 2013). Consequently, compound with potent antioxidant activity may be a promising antiglycative agent. Additionally, the side effects of synthetic AGEs inhibitors, such as aminoguanidine (AG), pyridoxamine, tenilsetam and metformin have been documented. For instance, the side effects and safety concerns of AG in clinical trials have been reported, including gastrointestinal disturbance, liver disorder, flu-like symptoms, rare vasculitis and kidney tumours (Freedman et al. 1999; Thornalley 2003). In light of the fact, much more attention has been drawn to natural origin or food-derived AGEs inhibitors due to minor or no adverse effects. Triterpenoid glycosides characterized as 30-carbon skeletal structure with various glycosyl residues are widely distributed in plant kingdom. Although antioxidant and antiglycative properties of triterpenoid glycosides have not

been reported frequently, some specific glycosides or extracts exhibited considerable antioxidant and/or antiglycative activities (Bi et al. 2012; Xi et al. 2010). Mogrosides, principal active components present in *Siraitia grosvenorii* (Swingle) C. Jeffery, are cucurbitane-type triterpene glycosides and have been granted as GRAS (generally recognized as safe) substances by U.S. Food and Drug Administration with low calorie and extreme sweetness. Recently, antiallergic, anticarcinogenic, and antibacterial activities of mogrosides have been reported (Jin and Lee 2012). Our previous studies showed that mogrosides were effective oxygen radicals scavengers in vitro (Chen et al. 2007) and restored decreased level of antioxidant enzymes in alloxan-induced type 1 diabetic mice model. (Qi et al. 2006, 2008). To our best knowledge, however there are no report on antiglycative activity of mogrosides, and the correlation between antiglycation with its antioxidant properties. Therefore, the present study aimed to explore the antioxidant and antiglycative activities of mogrosides and attempted to clarify the relationship between antiglycation and antioxidation.

Materials and methods

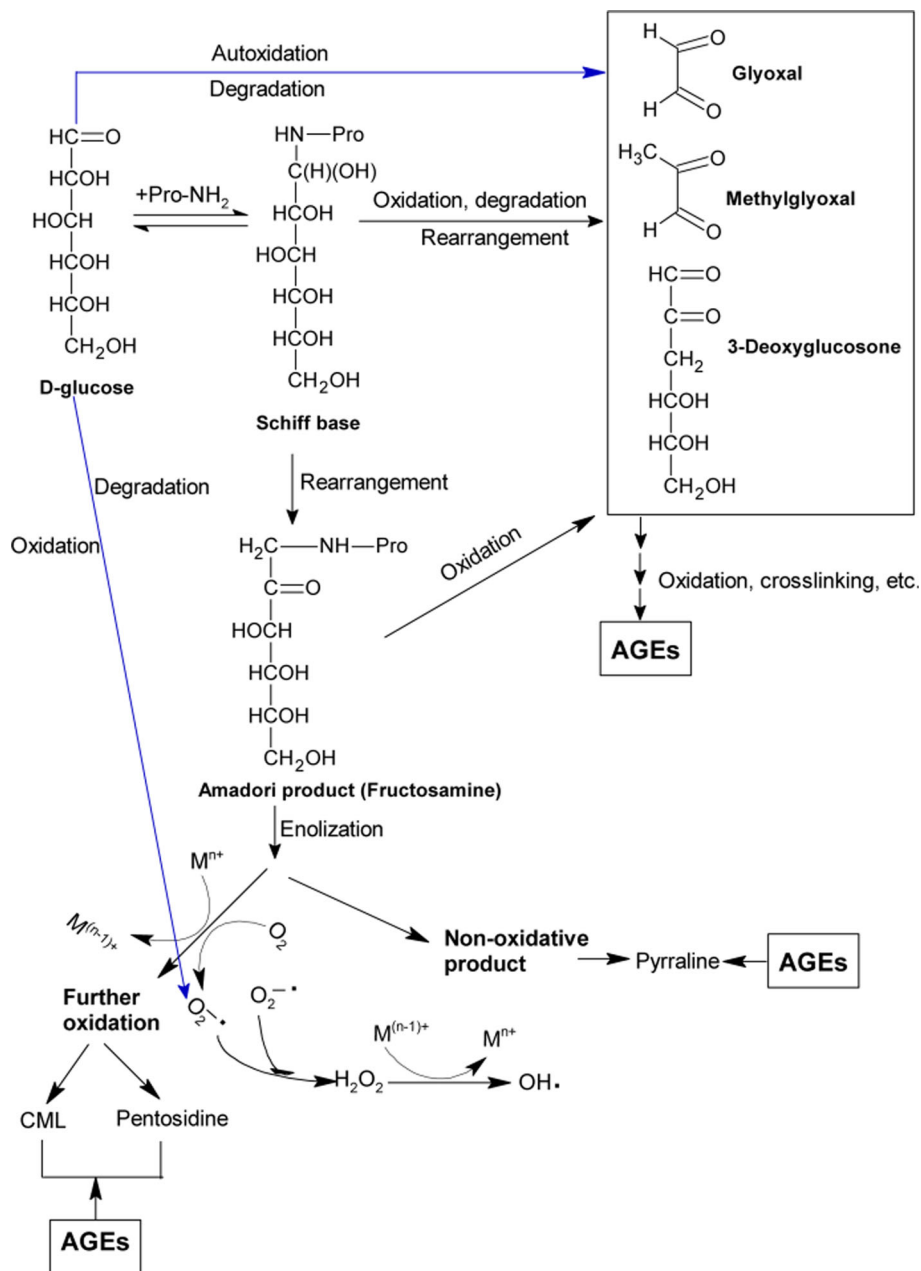
Chemicals and materials

Mogroside extract (MGE) was provided by Guilin Layn Natural Ingredients Corp. Gallic acid, Folin–Ciocalteu's reagent, bovine serum albumin (BSA), D-(+)-glucose, thioflavin T, L-cysteine and aminoguanidine hydrochloride (AG) were purchased from Sigma-Aldrich (St. Louis, MO, USA). N^ε-(carboxymethyl) lysine (CML) ELISA kit was purchased from Usen Life Science, Inc. (Wuhan, China). Fructosamine kit was obtained from Nanjing Jiancheng Bioengineering Institute (Nanjing, China). Solvents used for liquid chromatography were of HPLC grade. Water was produced from a Milli-Q purification system (Bedford, MA, USA). Mogroside standards were purchased from Shanghai Tauto Biotech Co., Ltd. (Shanghai, China) or Chengdu Must Biotechnology Co., Ltd. (Chengdu, China). All other reagents were of analytical grade.

In vitro BSA-glucose glycation model

In vitro glycation was performed according to a previous method with minor modifications (Meepprom et al. 2013). Briefly, 0.1 M phosphate buffered saline (PBS), pH 7.4 containing 10 mg/mL BSA, 0.5 M glucose and 0.02% sodium azide without or with 31, 125 and 500 µg/mL MGE was incubated in dark at 37 °C for 4 weeks. Equal amount of BSA, without glucose and MGE, dissolved in the same buffer was prepared as a blank. AG instead of MGE was

Fig. 1 General scheme of advanced glycation end products (AGEs) formation pathways in vivo. CML, N^ε-(carboxymethyl)lysine; M, transition metal; n, valence of transition metal



used as a positive control. Aliquots of these reaction mixtures were taken for determination of AGEs formation, fructosamine level, protein carbonyl and thiol group contents, and amyloid cross- β structure every week. At the last week, CML level was assayed.

AGEs formation

AGEs formation was assayed by measuring fluorescence intensity (FI) of the mixture using an Infinite 200 Pro microplate reader (Tecan, Männedorf, Switzerland) at excitation and emission wavelengths of 355 and 460 nm, respectively. The percent inhibition of AGEs formation

was calculated by FI of MGE- or AG-treated group divided by that of BSA-glucose group.

Fructosamine Content

The fructosamine level was analyzed by a commercial kit according to the manufacturer's protocol. Briefly, 100 μ L glycated BSA was incubated with 2.0 mL of 0.5 mM 4-nitro blue tetrazolium (NBT) in 0.1 M, pH 10.4 carbonate buffer at 37 $^{\circ}$ C for 15 min. The absorbance was measured at 530 nm. The fructosamine level was calculated by calibration curve with 1-deoxy-1-morpholino-fructose (1-DMF) as standard.

Protein carbonyl content

Protein carbonyl group was determined according to the reported method with minor modification (Meeprom et al. 2013). Briefly, 0.8 mL glycated aliquots was added to 3.2 mL 10 mM 2,4-dinitrophenylhydrazine (DNPH) solution (dissolved in 2.0 M hydrochloric acid solution). After incubation in the dark for 1 h, 4.0 mL of 20% (w/v) pre-cooled trichloroacetic acid (TCA) was added to the mixture to precipitate protein and was placed on ice for 5 min. Pellet was collected by centrifugation at 4 °C, 10,000×g for 10 min and washed three times with 2.0 mL of ethanol/ethyl acetate (1:1, v/v) mixture. Finally, it was resuspended in 2.0 mL of 6 M guanidine hydrochloride, and spectrophotometrically measured at 370 nm. The extinction coefficient of DNPH ($\epsilon = 2.2 \times 10^4 \text{ M}^{-1}\text{cm}^{-1}$) was employed to calculate the carbonyl content (nmol/mg protein) of glycated BSA.

Protein thiol group

Free thiol group of native and modified BSA was measured using earlier method with minor modifications (Meeprom et al. 2013). Briefly, 350 μL of glycated samples were mixed with 650 μL 5,5'-dithiobis-(2-nitrobenzoic acid) (DTNB) (5 mM) in 0.1 M PBS, pH 7.4 and was left at ambient temperature for 15 min. The absorbance was read at 410 nm and the free thiol concentration of samples was calculated from the L-cysteine standard curve and expressed as nmol thiol group/mg protein.

Amyloid cross- β structure

Amyloid cross- β structure in glycated BSA was measured according to earlier method with slight modifications (Meeprom et al. 2013). Briefly, 2.0 mL of 64 μM thioflavin T in 0.1 M PBS, pH 7.4 was added to 200 μL of the glycated samples and incubated at 25 °C for 60 min. FI was measured at an excitation wavelength and emission wavelength of 435 and 485 nm, respectively. Background FI was subtracted from the same buffer without thioflavin T.

CML level

An enzyme-linked immunosorbant assay (ELISA) kit was applied to determine CML level according to the manufacturer's protocol.

Oxygen radical absorbance capacity (ORAC) assay

The ORAC assay was performed in a black-walled microplate according to a literature method (Zheng et al. 2012).

DPPH radical scavenging assay

Briefly, 1.0 mL serially diluted MGE aqueous solutions (50–5000 $\mu\text{g}/\text{mL}$) were added to 3.0 mL DPPH methanolic solution (0.1 mM). The mixture was shaken to mix thoroughly and allowed to stand at room temperature in the dark for 30 min. The absorbance was measured at 517 nm against methanol blank. Distilled water instead of sample was used as control and methanol instead of DPPH solution as sample control for background elimination. Ascorbic acid was used as a positive control. The percent scavenging activity was calculated according to the following equation:

$$\begin{aligned} &\text{Scavenging of DPPH radical (\%)} \\ &= \frac{A_C - (A_S - A_{SC})}{A_C} \times 100 \end{aligned}$$

where A_C is the absorbance of control (distilled water instead of sample), A_S is the absorbance in the presence of the sample, and A_{SC} is the absorbance of sample control (methanol instead of DPPH solution).

ABTS radical scavenging assay

The method reported by Re et al. (1999) was employed with minor modification. Briefly, ABTS radical cation (ABTS^+) stock solution was prepared by mixing 7 mM ABTS aqueous solution with 2.45 mM potassium persulphate, and incubation in the dark at room temperature for 16 h. Prior to use, the ABTS^+ stock solution was diluted to an absorbance of 0.70 ± 0.02 at 734 nm with pH 7.4, 0.1 M sodium phosphate buffer. 0.2 mL sample diluted solutions (50–5000 $\mu\text{g}/\text{mL}$) was mixed well with 3.8 mL diluted ABTS^+ solution and kept in the dark for 6 min before measurement. The absorbance was recorded at 734 nm. Distilled water instead of diluted ABTS^+ solution was used as a blank and trolox as a positive control.

Reducing power assay

The reducing power was determined by the published method with slight modifications (Zheng et al. 2012). 2.0 mL sample aqueous solution (1.0 mg/mL) was mixed with 2.0 mL 0.2 M phosphate buffer (pH 6.6) and 2.0 mL of 0.1% potassium ferricyanide. After incubation at 50 °C for 20 min, 2.0 mL of TCA solution (10%) was added to the mixture. Supernatant was collected by centrifugation at 3000 rpm for 10 min. The supernatant (2.0 mL) was mixed well with equal volume distilled water and 0.4 mL ferric chloride solution (0.3%), and then the absorbance was measured at 700 nm. Ascorbic acid was used as a positive control.

Statistical analysis

All data is obtained from experiments independently performed in triplicate, and is presented as mean \pm standard deviation (SD) from triplicate experiments. Difference between groups is analyzed by one-way analysis of variance (ANOVA), followed by Duncan's post hoc multiple comparisons. $P < 0.05$ is defined as statistical significance.

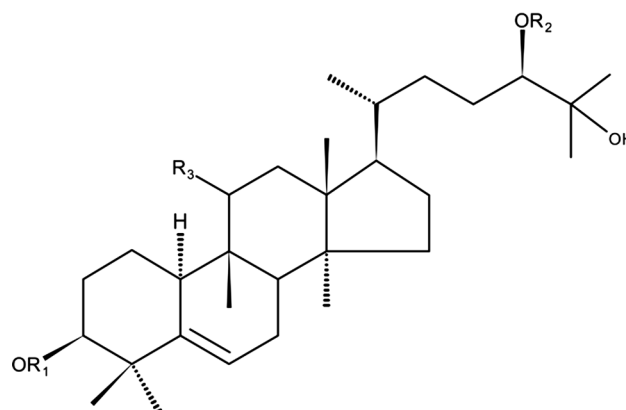
Results and discussion

Antioxidant activities of MGE

The chemical composition of MGE was as follows: moisture: 8.69 ± 0.21 g/100 g, total phenolics: 1.43 ± 0.18 g/100 g, total flavonoids: 0.38 ± 0.03 g/100 g, total mogrosides (colorimetric assay): 80.14 ± 0.50 g/100 g. Individual mogrosides were assayed by a HPLC method, and the content of 11-*O*-mogroside V (11-*O*-MGV), mogroside V (MG V), mogroside VI (MG VI), mogroside IV (MG IV), mogroside III (MG III) and mogroside IIA2 (MG IIA2) was 7.34 ± 0.16 , 44.52 ± 1.33 , 4.58 ± 0.45 , 0.97 ± 0.05 , 0.58 ± 0.03 and 0.32 ± 0.14 g/100 g, respectively (see Electronic Supplementary Material). The results showed that MGE was predominantly composed of mogrosides, particularly MG V, with minor phenolic compounds.

The antioxidant activities of MGE is shown in Table 1. The IC_{50} of MGE on DPPH and ABTS radicals scavenging assays were 1118.1 and 1473.2 μ g/mL, respectively, whereas IC_{50} of ascorbic acid in DPPH and trolox in ABTS was 9.6 and 47.9 μ g/ml, respectively. However, MGE exhibited strong inhibition on peroxy radicals generated by thermal decomposition of AAPH in ORAC assay. In addition, the data showed that MGE possessed weaker reducing power compared to ascorbic acid. These results indicated that MGE possessed considerable peroxy radicals-scavenging activities, with moderate DPPH and ABTS radicals-scavenging capacities. Furthermore, the content of total mogrosides was more than 80%, as well as a few amount of phenolic compounds (less than 2%) in MGE. In light of these results, it could be speculated that mogrosides possibly account for the antioxidant activities of MGE, in spite of no

direct evidence provided in the study. In fact, a variety of triterpene glycosides with antioxidant or radical-scavenging activities have been recently reported (Allouche et al. 2010; Bi et al. 2012; Xi et al. 2008, 2010). It is noteworthy that these antioxidant terpenoids are mostly pentacyclic triterpene glycosides, with either multiple hydroxyl groups or one hydroxyl group and at least one carboxyl group (Allouche et al. 2010), which may act as an electron or hydrogen donor. However, most types of reported mogrosides, including MG V and 11-*O*-MG V, are cucurbitane-type tetracyclic triterpene glycosides, and they are not easily prone to provide



mogrol: $R_1=R_2=H$, $R_3=\alpha-OH$

mogroside II A₂: $R_1=Glc$, $R_2=Glc$, $R_3=\alpha-OH$

mogrosidelll: $R_1=Glc$, $R_2=Glc$, $R_3=\alpha-OH$

siamenoside I : $R_1=Glc$, $R_2=Glc$, $R_3=\alpha-OH$

mogroside IV A: $R_1=R_2=Glc$, $R_3=\alpha-OH$

mogroside V : $R_1=Glc$, $R_2=Glc$, $R_3=\alpha-OH$

mogroside VI : $R_1=Glc$, $R_2=Glc$, $R_3=\alpha-OH$

11-oxomogroside V : $R_1=Glc$, $R_2=Glc$, $R_3=O$

Fig. 2 Chemical structures of major mogrosides from *Siraitia grosvenorii*

Table 1 Antioxidant activities of MGE

	DPPH (IC_{50} , μ g/mL)	ABTS (IC_{50} , μ g/mL)	ORAC (μ mol TE/g MGE)	Reducing power (A_{700nm}) ^a
MGE	1118.1 ± 38.6	1473.2 ± 85.2	851.8 ± 58.7	0.387 ± 0.006
Trolox	–	47.9 ± 2.3	–	
Ascorbic acid	9.6 ± 1.9	–	–	2.850 ± 0.003

^aAll of the concentrations of MGE, trolox and ascorbic acid are 1.0 mg/mL

active electron and hydrogen to quench radicals considering their chemical structures (Fig. 2), which corresponded with weaker DPPH and ABTS radicals-scavenging abilities, compared to ascorbic acid or trolox. On the contrary, ORAC assay showed that MGE was a powerful peroxy radicals scavenger. This discrepancies may be explained by the synergistic effect of mogrosides with minor phenolics present in MGE or synergistic effect among individual mogrosides with different chemical structures. Actually, Xi et al. (2010) reported that individual triterpene glycosides from *Aralia taibaiensis* interacted synergistically to enhance free radicals scavenging activity. Another explanation may be that minor polyhydroxy tetracyclic triterpenoids and pentacyclic triterpenoids in MGE, such as 5α , 6α -epoxymogroside I E1, 20-hydroxy-11-oxomogroside I A1, isomultiflorenol and β -amyrin (Jin and Lee 2012) contribute to peroxy radicals-scavenging activity. These explanations should be studied further.

Inhibitory effect of MGE on fluorescent AGEs formation

AGEs are a group of complex and heterogeneous compounds with or without fluorescence. Although not all fluorescent compounds formed during protein glycation have been isolated and fully characterized, a variety of fluorescent AGEs such as crossline, pentosidine and pyrrolopyridinium, was proven as fluorophores (Matiacevich et al. 2005; Obayashi et al. 1996). Therefore, specific fluorescence of AGEs has been recommended as an indicator for the formation of AGE-modified proteins (Matiacevich et al. 2005). As observed in Fig. 3a, compared to BSA group, FI of glycated BSA increased greatly with increased incubation time, indicating more fluorescent AGEs formation, while co-incubation with MGE inhibited fluorescent AGEs formation in a dose-dependent manner throughout the experimental

period. At the end of experimental period, the percent inhibition of MGE on AGEs formation was 11.6, 33.6 and 58.5% at corresponding concentrations to 31, 125, and 500 $\mu\text{g}/\text{mL}$, respectively, while inhibitory rate of AG was found to 81.6% at 500 $\mu\text{g}/\text{mL}$.

In view of the structural diversity of reported AGEs inhibitors, various antiglycation mechanisms have been proposed, including scavenging radicals, blocking reactive carbonyls, chelating transition metal ions, inhibiting the formation of Amadori products and breaking cross-link protein structures. With regard to AGEs inhibitor of natural origin, the first three mechanisms are predominant (Wu et al. 2011). Naturally occurring terpenoids, especially triterpene glycosides, are structurally difficult to donate electron or hydrogen to neutralize radicals or oxidants as phenolic compounds act (Chae et al. 2010). Consequently, they are not prone to contribute to antiglycation activity through their radicals-scavenging capacities. Actually, most triterpene glycosides which exhibited strong antiglycation properties perform as either reactive carbonyls blockers or metal ions chelators or both. The study of Chompoo et al. (2011) indicated that aldehyde groups of labdadiene, a diterpenoid from *Alpinia zerumbet*, blocked active carbonyls and subsequently inhibited AGEs formation, while maslinic acid, a pentacyclic triterpenic acids with vicinal hydroxyl groups at C-2 and C-3 positions, was reported to possess strong copper-chelating activity (Allouche et al. 2010). Although major mogrosides (Fig. 2) are structurally not in agreement with these characteristics described above, MGE exhibited potent antiglycative activity in the present study. The plausible explanation for this is unknown, but it seems that some mogrosides with specific chemical structures, such as carbonyl-containing 11-*O*-MG V and siraitic acid C, as well as polyhydroxy-containing 5α , 6α -epoxymogroside I E1, present in MGE contribute to antiglycation of MGE. Further studies are needed to confirm it.

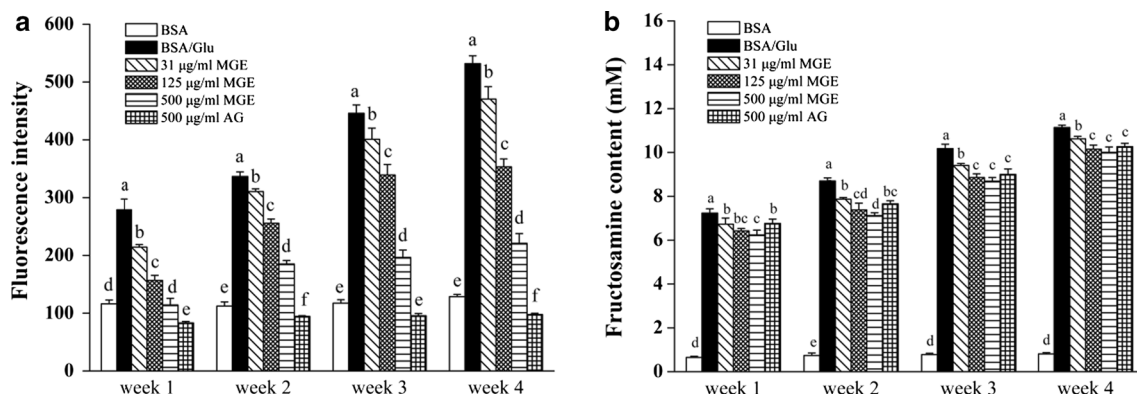


Fig. 3 Inhibitory effect of MGE on the formation of fluorescent AGEs (a) and fructosamine level (b) in BSA-glucose model. Results are expressed as mean \pm SD from triplicate experiments. Different

lowercase letters indicated statistical difference between groups ($P < 0.05$), the same as follow

Inhibitory effect of MGE on fructosamine

Fructosamine is an oxidative product from Amadori adducts during the early stage of nonenzymatic glycation, and used as a marker for protein glycation (Singh et al. 2001). The inhibition of MGE and AG on fructosamine formation is shown in Fig. 3b. The fructosamine level in BSA-glucose group increased dramatically, while the level in BSA group was very low and almost unchanged throughout the incubation period. In contrast, MGE dose-dependently suppressed the formation of fructosamine, and at week 4, the inhibitory rate of MGE (31–500 $\mu\text{g}/\text{mL}$) on fructosamine formation was 4.7–10.2%, while AG was 7.8% at the concentration of 500 $\mu\text{g}/\text{mL}$, which indicated that MGE was comparable to AG in preventing early protein glycation.

Inhibitory effect of MGE on protein oxidation

Protein oxidation, generated carbonyl groups and deplete free thiols (Singh et al. 2001; Zheng et al. 2016), can not only be mediated by free radicals and/or ROS from monosaccharide autoxidation and glycooxidation (Wolff and Dean 1987), but also by reactive α -oxoaldehydes, such as glyoxal (GO), methylglyoxal (MGO) and 3-deoxyglucosone (3-DG) derived from oxidative degradation of Amadori adducts or from lipid peroxidation.

Compared to BSA-glucose group, MGE effectively inhibited carbonyl increase and the effect of MGE at low and middle dose (31–125 $\mu\text{g}/\text{mL}$) was comparable to that of AG at 500 $\mu\text{g}/\text{mL}$ after 4 week incubation. High dose of MGE was more effective than AG and reversed carbonyl level near to BSA control (Fig. 4a). Likewise, the level of free thiol remarkably decreased in BSA-glucose group compared to BSA control, whereas co-incubation with MGE slightly increased free thiols (Fig. 4b). Increasing carbonylation has been considered as one of the

hallmarks of protein oxidation, which is induced by interaction with active carbonyls, mainly from degradation of Amadori products during in vitro glycation. Another hallmark of protein oxidation is depletion of free thiols which are susceptible to oxidative attack by ROS and form adducts with active dicarbonyls (Ahmed 2005; Quiney et al. 2011). It should be noted that AG co-incubation with glucose-BSA mixture for 4 weeks only retained a little more free thiols than glycated BSA (8.6%) (Fig. 4b), although it moderately reduced protein carbonylation (17.3%) (Fig. 4a). It may be explained by the fact that AG exerts anti-glycation effect mainly by neutralizing reactive carbonyls arising from glycooxidation (Thornalley 2003). However, free thiols are depleted by ROS generated from oxidative cleavage of Amadori adducts, which is independent on its carbonyls-trapping property (Glomb and Monnier 1995; Hunt et al. 1993).

Inhibitory effect of MGE on CML and amyloid cross- β structure formation

To determine the inhibitory effect of MGE on BSA glycooxidation, non-fluorescent and non-crosslinking CML level was tested by ELISA. As shown in Fig. 5a, CML level of non-glycated BSA was negligible (only 11.70 ng/mL), whereas it rose sharply in glycated BSA ($P < 0.05$) at the fourth week. AG at 500 $\mu\text{g}/\text{mL}$ strongly inhibited CML formation, while MGE treatment dose-dependently reduced CML level and the inhibitory activity of MGE (500 $\mu\text{g}/\text{mL}$) was comparable to AG at 500 $\mu\text{g}/\text{mL}$, suggesting potent inhibition of MGE on CML formation.

The oxidative cleavage of Amadori products (Ođjakova et al. 2012) and Schiff base (Glomb and Monnier 1995), as well as glucose autoxidation-mediated pre-Amadori adducts pathway (Wells-Knecht et al. 1995a, b), are involved in CML formation. Regardless of the three pathways, ROS are involved in the formation of CML

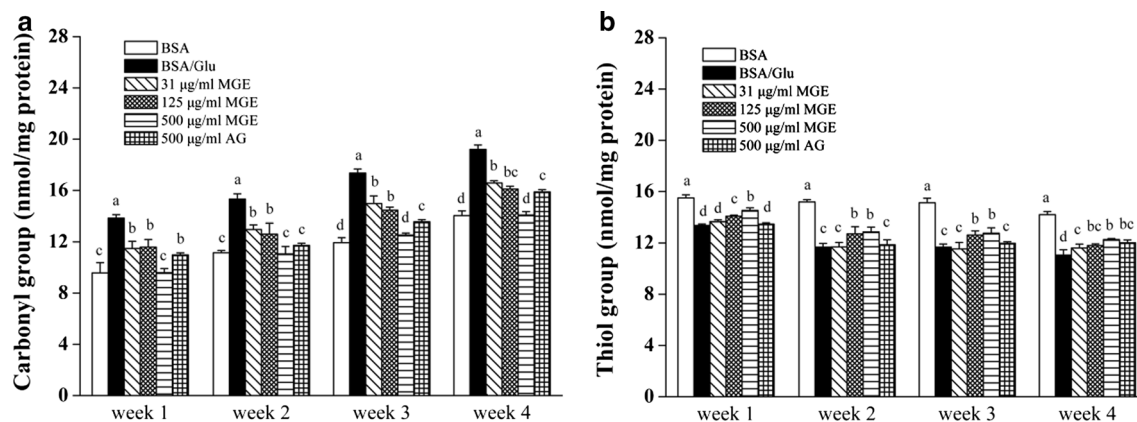


Fig. 4 Inhibitory effect of MGE on protein oxidation in BSA-glucose model. Protein carbonyl level (a) and free thiols level (b)

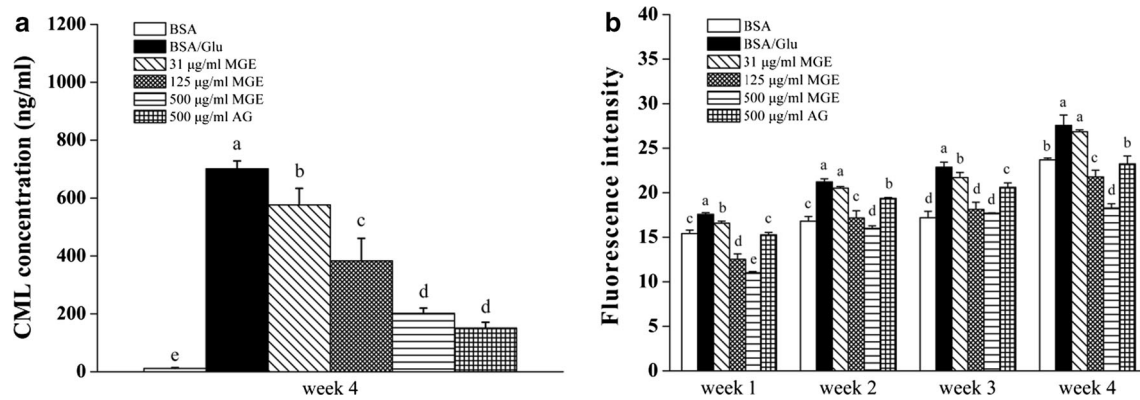


Fig. 5 Inhibitory effect of MGE on CML formation (a) and amyloid cross- β structure (b) in BSA-glucose model

(Odjakova et al. 2012). Nagai et al. (1997) found that hydroxyl radical directly accelerated CML formation from Amadori products in vitro. Therefore, it is credible that any compound which possesses strong radicals-, especially hydroxyl radical-scavenging activities, can inhibit CML formation. In the current study, although the scavenging capacity of MGE on hydroxyl radical was not tested, it showed considerable peroxy radicals and moderate DPPH and ABTS radicals scavenging activities, which may contribute to inhibition of CML (Fig. 5a). Actually, our previous study showed that both MG V and 11-*O*-MG V were highly reactive to scavenge radicals in chemiluminescence systems, and MG V was more effective than 11-*O*-MG V on hydroxyl radical, while 11-*O*-MG V was more potent on hydrogen peroxide and superoxide anion radicals (Chen et al. 2007). Therefore, it can be concluded that the radical-scavenging properties of MGE, at least in part, contribute to its inhibition on CML formation.

Amyloid cross- β structure, as a result of irreversible protein–protein crosslinking and abnormal aggregation, is predominantly driven by nonenzymatic reactions between reactive carbonyls and amino acid residues of protein (Liu et al. 2016). Two effective approaches have been proposed to reduce this crosslinking. One method acts as a carbonyl-quenching agent by inhibiting the formation of amyloid structure, such as AG, and other functions as a dicarbonyl-breaking agent, for example, 4,5-dimethyl-3-phenacylthiazolium chloride (DPTC) by cleaving crosslinked structure (Ulrich and Cerami 2001). As observed in Fig. 5b, compared to non-glycated BSA, glycated BSA exhibited elevated amyloid cross- β structure by 16.3% at the end of this study, while co-incubation with MGE (31–500 μ g/mL) reduced the level of amyloid cross- β structure ranging from 2.6–33.6%. Furthermore, treatment with MGE at 125 and 500 μ g/mL reversed amyloid cross- β structure near to BSA group, and were more effective than AG at 500 μ g/mL. Although the precise antiglycation mechanism needs to be studied further, some carbonyl-containing mogrosides,

such as 11-*O*-MG V (Fig. 2), may be responsible for the antiglycative properties of MGE. It is possible that carbonyl groups present in 11-*O*-MG V compete with glucose to react with amino groups of protein, which inhibits formation of Schiff base, and/or restricts amino groups to attach glucose (Odjakova et al. 2012). Similarly, carbonyl-containing labdadiene, a diterpenoid compound, showed more antiglycative activity than other diterpenoids without free carbonyl groups (Chompoo et al. 2011).

Conclusion

MGE exhibited considerable antioxidant capacity by scavenging peroxy radicals and potent antiglycative activities by alleviating glucose-mediated protein glycoxylation and crosslinking in vitro. Although there is no direct evidence that antioxidant properties of MGE contribute to its antiglycation activity, our findings indicate that MGE may be used as a new potential antiglycation agent for treatment of diabetes and its complications.

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Compliance with ethical standards

Conflict of interest The authors declare that there are no competing interests.

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