

Saponins in Yerba Mate Tea (*Ilex paraguariensis* A. St.-Hil) and Quercetin Synergistically Inhibit iNOS and COX-2 in Lipopolysaccharide-Induced Macrophages through NF κ B Pathways

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Yerba mate tea (*Ilex paraguariensis*) is growing in popularity around the world. The objective of this study was to investigate the potential anti-inflammatory effect of yerba mate tea (MT) extracts as well as some of its phytochemicals and their interactions. MT and decaffeinated MT extracts [1–300 μ M chlorogenic acid (CHA) equiv]; CHA, caffeine from MT (matein), and mate saponins (1–300 μ M); quercetin (1–200 μ M); and ursolic and oleanolic acids (1–100 μ M) were tested by measuring their ability to inhibit COX-2/PGE₂ and iNOS/NO pathways in LPS-induced RAW 264.7 macrophages. Mate saponins (IC₅₀ = 20 μ M) and oleanolic acid (IC₅₀ = 80 μ M) significantly inhibited iNOS/NO pathways, whereas ursolic acid showed low or no inhibition at 100 μ M. Quercetin was the most potent inhibitor of pro-inflammatory responses at a concentration 10 times lower than the concentrations used of other compounds (IC₅₀ = 11.6 μ M for NO, 7.9 μ M for iNOS, and 6.5 μ M for PGE₂). Combination of quercetin/mate saponins (0.001:0.004, molar ratio) resulted in synergistic interaction inhibiting both NO and PGE₂ production. It also suppressed IL-6 and IL-1 β production and resulted in reduction of LPS-induced nuclear translocation of nuclear factor- κ B subunits. MT extract did not have a potent anti-inflammatory effect perhaps due to the antagonistic effect of some of its compounds. However, whole MT consumption still has a promising anti-inflammatory outcome mainly through the PGE₂/COX-2 pathway. To the authors' knowledge, this is the first study demonstrating the efficacy, interactions, and mechanisms of some MT phytochemicals in inhibiting pro-inflammatory responses.

KEYWORDS: Cyclooxygenase-2 (COX-2); prostaglandin E₂ (PGE₂); NF κ B; quercetin; RAW264.7 macrophages; saponins; yerba mate tea phytochemicals

INTRODUCTION

Yerba mate tea (*Ilex paraguariensis*) (MT) is rich in many phytochemical compounds such as theobromine, caffeine (matein), caffeoylquinic acid derivatives, dicaffeoylquinic acid derivatives, rutin, luteolin, quercetin, and saponins (1). **Figure 1** presents the chemical structures of some of the important phytochemicals in MT.

Several studies have shown that flavonoids in MT have anti-inflammatory activity; this has important health implications because chronic inflammation can develop into several diseases such as arteriosclerosis, cancer, diabetes, and neurodegenerative diseases (2–5). Published reports suggest that MT has chemopreventive activity against some types of cancer (6–8). However, at the cellular level, it is unclear which compounds in MT are responsible for this effect. Administering agents that cause colitis in healthy rodents or genetically engineered cancer-prone mice accelerated the development of colorectal cancer (9). These observations support the role of inflammation in carcinogenesis. Anti-inflammatory properties of phytochemicals such as

quercetin (10–13) and saponins (14) are documented. However, the anti-inflammatory mechanism of MT and its components has not been studied.

Pro-inflammatory cytokines are involved in the formation of toxic peroxynitrite by increasing the activity of inducible nitric oxide synthase (iNOS). Prostaglandin E₂ (PGE₂) is also an important mediator of inflammatory reaction produced by the enzyme cyclooxygenase-2 (COX-2). COX-2 and iNOS are induced during inflammatory conditions and are overexpressed in colonic tumors in humans, as well as in animals (9). iNOS has been shown to be involved in regulating COX-2 production, which plays an important role in colon tumorigenesis. Hence, suppression of enzyme induction and the activities of iNOS/COX-2 is an important approach to preventing carcinogenesis in several organs including the stomach and colon (15).

Interleukin (IL)-1 β and IL-6 are pro-inflammatory cytokines that can promote acute to chronic inflammation through activation of nuclear factor- κ B (NF κ B). NF κ B is a transcription factor that has five protein subunits: RelA (p65), Rel B, c-Rel, NF κ B1 (p50), and NF κ B2 (p52). These subunits may form homo- or heterodimers. The activated NF κ B is a heterodimer and consists of p65 and p50. In human cells, NF κ B is bound to the inhibitor

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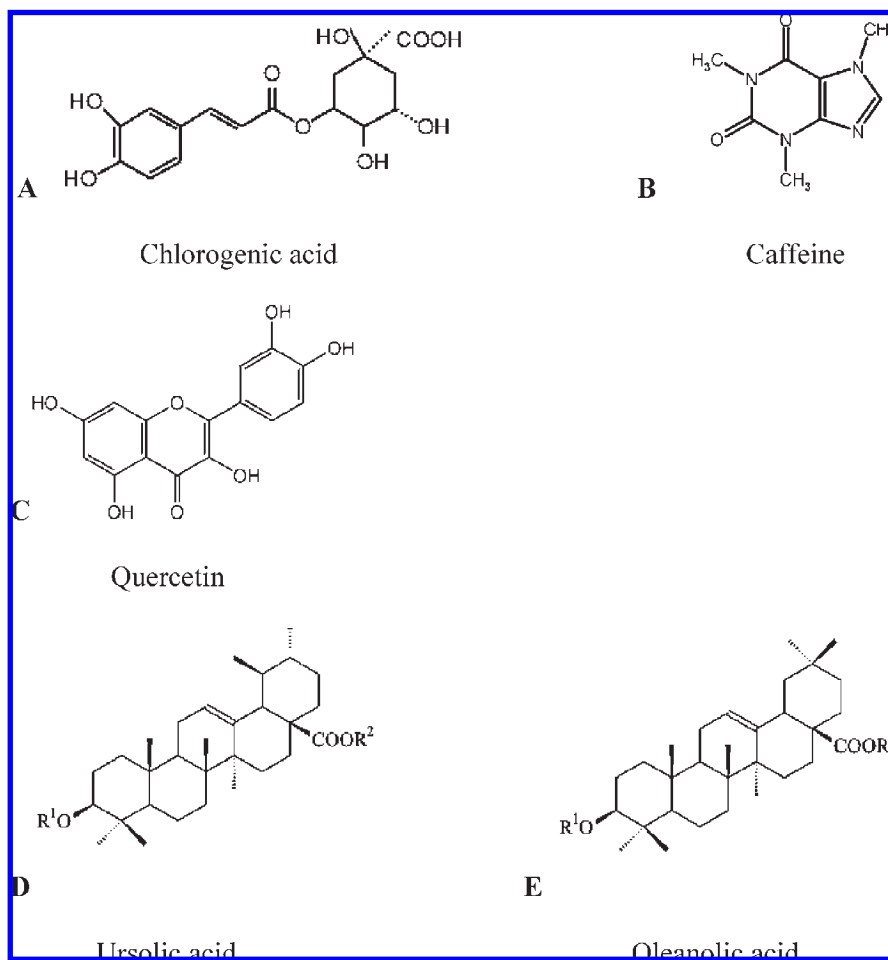


Figure 1. Chemical structures of (A) chlorogenic acid, (B) caffeine (matein), (C) quercetin, (D) ursolic acid, and (E) oleanolic acid.

κ B α ($I\kappa$ B α) in the cytoplasm (15). $I\kappa$ B kinase complex (IKK) can be activated in response to many different signals which result in phosphorylation of $I\kappa$ B α followed by its ubiquitination and proteosomal degradation, causing the $I\kappa$ B α to dissociate from NF κ B. Degradation of $I\kappa$ B α releases NF κ B (p65 and p50), resulting in translocation of the activated NF κ B into the nucleus, which then regulates the transcription of target genes (16). NF κ B regulates several genes involved in immunity, inflammation, and apoptosis and promotes the expression of cytokines such as IL-1 β , IL-6, and TNF α , and up-regulates inflammatory enzymes such as COX-2 and iNOS. Thus, NF κ B is an emerging target for the prevention of cancer and inflammation (15). For instance, ursolic acid reduces pro-inflammation cytokines through down-regulation of NF κ B (17).

We hypothesized that phytochemicals present in MT will be able to inhibit inflammation through NF κ B pathways. The objective of this research was therefore to determine the effect of selected MT phytochemicals for their ability to suppress inflammation. The anti-inflammatory properties of MT extract, decaffeinated MT extract, chlorogenic acid (CHA), quercetin, matein, mate saponins, ursolic acid, and oleanolic acid were evaluated in vitro using lipopolysaccharide-induced RAW 264.7 macrophages. COX-2/PGE₂ and iNOS/NO pathways as well as the interaction between MT phytochemicals in inhibiting NO and PGE₂ production were investigated. On the basis of the results, a mechanism of action was suggested.

MATERIALS AND METHODS

Chemicals. Chlorogenic acid ($\geq 95\%$ CHA), ursolic acid ($\geq 90\%$), quercetin ($\geq 98\%$), oleanolic acid ($\geq 97\%$), sodium nitrite, sulfanilamide,

N-1-(naphthyl)ethylenediamine-diHCl, and lipopolysaccharide (LPS) from *Escherichia coli* O55:B5) were purchased from Sigma (St. Louis, MO). Macrophage RAW 264.7 cell line and Dulbecco's Modified Eagle Medium with L-glutamine (DMEM) were purchased from American Type Culture Collection (Manassas, VA). Fetal bovine serum was purchased from Invitrogen (Grand Island, NY). COX-2, iNOS, NF κ B p50, and NF κ B p65 mouse monoclonal antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Anti-mouse IgG conjugated horseradish peroxidase secondary antibody was purchased from GE Healthcare (Buckinghamshire, U.K.). Luciferase assay kit and TransFast transfection reagent were purchased from Promega (Madison WI) and pNF- κ B-luciferase vector (pNF- κ B luc vector) was purchased from Clontech (Mountain View CA). All other chemicals were purchased from Sigma, unless otherwise specified.

Mate Tea Preparation. Organically grown mate (*Ilex paraguariensis*) leaves from Paraguay were kept in plastic bags and refrigerated at 4 °C until used. MT was prepared by using the traditional North American procedure of preparing tea. Briefly, 2.5 g of dry tea leaves (DL) was soaked in 250 mL of boiling water for 10 min. The infusion was cooled to room temperature before filtration using Whatman paper no. 2 to eliminate dry tea leaves and then freeze-dried in a Labconco, Shell Freeze System (Fisher Scientific, Pittsburgh, PA). The freeze-dried material called MT extract, equivalent to instant tea, was kept at -20 °C protected from light. The phytochemical composition of MT under investigation was previously published by Heck et al. (35).

Preparation of Decaffeinated Mate Tea. Matein and decaffeinated MT were obtained by supercritical CO₂ extraction of MT leaves. The supercritical extraction apparatus was provided by the U.S. Department of Agriculture (Peoria, IL). Matein was extracted from organic MT using a water-saturated CO₂ supercritical extraction. The temperature for the extraction vessel was 70 °C with a pressure of 398 bar. The temperature of the receiver vessel was 50 °C with a pressure of 75 bar. The flow rate was

466 mL/min of liquid CO₂. A total of 1.5 kg of MT leaves was placed in the CO₂ supercritical extractor, and it took approximately 8 h for the total procedure (18). The final product was freeze-dried.

Extraction and Purification of Mate Saponins. Mate saponins were isolated from the MT infusions as performed by Gnoatto et al. (19). Briefly, MT infusion was prepared by boiling 15 g of dry leaves for 10 min in 100 mL of water. The infusion was then filtered, brought to 100 mL with water, treated with hydrochloric acid, and then refluxed for 3 h. Mate saponins were extracted four times with 50 mL of chloroform. The chloroform fraction was evaporated to dryness (77.6 mg). A 1050 Hewlett-Packard (Palo Alto, CA) gradient liquid chromatograph equipped with a 1050 HP autosampler, a gradient pump, and a photodiode array detector was used to estimate saponin concentration. HPLC analysis was performed using a C₁₈ RP guard column and a C₁₈ RP Phenomenex Prodigy ODS column (250 mm × 4.6 mm × 5 μm). The variable-wavelength detector was set at 203 nm. The mobile phase consisted of acetonitrile/water (70:30 v/v). Flow rate = 1.0 mL min⁻¹ and sensitivity = 0.001 AUFS. Mate saponins were calculated to have 73% purity. A molecular mass of 1000 g/mol was used in the calculation of the molar concentration of mate saponin.

Macrophage Culture and Cell Viability Assay. Macrophage cell line RAW 264.7 was cultured in growth medium containing DMEM, 1% penicillin/streptomycin, 1% sodium pyruvate, and 10% fetal bovine serum at 37 °C in 5% CO₂/95% air. The cell viability assay was conducted using the CellTiter 96 Aqueous One Solution proliferation assay kit using the novel tetrazolium compound 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium, inner salt (MTS), and an electron coupling reagent, phenazine ethosulfate (PES) (Promega Corp., Madison, WI). Briefly, 5 × 10⁴ cells were seeded in a 96-well plate, and the total volume was adjusted to 200 μL with growth medium. Cells were allowed to grow for 24 h at 37 °C in 5% CO₂/95% air. Cells were then treated with different concentrations of MT phytochemicals for 24 h, the growth medium was replaced by 100 μL of fresh growth medium, and 20 μL MTS/PES was added to each well. The plate was incubated for 2 h at 37 °C, and the absorbance was read at 515 nm. The percentage of viable cells was calculated with respect to cells treated with 0.3% DMSO (for MT extract, decaffeinated MT extract, CHA, matein, quercetin, ursolic acid, oleanolic acid) and 0.3% 1:1 DMSO/methanol (for mate saponins). Solvents used at these concentrations showed no cytotoxicity (viability > 80%). DMSO at < 0.3% presented 95.8% cell viability.

Treatments. Cell treatment was conducted by seeding 2 × 10⁵ cells in a 6-well plate, and cells were allowed to grow for 48 h at 37 °C in 5% CO₂/95% air. After 48 h of incubation, the cells were treated with different concentrations of MT phytochemicals (1, 10, 100, 200, 300 μM) and 1 μg/mL of LPS for 24 h. All MT phytochemicals were sterilized using 0.22 μm nylon filters before the cells were treated. After 24 h of treatment, the spent medium was collected and analyzed for NO, PGE₂, IL-6, and IL-1β. Cell lysates were used to study the effect of MT phytochemicals on the expressions of COX-2 and iNOS.

Nitrite and PGE₂ Measurement. After 24 h of treatment and LPS induction, culture supernatant was collected. For nitrite measurement, 100 μL of the spent medium was plated in 96-well plate and an equal amount of Griess reagent (1% sulfanilamide and 0.1% *N*-1-(naphthyl)ethylenediamine-dihCl in 2.5% H₃PO₄) was added. The plate was incubated for 5 min, and the absorbance was measured at 550 nm. The amount of NO was calculated using sodium nitrite standard curve. For PGE₂ measurement, ELISA kit monoclonal was used following the manufacturer's instructions (Cayman Chemical, Ann Arbor, MI).

Interaction Study and Isobolographic Analysis. Possible interactions between MT phytochemicals in inhibiting NO and PGE₂ production by LPS-induced RAW 264.7 were tested. The molar ratio of phytochemicals tested was approximately equal to its concentration in one cup of tea (1.5 g of DL/150 mL of water); 2.3:0.86 (CHA/matein), 2.3:0.004 (CHA/mate saponins), 2.3:0.001 (CHA/quercetin), 0.004:0.001 (mate saponins/quercetin), 0.004:0.86 (mate saponins/matein), 0.001:0.86 (quercetin/matein). Calculated and experimental percent inhibition values from combination studies were compared to determined expected interactions. Experimental values that were significantly different from the calculated values at 95% confidence interval were defined as synergistic or "more than additive" interactions as previously reported (20). To further understand the interactions, an isobolographic analysis was performed (20). IC₃₅

values for MT phytochemicals were used in constructing isobolograms. A straight line connecting the IC₃₅ values of these MT phytochemicals predicts the IC₃₅ value of an additive effect. If results of the isobolographic analysis deviated significantly to the left of the additivity line with 95% confidence interval, the interaction was confirmed as a synergy; when it deviated to the right, the interaction was antagonistic. If it fell within the confidence limit for the additivity line, the interaction was said to be additive (20). The IC₃₀ and IC₅₀ values for MT phytochemicals were calculated by linear regression using Prism 4.0 (GraphPad Software, San Diego, CA).

Measurement of iNOS and COX-2 Protein Expressions. COX-2 and iNOS expressions were determined in cell lysates. Briefly, treated cells were washed with ice-cold DMEM and ice-cold phosphate-buffered saline before treatment with 200 μL of Laemmli buffer (Bio-Rad Laboratories, Hercules, CA) with 5% β-mercaptoethanol as lysing buffer. The cell lysates were boiled for 5 min, and approximately 25 μg of proteins was loaded in 4–20% Tris-HCl ready gels (Bio-Rad Laboratories) for protein separation. Separated proteins were transferred in PVDF membrane and blocked with 5% nonfat dry milk in 0.1% Tris-buffered saline Tween 20 (TBST) for 1 h at 4 °C. After blocking, the membrane was washed with 0.1% TBST (five times, 5 min each) and incubated overnight with either COX-2 or iNOS mouse monoclonal antibody (1:1000) at 4 °C. The membrane was washed again and incubated with anti-mouse IgG antibody conjugated to horseradish peroxidase secondary antibody for 3–4 h at room temperature. After incubation and repeated washings, the expression of COX-2 and iNOS was visualized using a chemiluminescent reagent (GE Healthcare) following the manufacturer's instructions. The membrane picture was taken with a Kodak Image station 440 CF (Eastman Kodak Co., New Haven, CT).

IL-6 and IL-1β Measurement. For IL-6 and IL-1β measurements, a Bio-Plex singleplex cytokine assay was used (Bio-Rad Laboratories). Briefly, a 96-well filter plate was prewetted with 100 μL of assay buffer and the buffer was removed by vacuum filtration. Then, 50 μL of the bead working solution (IL-6 and IL-1β) was added, and the buffer was removed by vacuum filtration. The plate was washed with 100 μL of wash buffer, and 50 μL of diluted standard or sample was added to each well. The plate was then covered with sealing tape and aluminum foil and incubated at room temperature for 30 min with shaking. After incubation, the buffer was removed and washed with 100 μL of wash buffer three times. Twenty-five microliters of detection antibody was added, and the plate was incubated at room temperature for 30 min with shaking. The plate was washed again, and 50 μL of streptavidin-PE was added and incubated for 10 min at room temperature with shaking. After incubation with streptavidin PE, the plate was washed three times with wash buffer, and the beads were resuspended with 125 μL of assay buffer. The plate was then read using a Bioplex suspension array system using high photomultiplier tube setting standards measuring 100 beads per region. The concentrations of IL-6 and IL-1β were calculated using IL-6 and IL-1β standard curves, respectively.

Transfection of RAW 264.7 Macrophage Cell Line with pNF-κB-Luciferase Vector. RAW 264.7 macrophage cell line was transiently transfected with a pNF-κB-luciferase vector or pTAL-Luc vector (Clontech) using TransFast transfection reagent (Promega) following the manufacturer's protocol. The pNF-κB-luciferase vector contained the firefly luciferase gene from *Photinus pyralis*, whereas the pTAL-Luc vector served as a negative control. Briefly, 5 × 10⁴ cells were plated in a 24-well plate and allowed to grow to 80–90% confluency for 24 h at 37 °C in 5% CO₂/95% air. The day before transfection, the transfection reagent was resuspended in 400 μL of nuclease-free water and stored at -20 °C overnight. The cells were then treated with 200 μL of DNA-Transfast reagent mixture and incubated for 1 h at 37 °C in 5% CO₂/95% air. After 1 h of incubation, each well was then overlaid with 1 mL of complete growth medium, and transfection was carried out for 48 h at 37 °C in 5% CO₂/95% air.

Measurement of Luciferase Activity. After transfection, cells were treated with quercetin (1 μM), mate saponins (4 μM), quercetin/mate saponins combination (1:4 μM), and 1 μg/mL LPS for 24 h. Luciferase activity from the cells was measured using a luciferase assay system (Promega) following the manufacturer's protocol and a Femtomaster FB 12 luminometer (Zylux Corp.). Briefly, growth medium was removed

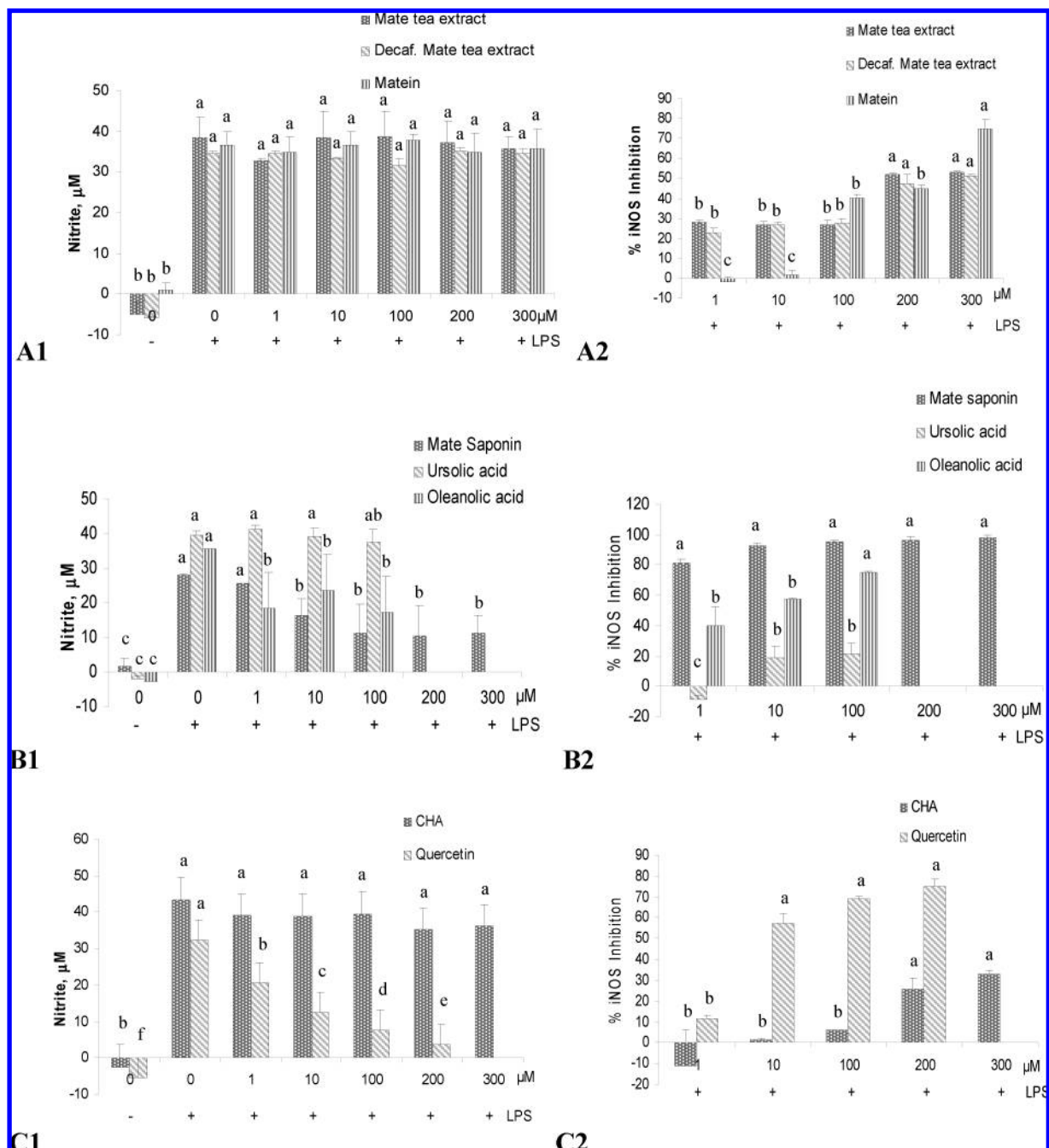


Figure 2. Effect of different concentrations of MT phytochemicals on NO (1) production and iNOS (2) expression by LPS-induced RAW 264.7 cells: (A) MT extract, decaffeinated MT extract, matein; (B) mate saponins, ursolic acid, oleanolic acid; (C) CHA, quercetin. The data represent the mean \pm SD from two independent experiments and at least a triplicate analysis. Different letters indicate significant differences, $P < 0.05$.

and the cells washed with 1 mL of ice-cold PBS. After complete removal of ice-cold PBS, 100 μ L of passive lysis buffer was added and the plate was incubated at room temperature for 15 min with shaking. After incubation, luciferase activity was measured by adding 20 μ L of cell lysate in 100 μ L of luciferase assay reagent and immediately reading the flash using the Fentomaster luminometer. Luciferase activity was normalized on the basis of protein content as determined by Bio-Rad protein assay.

Western Blotting of NF κ B p50 and p65 Protein Expression.

NF κ B p50 and p65 expressions were determined in cell cytoplasm and nucleus. Nuclear and cytoplasmic proteins were isolated with a buffer extraction system and centrifugation NE-PER nuclear and cytoplasmic extraction reagents (Pierce Biotechnology, Rockford, IL) according to the manufacturer's recommendations. Briefly, RAW 264.7 macrophages were plated at a density of 2×10^5 cells per well and allowed to confluence for 48 h at 37 $^{\circ}$ C in 5% CO $_2$ /95% air. Confluent cells were then treated with

quercetin (1 μ M), mate saponins (4 μ M), and quercetin/mate saponins combination (1:4 μ M) and 1 μ g/mL LPS for 24 h. After 24 h of treatment, cells were harvested in 200 μ L of 0.25% trypsin solution. The lysates were then transferred in a 1.5 mL microcentrifuge tube and pelleted by centrifugation at 500g for 2–3 min. After centrifugation, the cell pellet was treated with 200 μ L of ice-cold cytoplasmic extraction reagent I, vortexed for 15 s, and incubated on ice for 10 min and then with treated 11 μ L of ice-cold cytoplasmic extraction reagent II, vortexed for 5 s, and incubated on ice for 1 min. After incubation, the tube was vortexed for 5 s and centrifuged at 16000g for 5 min at 4 $^{\circ}$ C. The supernatant (cytoplasmic extract) was immediately transferred to a clean prechilled 1.5 mL microcentrifuge tube. The pellet was then resuspended in 100 μ L of nuclear extraction reagent, vortexed for 15 min, and incubated on ice for 40 min with vortexing for 15 s every 10 min. The mixture was centrifuged at 16000g for 10 min at 4 $^{\circ}$ C, and the supernatant (nuclear extract) was

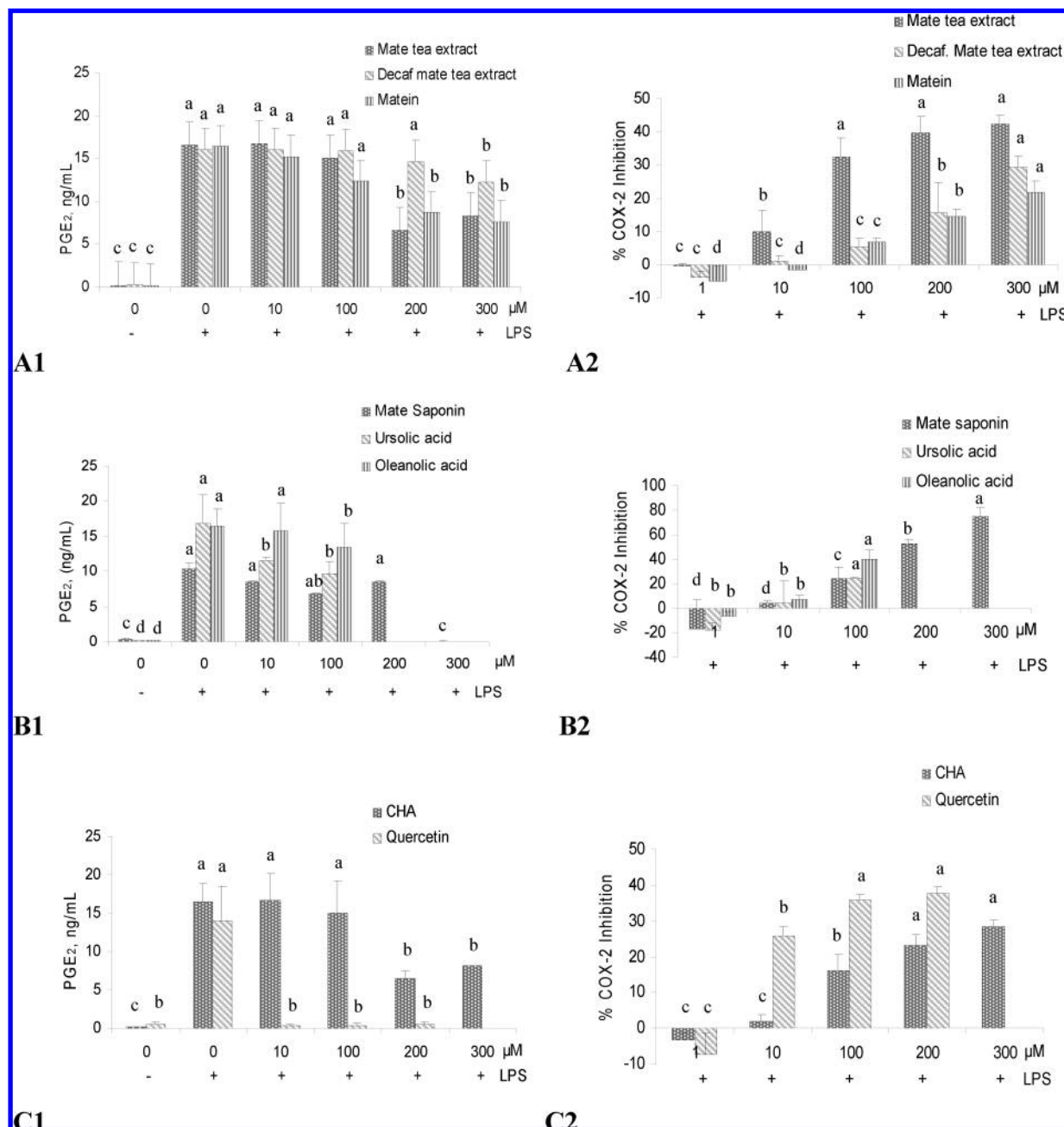


Figure 3. Effect of different concentrations of MT phytochemicals on PGE₂ production (1) and COX-2 expression (2) by LPS-induced RAW 264.7 cells: (A) MT extract, decaffeinated MT extract, matein, (B) mate saponins, ursolic acid, oleanolic acid, (C) CHA, quercetin. The data represent the mean \pm SD from two independent experiments and at least a triplicate analysis. Different letters indicate significant differences, $P < 0.05$.

collected. Both cytoplasmic and nuclear extracts were stored at -80°C until use for Western blot analysis. For the determination of p50 and p65 nuclear translocation, approximately $10\ \mu\text{g}$ of protein was loaded in 4–20% Tris-HCl ready gels (Bio-Rad Laboratories) for protein separation. The separated proteins were transferred in PVDF membrane and blocked with 5% nonfat dry milk in 0.1% Tris-buffered saline Tween 20 (TBST) for 1 h at 4°C . After blocking, the membrane was washed with 0.1% TBST (five times, 5 min each) and incubated overnight with either p50 or p65 mouse monoclonal antibody (1:200) at 4°C . The membrane was washed again and incubated with anti-mouse IgG horseradish peroxidase conjugate secondary antibody for 3–4 h at room temperature. After incubation and repeated washing, the expression of p50 and p65 was visualized using chemiluminescent reagent (GE Healthcare) following the manufacturer's instructions. The membrane picture was taken with a Kodak Image station 440 CF (Eastman Kodak Co.).

Statistical Analysis. Data are presented as means \pm SD for the indicated number of independently performed experiments. Data were

analyzed using one-way ANOVA, and means were considered to be significantly different at $P < 0.05$ as determined by least significant differences (LSD).

RESULTS

Effect of Different Mate Tea Phytochemicals on Viability of RAW 264.7 Cells. Cells showed a survival rate of $> 80\%$ when incubated with MT extract (92.8%), decaffeinated MT extract (93.5%), CHA (91.2%), matein (91.2%), and mate saponins (82.7%), at a concentration $\leq 300\ \mu\text{M}$. Therefore, in this study, a concentration of 1– $300\ \mu\text{M}$ was used to treat the cells and thus prevent the compounds from having a cytotoxic effect. Concentrations ranging from 1 to $200\ \mu\text{M}$, from 1 to $100\ \mu\text{M}$, and from 1 to $100\ \mu\text{M}$ were used for quercetin, ursolic acid, and oleanolic acid, respectively, because their survival rates were 96.4, 80.8, and 87.6%, respectively.

Table 1. IC₁₀, IC₃₅, and IC₅₀ of Mate Tea Compounds against NO and iNOS Production and Expression^a

compound	NO ^b (μM)			iNOS ^b (μM)		
	IC ₁₀	IC ₃₅	IC ₅₀	IC ₁₀	IC ₃₅	IC ₅₀
mate tea extract	>300	>300	>300	16.5 ± 10.2	198.0 ± 10.2	>300
decaffeinated mate tea extract	>300	>300	>300	24.8 ± 3.5	208.0 ± 3.5	>300
chlorogenic acid	123.4 ± 1.6	>300	>300	198.8 ± 0.8	250.8 ± 0.8	>300
matein	>300	>300	>300	147.5 ± 0.3	203.0 ± 0.3	223.7 ± 0.3
quercetin	0.5 ± 0.3	1.5 ± 0.3	11.6 ± 0.3	1.1 ± 0.1	4.5 ± 0.1	7.9 ± 0.1
mate saponins	4.8 ± 0.5	10.1 ± 0.5	20.8 ± 0.5	2.3 ± 0.5	6.2 ± 0.5	8.4 ± 0.5
ursolic acid	101.5 ± 1.6	>100	>100	81.2 ± 0.5	>100	>100
oleanolic acid	16.7 ± 5.4	37.8 ± 5.4	79.7 ± 5.4	8.3 ± 0.5	22.6 ± 0.5	30.9 ± 0.5

^a IC₁₀, IC₃₅, and IC₅₀ are the concentrations (μM) that resulted in 10, 35, and 50% reduction, respectively, of production/expression of pro-inflammatory responses (mean ± SD, *n* = 2). ^b Abbreviations: NO, nitric oxide; iNOS, inducible nitric oxide synthase.

Table 2. IC₁₀, IC₃₅, and IC₅₀ of Mate Tea Compounds against PGE₂ and COX-2 Production and Expression^a

compound	PGE ₂ ^b (μM)			COX-2 ^b (μM)		
	IC ₁₀	IC ₃₅	IC ₅₀	IC ₁₀	IC ₃₅	IC ₅₀
mate tea extract	173.6 ± 2.0	220.0 ± 2.0	280.6 ± 2.0	111.8 ± 0.2	206.3 ± 0.2	>300
decaffeinated mate tea extract	225.0 ± 3.2	>300	>300	205.5 ± 2.5	>300	>300
chlorogenic acid	100.1 ± 2.4	130.6 ± 2.4	231.6 ± 2.4	186.4 ± 0.2	>300	>300
matein	137.5 ± 0.5	215.6 ± 0.5	252.6 ± 0.5	213.7 ± 0.5	>300	>300
quercetin	0.5 ± 2.2	1.5 ± 2.2	6.5 ± 2.2	52.2 ± 0.2	179.9 ± 0.2	>200
mate saponins	122.7 ± 16.3	208.7 ± 16.3	227.1 ± 16.3	153.3 ± 0.4	209.6 ± 0.4	227.7 ± 0.4
ursolic acid	74.2 ± 12.5	>100	>100	>100	>100	>100
oleanolic acid	>100	>100	>100	>100	>100	>100

^a IC₁₀, IC₃₅, and IC₅₀ are the concentrations (μM) that resulted in 10, 35, and 50% reduction, respectively, of production/expression of pro-inflammatory responses (mean ± SD, *n* = 2). ^b Abbreviations: PGE₂, prostaglandin E₂; COX-2, cyclooxygenase-2.

Effect of Different Mate Tea Phytochemicals on NO Production and iNOS Expression by LPS-Induced RAW 264.7 Cells. Figure 2 shows the effect of different concentrations of MT phytochemicals on NO production (A1, B1, C1) and iNOS expression (A2, B2, C2) by LPS-stimulated RAW 264.7 cells.

MT extract, decaffeinated MT extract, and matein did not show any effect on NO production (Figure 2A1) but inhibited iNOS expression (Figure 2A2) at the conditions tested. Mate saponins and oleanolic acid significantly decreased NO production at 10 and 1 μM, respectively (Figure 2B1). Mate saponins and oleanolic acid significantly inhibited iNOS expression (80%) at 1 and 100 μM, respectively (Figure 2B2). Ursolic acid did not present a significant inhibition of NO and did present an extremely low inhibition for iNOS. Quercetin significantly inhibited NO production, in a dose-dependent manner, starting at a very low concentration (1 μM) (Figure 2C1). CHA had no effect in reducing NO production. Quercetin significantly inhibited iNOS expression (60%) at 10 μM, whereas CHA inhibited only 30% at a higher concentration (200 μM) (Figure 2C2).

Effect of Different Mate Tea Phytochemicals on PGE₂ Production and COX-2 Expression by LPS-Induced RAW 264.7 Cells. Figure 3 shows the effect of different concentrations of MT phytochemicals on PGE₂ production (A1, B1, C1) and COX-2 expression (A2, B2, C2) by RAW 264.7 macrophages induced by LPS.

MT extract (> 50%), decaffeinated MT extract (20%), and matein (50%) inhibited PGE₂ production at 200, 300, and 200 μM, respectively (Figure 3A1). MT extract significantly inhibited COX-2 expression (30%) at 100 μM, but decaffeinated MT extract (30%) and matein (25%) inhibited at concentration of 300 μM (Figure 3A2). Ursolic acid significantly decreased PGE₂ production (20%) at 10 μM, whereas mate saponins (90%) and oleanolic acid (10%) showed inhibition at higher concentrations (300 and 100 μM, respectively) (Figure 3B1). Mate saponins (50%), ursolic acid (20%), and oleanolic acid (40%) inhibited

COX-2 expression at concentrations of 200, 100, and 100 μM, respectively (Figure 3B2). Quercetin significantly inhibited PGE₂ production (80% at 10 μM) (Figure 3C1) and COX-2 expression (25% at 10 μM), whereas CHA showed PGE₂ inhibition (50%) at higher concentration (200 μM) (Figure 3C1).

Tables 1 and 2 present the concentrations (μM) of each MT phytochemical that resulted in 10, 35, and 50% reductions against pro-inflammatory parameters. Quercetin was the most potent inhibitor of pro-inflammatory responses at a concentration 10 times lower than the concentrations used for the other compounds (IC₅₀ = 11.6 μM for NO, 7.9 μM for iNOS, and 6.5 μM for PGE₂). Mate saponins had a high capacity in preventing inflammation through the NO/iNOS pathway (IC₅₀ = 20.8 μM for NO and 8.4 μM for iNOS). Among the eight MT phytochemicals studied, quercetin and mate saponins exhibited the lowest IC₅₀ values for inhibition of iNOS, NO, and PGE₂ production, indicating that they are the most potent phytochemicals in MT in preventing inflammation.

Interaction Study and Isobolographic Analysis. Figures 4 and 5 show the effect of combined MT phytochemicals on the production of NO (A1, B1, C1) and PGE₂ (A2, B2, C2) by LPS-induced RAW 264.7. More than additive effect was observed in the inhibition of NO production when CHA and quercetin were combined (Figure 4A1). Additive effects in inhibiting PGE₂ production were observed in the combination of CHA/quercetin (Figure 4A2). Combinations of CHA/mate saponins (Figure 4B1, B2) and CHA/matein (Figure 4C1, C2) resulted in an antagonistic effect in inhibiting NO and PGE₂ production. Combinations of quercetin/matein (Figure 5A1) resulted in an antagonistic effect in inhibiting NO production but resulted in an additive effect in PGE₂ production (Figure 5A2). Combinations of quercetin/mate saponins (Figure 5 B1, B2) resulted in synergistic effects in inhibiting both NO and PGE₂ production. More than an additive effect was observed in the inhibition of NO production when mate saponins/matein were combined (Figure 5C1). A slight decrease

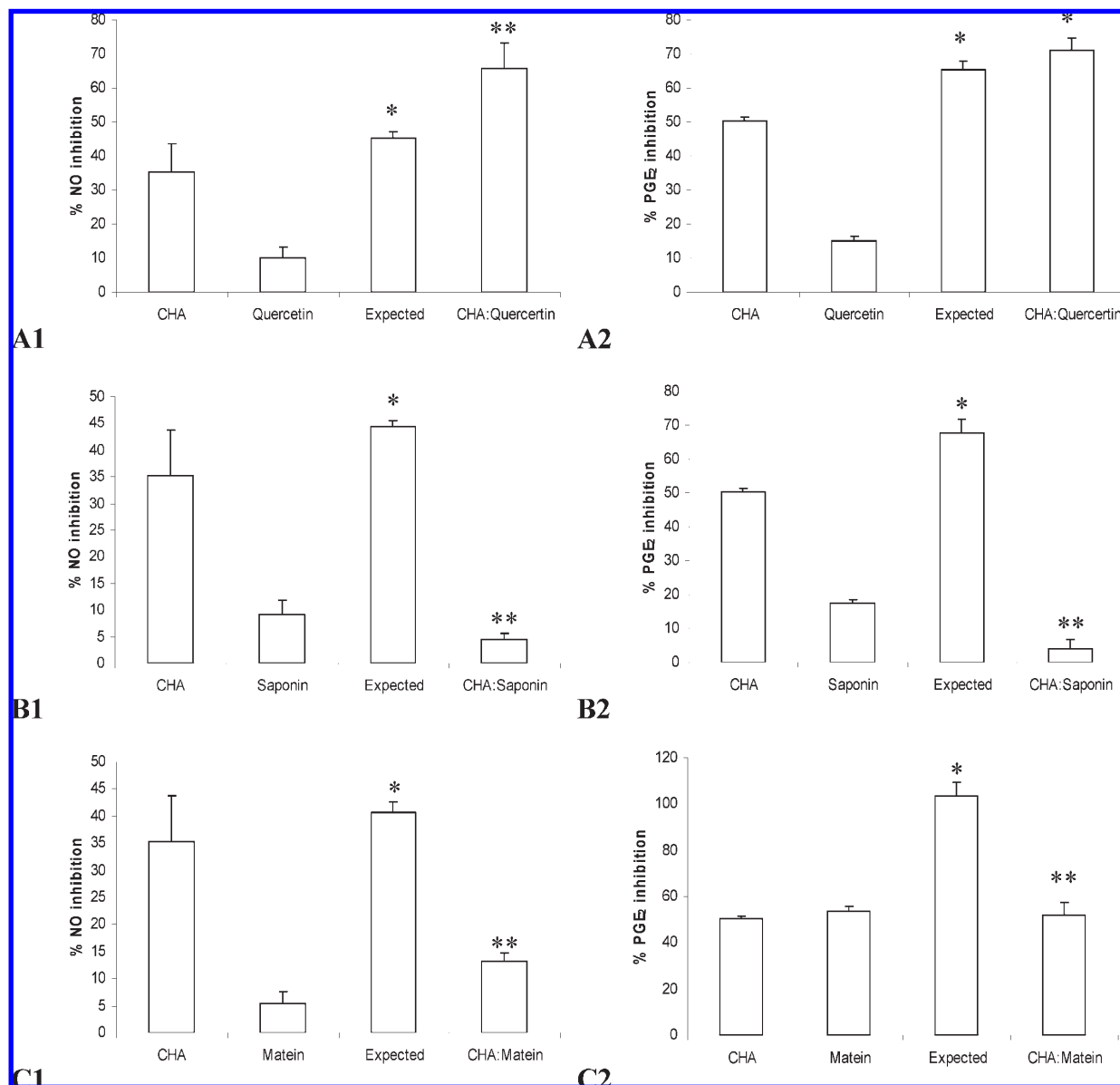


Figure 4. Inhibition (%) of NO (1) and PGE₂ (2) production by LPS-induced RAW 264.7 macrophages after treatment with (A1, A2) CHA (200 μ M), quercetin (1 μ M), and combination of CHA/quercetin (2.3:0.001); (B1, B2) CHA (200 μ M), saponins (4 μ M), and combination of CHA/saponins (2.3:0.004); and (C1, C2) CHA (200 μ M), matein (200 μ M), and combination of CHA/matein (2.3:0.86). The data represent the mean \pm SD from two independent experiments and at least a triplicate analysis. Significant differences between observed values and additive interaction are reported at $P < 0.05$.

between the combinations of mate saponins/matein in inhibiting PGE₂ production was also shown but was not statistically significant (Figure 5C2). Interactions were confirmed by isobolographic analysis as shown in Figure 6. Quercetin and mate saponins synergistically interacted to inhibit NO (Figure 6A1) and PGE₂ (Figure 6A2), whereas CHA/mate saponins (Figure 6B1) and CHA/matein (Figure 6B2) presented an antagonistic interaction.

Effect of Mate Saponins and the Combination of Quercetin and Mate Saponins on IL-1 β and IL-6 Protein Levels and on the Transcriptional Activity of NF κ B and NF κ B Nuclear Translocation by LPS-Induced Macrophages. Mate saponins and the combination of quercetin and mate saponins significantly inhibited IL-1 β (Figure 7A) and IL-6 (Figure 7B). Transient transfection experiments using the NF κ B vector indicated that quercetin and mate saponins inhibited the transcriptional activity of NF κ B but that the combination of quercetin and mate saponins showed no statistical differences by measuring firefly luciferase activity (Figure 8A). Quercetin, mate saponins, and their combination

significantly inhibited nucleus translocation of the NF κ B p65 subunit (Figure 8B). Although quercetin and mate saponins did not statistically inhibit the nucleus translocation of NF κ B p50 subunit (Figure 8C), their combination significantly inhibited the nucleus translocation of this subunit.

DISCUSSION

MT has been reported to have several biological functions, mostly related to antioxidant activity (6, 7, 21–26), interfering with glucose absorption, which might help lower the risk for type 2 diabetes (27) and obesity (28). Although a recent paper has shown that MT could be useful as an anti-inflammatory agent in lungs damaged by cigarette smoke exposure (29), the anti-inflammation mechanism of MT has not been studied yet. To our knowledge, this is the first study that evaluated and compared the inflammatory properties of different MT phytochemicals and investigated the potential interactions between phytochemicals using iNOS/NO and COX-2/PGE₂ mechanistic pathways.

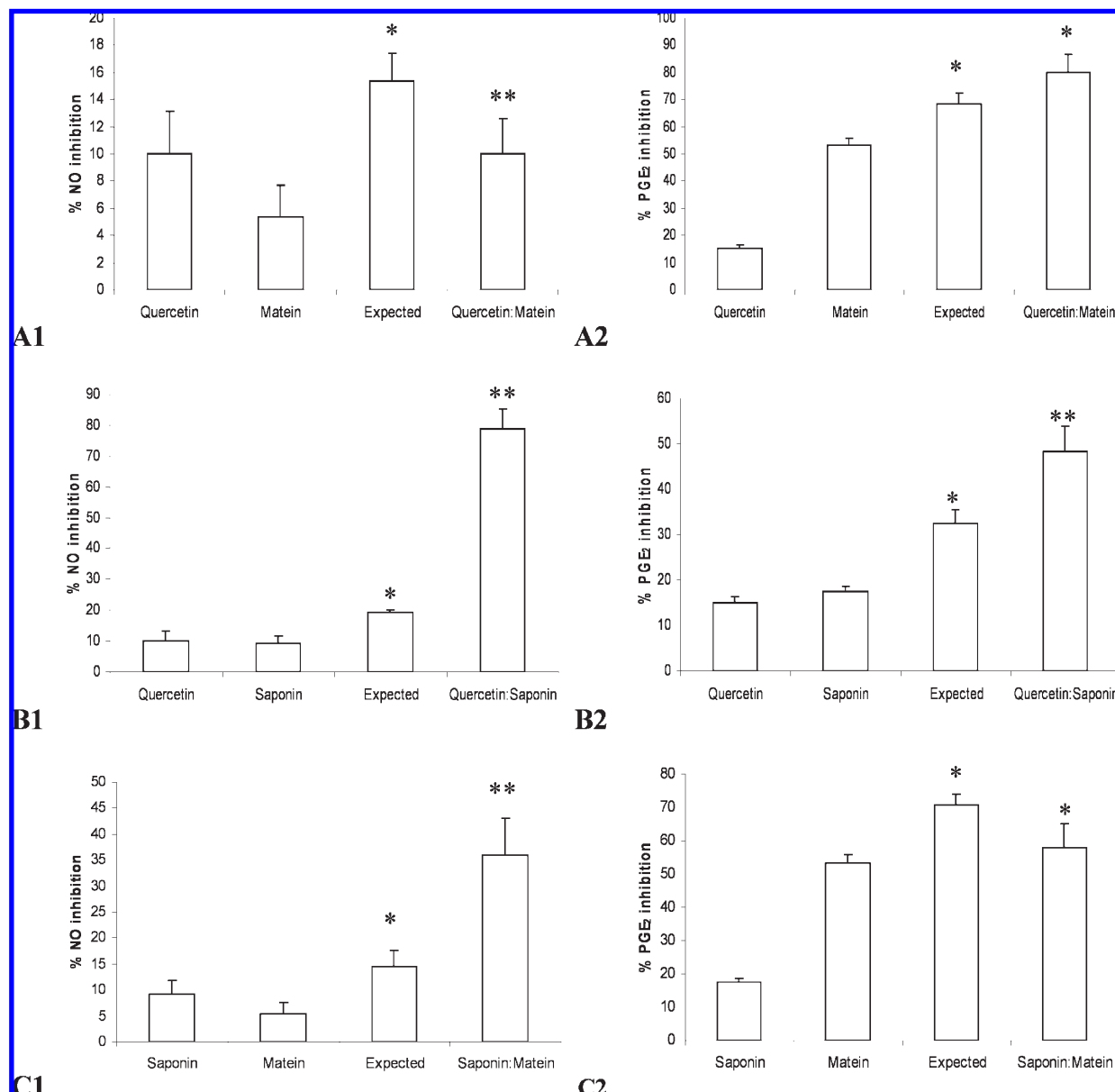


Figure 5. Inhibition (%) of NO (1) and PGE₂ (2) production by LPS-induced RAW 264.7 macrophages after treatment with (A1, A2) quercetin (1 μ M), matein (200 μ M), and combination of quercetin/matein (0.001:0.86); (B1, B2) quercetin (1 μ M), saponins (4 μ M), and combination of quercetin/saponins (0.001:0.004); and (C1, C2) saponins (4 μ M), matein (200 μ M), and combination of saponins/matein (0.004:0.86). The data represent the mean \pm SD from two independent experiments and at least a triplicate analysis. Significant differences between observed values and additive interaction are reported at $P < 0.05$.

The results suggest that different MT phytochemicals have different capacities in inhibiting inflammation through iNOS/NO and COX-2/PGE₂ pathways. MT extract, decaffeinated MT extract, and matein showed no reduction of NO production and a slight inhibition of iNOS expression in a concentration-dependent manner. The observed slight inhibition in iNOS expression showed that these compounds are capable of inhibiting the translation of the enzyme. However, the amount of NO produced was not significantly inhibited by these compounds, which might be attributed to their incapability of inhibiting the enzyme activity. Even though the expression of iNOS was slightly inhibited, the enzyme was still present and as such NO was still produced. In addition, matein did not show a potent anti-inflammatory effect. Thus, the inhibitory effect on pro-inflammatory responses by MT may not be attributed to matein. Recently, caffeine (5–50 mg/kg) did not show an inhibitory effect on PGE₂ synthesis in inflamed tissue of male Sprague–Dawley rats (30).

MT contains a significant amount of triterpenoid saponins, which exist as ursolic and oleanolic acid (19). Our results showed that mate saponins and ursolic acid inhibited inflammation through the iNOS/NO pathway more than through the COX-2/PGE₂ pathway. Sur et al. (14) found that two saponins (10 mg/kg) isolated from aqueous alcohol extract of tea root had an anti-inflammatory effect by inhibiting carrageenan-induced paw edema in rats. Crude saponin extracts and purified saponin extracts (100 mg/kg) of *Hedera colchica* were shown to be potent inhibitors of acute and chronic inflammation in carrageenan- and cotton pellet-induced inflammation models in rats (31). It has been reported that ursolic acid (100 μ M) suppressed the expression of LPS-induced pro-inflammatory mediators in RAW 264.7 macrophages that resulted from inhibition of NF κ B activation (17, 32). Although ursolic acid is known to have numerous pharmacological activities, the potency of ursolic acid is relatively weak (32), an observation in accordance with our results.

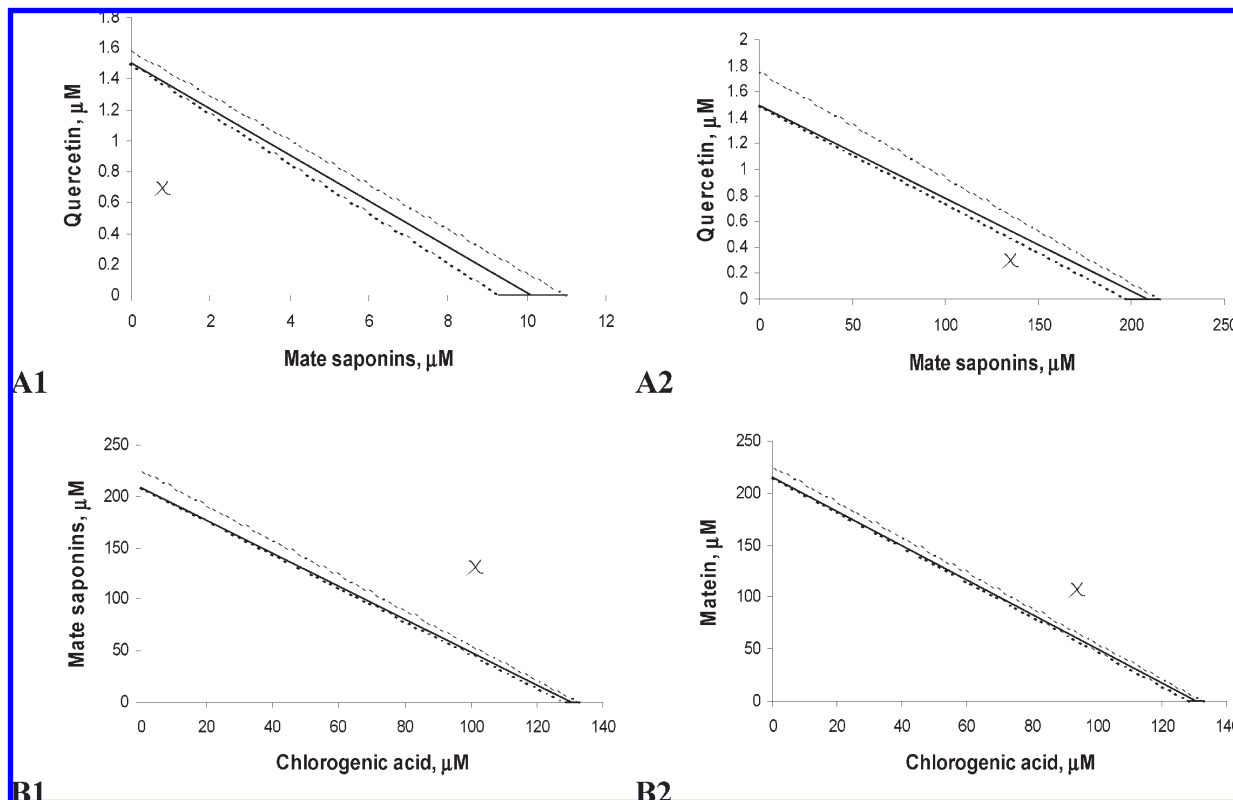


Figure 6. IC_{35} isobolograms for the synergistic interaction of (A) quercetin and mate saponins in inhibiting NO (1) and PGE_2 (2) production and for the antagonistic interaction of (B1) CHA and mate saponins and (B2) CHA and matein in inhibiting PGE_2 production by LPS-induced RAW 264.7 cells. The line of additivity (solid line) is shown with its 95% confidence interval (dotted lines).

CHA had a weak inhibition on pro-inflammatory markers in RAW 264.7 cells, even though it showed a very strong antioxidant capacity *in vitro*. These results are in agreement with those of Wang and Mazza (33) using the same model system. We found that CHA inhibited only the COX-2/ PGE_2 pathway; this is consistent with the findings of Shan et al. (34). COX-2 can be activated by both AP-1 and $NF\kappa B$, but iNOS is activated only by $NF\kappa B$ and negatively regulated by AP-1. This may explain why CHA did not alter LPS-induced iNOS expression.

Quercetin revealed a very potent anti-inflammatory effect via inhibition of NO/iNOS pathway. This finding is in accordance with Mu et al. (10), who showed that quercetin (10 μM) significantly reduced LPS-induced RAW 264.7 macrophage NO production in a concentration-dependent manner through down-regulating the expression of iNOS. Furthermore, Liang et al. (11) found that quercetin (12.5 μM) slightly enhanced the COX-2 protein level, but significantly inhibited iNOS protein level expression in LPS-induced macrophages. They suggested that the two OH groups on the B ring (Figure 1) decrease the inhibitory potency of quercetin on COX-2 activity (11). Our results showed that quercetin had less capacity to inhibit COX-2 expression. However, quercetin has been shown to suppress COX-2 expression in J774A.1 macrophages (50 μM) (12) and in rat aberrant crypt foci formation (4.5 g/kg) (13).

The combined concentrations of MT phytochemicals used in this study were equivalent to the concentrations in one cup of tea (1.5 g of DL/150 mL of water). We found that the mixture of quercetin/mate saponins exhibited the highest potency in inhibiting pro-inflammatory responses, among other combinations tested. The presence of quercetin in the mixture resulted in higher potency as compared to individual MT phytochemicals, suggesting a possible synergistic and/or additive effect. On the other

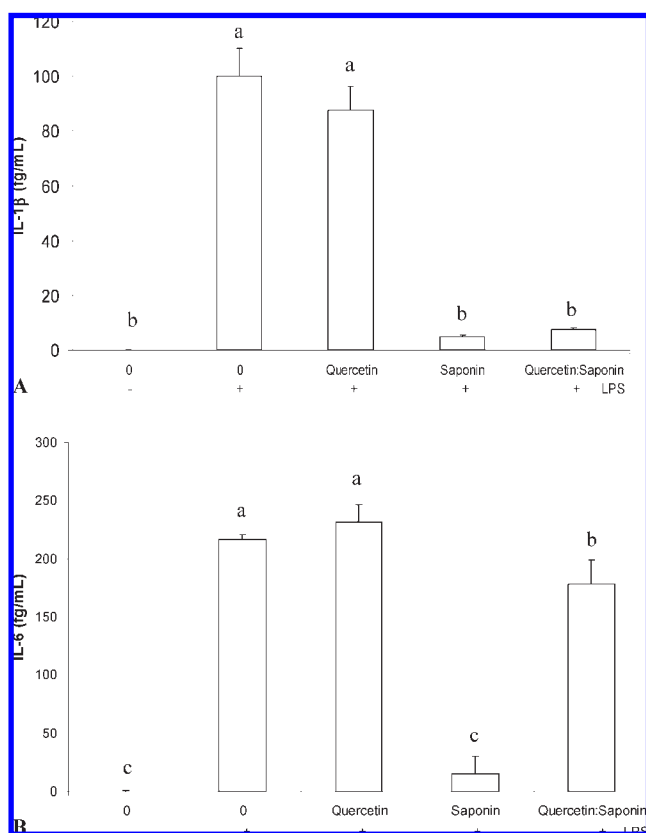


Figure 7. Effect of quercetin, mate saponins, and their combination on (A) IL-1 β and (B) IL-6 production by LPS-induced RAW 264.7 macrophages. The data represent the mean \pm SD from two independent experiments. Different letters indicate significant differences, $P < 0.05$.

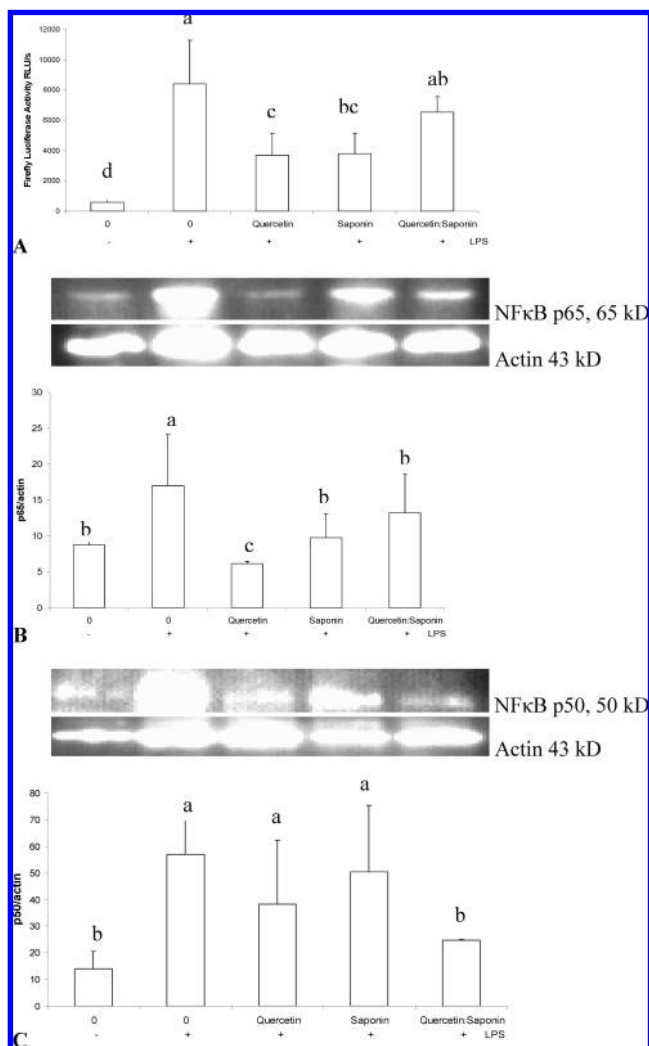


Figure 8. Effect of quercetin, mate saponins, and their combination on NF κ B transactivation by LPS-induced RAW 264.7 macrophages as measured by luciferase assay (A) and effect of quercetin, mate saponins, and their combination on (B) p65 nuclear translocation and (C) p50 nuclear translocation by LPS-induced RAW 264.7 macrophages. The data represent the mean \pm SD from two independent experiments. Different letters indicate significant differences, $P < 0.05$.

hand, combination of CHA/mate saponins and CHA/matein resulted in an antagonistic inhibition of NO and PGE₂ production. The MT extract not having a potent anti-inflammatory effect may be due to the antagonistic effect of some of its compounds. However, whole MT consumption still has a promising anti-inflammatory effect mainly through the PGE₂/COX-2 pathway.

The data suggest that the anti-inflammatory effect of MT might be due to the presence of quercetin and mate saponins even though their concentration in MT is low (35). The synergistic or antagonistic effects of the mixtures may depend on the chemical structures of the molecules and the possible formation of stable intermolecular complexes (36). These interactions could be due to attachment between the aromatic ring of phenolic acid and the B ring of flavonol, and hydrogen-bonding effects would also help to form the complex. Further studies in vivo are important to determine the bioavailability and the potential adverse effects (37).

Mate saponins and the combination of quercetin and mate saponins had a strong anti-inflammatory activity due to their

ability to inhibit the overexpression of proinflammatory cytokine IL-1 β , suggesting inhibition in the activation IKK, the enzyme responsible for the phosphorylation of I κ B protein kinase. In addition, quercetin, mate saponins, and their combination inhibited the nuclear translocation of NF κ B p65 in accordance with the decrease in NO and PGE₂ and reduced production of IL-6 by saponins and the mixture.

Quercetin and mate saponins reduced significantly ($P < 0.05$) the activation of NF κ B in transiently transfected RAW 264.7 macrophages, suggesting a reduction in the binding of NF κ B to its target DNA, so inhibiting transcription of genes with pro-inflammatory properties. Furthermore, the combination significantly inhibited nuclear translocation of NF κ B p50.

In summary, our results suggest that quercetin and mate saponins are the most potent inflammation inhibitors in MT and, when combined, they have a synergistic effect. The quercetin and saponin combination, at a concentration found in one cup of MT, inhibited inflammation through suppression of NF κ B pathways by inhibiting production of IL-1 β , thereby suggesting prevention in the activation of I κ B kinase, by blocking the translocation of p50 and p65 NF κ B subunits resulting in the decreased production of IL-6, PGE₂, and NO. To our knowledge this is the first study demonstrating the efficacy and interactions of phytochemicals in MT inhibiting pro-inflammatory markers.

ABBREVIATIONS USED

CHA, chlorogenic acid; COX-2, cyclooxygenase-2; DL, dry leaves; IL-1 β , interleukin-1 β ; IL-6, interleukin-6; iNOS, inducible nitric oxide synthase; NO, nitric oxide; NF κ B, nuclear factor- κ B; PGE₂, prostaglandin E₂; MT, mate tea.

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