Genkwanin Inhibits Proinflammatory Mediators Mainly through the Regulation of miR-101/MKP-1/MAPK Pathway in LPS-Activated Macrophages

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Abstract

Genkwanin is one of the major non-glycosylated flavonoids in many herbs with anti-inflammatory activities. Although its anti-inflammatory activity in vivo has been reported, the potential molecular mechanisms remain obscure. In this study, by pharmacological and genetic approaches, we explore the anti-inflammatory effects of genkwanin in LPS-activated RAW264.7 macrophages. Genkwanin potently decreases the proinflammatory mediators, such as iNOS, TNF-α, IL-1β and IL-6, at the transcriptional and translational levels without cytotoxicity, indicating the excellent anti-inflammatory potency of genkwanin in vitro. Mechanism study shows that genkwanin significantly suppresses the p38- and JNK-mediated AP-1 signaling pathway and increases the mitogen-activated protein kinase (MAPK) phosphatase 1 (MKP-1) expression at the posttranscriptional level. We also confirmed that microRNA-101 (miR-101) is a negative regulator of MKP-1 expression. Moreover, regardless of miR-101-deficient cells or miR-101-abundant cells, the suppression effects of genkwanin on supernatant proinflammatory mediators’ levels are far less than that in respective negative control cells, suggesting that genkwanin exerts anti-inflammatory effect mainly through reducing miR-101 production. However, genkwanin can’t affect the level of phospho-Akt (p-Akt), indicating that the phosphorylation of Akt may be not responsible for the effect of genkwanin on miR-101 production. We conclude that genkwanin exerts its anti-inflammatory effect mainly through the regulation of the miR-101/MKP-1/MAPK pathway.

Introduction

Genkwanin is one of the major non-glycosylated flavonoids in some herbs which have anti-inflammatory activities, such as Genkwa Flos (Daphne Genkwa Sieb. et Zucc.) [1], rosemary (Rosmarinus officinalis L.) [2] and the leaves of Cistus laurifolius L. [3]. Previous pharmacological studies have found that genkwanin has a variety of pharmacological effects including anti-bacterial [4,5], antiplasmodial [6], radical scavenging [7], chemopreventive [8] and inhibiting 17β-Hydroxysteroid dehydrogenase type 1 [9] activities. Although Pelzer et al. [10] has reported that genkwanin could inhibit the development of cotton-pellet-induced granuloma in rat, the potential molecular mechanisms of the anti-inflammatory activity of genkwanin remain obscure.

Inflammation is a central feature of many pathophysiological conditions in response to tissue injury and host defenses against invading microbes [11]. Key events in the inflammatory process include the expression of inflammatory cytokines, chemokines, and other mediators [12]. Macrophages play a central role in host defense against pathogen microbes by recognizing bacterial components and resulting in the activation of an arsenal of antimicrobial effectors and initiation of the inflammatory cascade [13,14]. LPS, a major component of the outer membranes in Gram-negative bacteria, can be recognized by a TLR4 receptor complex [15]. Stimulation of TLR4 by LPS triggers the recruitment of adaptor protein MyD88, which in turn transmits a series of signaling cascades that lead to the activation of mitogen-activated protein kinase (MAPK) [16-19]. The MAPK is a group of highly conserved serine/threonine protein kinases, including p38, ERK1/2 and JNK. Once activated, MAPK phosphorylate downstream protein kinases and transcription factors, leading to the production of proinflammatory cytokines, such as iNOS, TNF-α, IL-1β and IL-6, etc.

The objective of the present paper is to clarify the anti-inflammatory mechanisms of genkwanin in LPS-activated RAW264.7 macrophages. Our results indicate that genkwanin suppresses the production of inflammatory mediator in LPS-activated RAW264.7 macrophages mainly through mediating microRNA-101 (miR-101)/MAPK phosphatase 1 (MKP-1)/MAPK pathway.

Materials and Methods

Materials

The murine macrophage RAW264.7 cell line was purchased from American Type Culture Collection (ATCC, Rockville, MD, USA). Genkwanin (≥98%) was purchased from Rochen Co. (Shanghai, China) and dissolved in DMSO at the concentration of...
was replaced with the medium already mixed with genkwanin. Concentrations of DMSO were performed genkwanin treatment, genkwanin was first added to the culture medium and then mixed thoroughly (the final concentration for 2 h and then exposed to LPS (10 ng/mL) for 12 h. After being washed twice by PBS, the cells were harvested and plated in a 48-well plate, and incubated in the 10 ng/mL) for 12 h. After being washed twice by PBS, the cells were harvested and plated in a 48-well plate, and incubated in the system and lipofectamineTM 2000 reagent were purchased from Colorado, USA). The plasmids for NF-κB and Akt were from Cell Signaling Technology (Danvers, Massachusetts). The plasmids for NFκB-TA-luc, AP1-TA-luc and their controls GL6-TA were from Beyotime Institute of Biotechnology (Nanjing, Jiangsu, China). The luciferase assay system and lipofectamineTM 2000 reagent were purchased from Promega Co. (Madison, Wisconsin, USA) and Invitrogen (New York, California, USA), respectively. Horseradish peroxidase-conjugated anti-rabbit or mouse IgG secondary antibodies were obtained from Jackson ImmunoResearch Laboratories, Inc. (Lancaster, Pennsylvania, USA). miR-101 qPCR kit with U6 snRNA (control), dsRNA mimic and ssRNA inhibitor for mmu-miR-101a were obtained from GenePharma (Shanghai, China). All other reagents were of analytical grade.

Cell culture and treatment
RAW264.7 macrophages were grown in DMEM supplemented with 10% heat-inactivated FBS and 1.0% penicillin-streptomycin solution in a humidified incubator with 5.0% CO2 at 37°C. When performing genkwanin treatment, genkwanin was first added to the culture medium and then mixed thoroughly (the final concentrations were ≤0.15%). The cell culture medium was replaced with the medium already mixed with genkwanin.

Measurement of nitrite
Cells were pretreated with genkwanin at the indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 24 h. The nitrite concentration in the medium was measured as an indicator of NO production according to the Griess reaction. 100 µL of each supernatant was mixed with the same volume of Griess reagent (1% sulfanilamide in 5% phosphoric acid and 0.1% N-1-naphthylethylenediamine dihydrochloride in water). The absorbance was measured at a wavelength of 540 nm after incubation for 10 min. The nitrite concentration was calculated with reference to a standard curve of sodium nitrite generated with L-NAME was used as a positive control.

Measurement of iNOS enzyme activity
The activity of iNOS was assayed as previously described [20] with slight modifications. The cells were plated in a 25 cm² culture flask and incubated with LPS (Sigma, Escherichia coli 055: B5; 10 ng/mL) for 12 h. After being washed twice by PBS, the cells were harvested and plated in a 48-well plate, and incubated in the presence or absence of genkwanin at different concentrations for a further 12 h. The iNOS activity was assayed by measuring the nitrite level in the supernatant by Griess method. L-NAME was used as a positive control.

ELISA for TNF-α, IL-1β and IL-6
For the measurements of TNF-α, IL-6 and IL-1β, RAW264.7 macrophages were pretreated with genkwanin for 2 h and then stimulated with LPS (10 ng/mL) for 24 h. TNF-α, IL-6 and IL-1β in the cell supernatants were assayed using ELISA kits according to the manufacturer’s instructions. The concentrations were calculated from the standard curves.

Quantitative real-time PCR (RT-qPCR)
The cells were pretreated with genkwanin at the indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 4 h. The RNA extraction and RT-qPCR assays for the mRNA levels of iNOS, TNF-α, IL-1β and IL-6 were performed as we previously described [21]. RT-qPCR assay for the miR-101 level was performed according to the manufacturer’s protocol. The cycling conditions were as follows: 95°C for 3 min, and then 40 cycles of 95°C for 12 s, 60°C for 40 s. The levels of iNOS, TNF-α, IL-1β and IL-6 were normalized to β-actin. The level of miR-101 was normalized to U6 snRNA.

Transfection of plasmids, dsRNA mimic and ssRNA inhibitor for mmu-miR-101a
The sense strand and the antisense strand of dsRNA mimic for mmu-miR-101a were 5'-UAC AGU GUG AUU ACU ACU GAA-3' and 5'-GAG UUA UCA CAG UAC UGU AUU-3', respectively. The strand of ssRNA inhibitor against mmu-miR-101a was 5'-UUC AGU UAU CAC AGU ACU GUA-3'. Plasmids for pNFκB-TA-luc, pAP1-TA-luc and their controls pGL6-TA, and RNAs were transfigured into RAW264.7 macrophages using lipofectamineTM 2000 as described in the manufacturer’s protocol.

Luciferase reporter assay
RAW264.7 macrophages were transfigured with the luciferase reporter pAP1-TA-luc (A) and pNFκB-TA-luc (B). 24 h after transfection, the cells were pretreated with different concentrations of genkwanin for 2 h and then exposed to LPS (10 ng/mL). After 6 h, the cells were lysed and luciferase activity was measured using the Luciferase Assay System.

Western blot analysis
RAW264.7 macrophages were pretreated with genkwanin at indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 1 h (for p-p38, p-JNK, p-ERK1/2, MKP-1 and p-Akt assays) or 24 h (for iNOS assay). The western blot assay was performed as we previously described [21]. Proteins in nucleus or cytoplasm were extracted and separated by SDS-PAGE and transferred to polyvinylidene difluoride membranes. The membranes were blocked at room temperature for 4 h with 5.0% nonfat dry milk, and then incubated with each primary antibody at 4°C overnight. After washing, the membranes were incubated with HRP-conjugated secondary antibodies for 2 h at room temperature. The blots were visualized using enhanced chemiluminescence, and data were analysed using the Gel Doc EQ System (Bio-Rad).

Statistical analysis
Data represent the mean ± SD of at least three independent experiments, each experiment was performed in triplicate. One-way ANOVA was used to determine the statistical significance between different groups. A student t-test was used when only two groups were compared. Differences were considered to be significant at p < 0.05.

Results
Genkwanin inhibits LPS-induced NO production and suppresses iNOS at the transcriptional and translational levels
Cell viability analysis showed that genkwanin did not affect the cell viability up to a concentration of 50 µM (Figure S1). NO, a
small diffusible molecular generated by iNOS in activated macrophages, is closely related to many inflammatory diseases [22–25]. Thus, to investigate the effect of genkwanin on inflammation, we first measured supernatant NO production in LPS-stimulated RAW264.7 macrophages. As shown in Figure 1A, genkwanin inhibited the LPS-induced production of NO in a concentration-dependent manner. As we known, iNOS only expresses in the present of external stimulus [26]. To assay the effect of genkwanin on iNOS enzyme activity, we pretreated the cells with LPS for 12 h and then removed LPS. In a 48-well plate, the re-plated cells could not produce new iNOS in the absent of LPS. Under this condition, all change of NO production was attributed to the change of iNOS enzyme activity rather than iNOS mass. As shown in Figure 1B, genkwanin could not significantly affect the activity of iNOS.

Thus, we next investigated the inhibitory effects of genkwanin on iNOS mRNA and protein levels. As shown in Figure 1C–D, LPS stimulation of RAW264.7 macrophages resulted in a dramatic increase in iNOS at the transcriptional (Figure 1C) and translational (Figure 1D) levels. Treatment with genkwanin concentration-dependently inhibited the LPS-induced increase in iNOS mRNA expression and protein levels.

Genkwanin suppresses LPS-induced TNF-α, IL-1β and IL-6 at the transcriptional and translational levels

The effect of genkwanin on the production of proinflammatory cytokines was examined. As shown in Figure 2A, genkwanin suppressed the productions of TNF-α, IL-1β and IL-6 in LPS-stimulated RAW264.7 macrophages in a concentration-dependent manner. We next analysed the effects of genkwanin on the mRNA quantities of TNF-α, IL-1β and IL-6 by RT-qPCR. As shown in Figure 2B, genkwanin consistently down-regulated the LPS-induced transcription of TNF-α, IL-1β and IL-6 mRNA in a concentration-dependent manner.

Genkwanin suppresses the LPS-induced phosphorylation of p38 and JNK via the up-regulation of MKP-1 expression

Since the induction of proinflammatory mediators by LPS is known to be predominantly regulated by NF-κB and AP-1 [27–

Figure 1. Effects of genkwanin on NO production and iNOS in LPS-activated RAW264.7 macrophages. (A) Effects of genkwanin on LPS-induced NO production. Cells were pretreated with genkwanin at the indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 24 h. After treatment, nitrite levels in the medium were measured by Griess reaction. L-NAME was used as a positive control. (B) Effects of genkwanin on iNOS enzyme activity. Cells were pretreated with LPS (10 ng/mL) for 12 h and then exposed to genkwanin at the indicated concentrations for a further 12 h without LPS. Nitrite levels in the medium were measured. L-NAME was used as a positive control. (C) Effects of genkwanin on iNOS mRNA expression. Cells were pretreated with genkwanin for 2 h and then exposed to LPS (10 ng/mL) for 4 h. mRNA of iNOS was determined by RT-qPCR analysis. (D) Effects of genkwanin on iNOS protein levels. Cells were pretreated with genkwanin at the indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 24 h. After treatment, cellular proteins were prepared and the iNOS protein levels were determined by Western blot analysis. Bars represent mean ±SD of three independent experiments. ##p<0.01 vs. normal control group; **p<0.01 vs. LPS alone.

doi:10.1371/journal.pone.0096741.g001
Figure 2. Effects of genkwanin on TNF-α, IL-1β and IL-6 in LPS-activated RAW264.7 macrophages at the transcriptional and translational levels. (A) The cells were pretreated with the indicated concentrations of genkwanin for 2 h and then exposed to LPS (10 ng/mL) for 24 h. The levels of TNF-α, IL-1β and IL-6 in the supernatant were determined by ELISA. (B) The cells were pretreated with genkwanin at the indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 4 h. The mRNA expressions of TNF-α, IL-6 and IL-1β were determined by RT-qPCR analysis. ##p<0.01 vs. normal control group; **p<0.01 vs. LPS alone. Bars represent mean ± SD of three independent experiments.

doi:10.1371/journal.pone.0096741.g002

Figure 3. Effects of genkwanin on LPS-induced AP-1 and NF-κB activities. RAW264.7 macrophages were transfected with the luciferase reporter pAP-1-TA-luc (A) and pNF-κB-TA-luc (B). 24 h after transfection, the cells were pretreated with genkwanin for 2 h and then exposed to LPS (10 ng/mL) at the indicated concentrations. After 6 h, the luciferase activity was determined. ###p<0.01 vs. normal control group; **p<0.01 vs. LPS alone. Bars represent mean ± SD of three independent experiments.

doi:10.1371/journal.pone.0096741.g003
we investigated if genkwanin exerts anti-inflammatory activities by affecting these two pathways. As shown in Figure 3, genkwanin significantly suppressed the AP-1 signaling pathway (Figure 3A) but had little effect on the NF-κB signaling pathway (Figure 3B). Thus, we next explored the MAPK, which mainly act upstream of AP-1, to determine the target of genkwanin. MAPK signal transduction pathways are classified into three components. Hence, the effects of genkwanin on LPS-induced phosphorylation of p38, ERK1/2 and JNK were investigated. The results indicate that genkwanin suppresses the phosphorylation of p38 and JNK in a concentration-dependent manner, but little affects ERK1/2 phosphorylation (Figure 4A).

Originally identified as an immediate early gene, MKP-1 was then found to be a dual specificity phosphatase acting as a negative regulator of ERK1/2, JNK and p38 MAPK activities, with predominant effects on the latter two [31–36]. Thus, we next explored the effect of genkwanin on MKP-1. As shown in Figure 4B–C, LPS stimulation induced the expression of MKP-1. As shown in Figure 4B–C, LPS stimulation induced the expression of MKP-1. Pretreatment with genkwanin markedly up-regulated the expression of MKP-1 without affecting MKP-1 mRNA.

Genkwanin exerts anti-inflammatory effects mainly through decreasing miR-101 production

Our above results have demonstrated that genkwanin up-regulates MKP-1 at the posttranscriptional level. It was previously found that MKP-1 as a target of miR-101 which can repress MKP-1 protein expression [37]. Thus, we next evaluated the effect of genkwanin on miR-101 expression by RT-qPCR. As shown in Figure 5A, LPS induced the expression of miR-101, but pretreatment with genkwanin significantly decreased miR-101. We also analysed the effect of miR-101 on the protein level of MKP-1 in LPS-activated RAW264.7 macrophages which had been transfected with dsRNA mimic or siRNA inhibitor for mmu-miR-101a (miR-101 mimic or miR-101 inhibitor). As shown in Figure 5B, transfection with miR-101 mimic significantly inhibited

![Figure 4. Effects of genkwanin on LPS-induced MAPK and MKP-1.](image-url)

(A) Effects of genkwanin on the LPS-induced phosphorylation of p38, JNK and ERK1/2. RAW264.7 macrophages were pretreated with genkwanin at indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 1 h. Total proteins were extracted for the Western blot analysis. (B) Effects of genkwanin on MKP-1 mRNA in LPS-activated RAW264.7 macrophages. The cells were pretreated with genkwanin at the indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 1 h. The mRNA expression of MKP-1 was determined by RT-qPCR analysis. (C) Effects of genkwanin on MKP-1 protein level in LPS-activated RAW264.7 macrophages. Cells were pretreated with genkwanin at indicated concentrations for 2 h and then exposed to LPS (10 ng/mL) for 1 h. Cell lysates were analysed by Western blot. **p<0.01 vs. normal control group; ***p<0.01 vs. LPS alone. Bars represent mean ±SD of three independent experiments.

doi:10.1371/journal.pone.0096741.g004
the production of MKP-1 protein, while miR-101 inhibitor markedly increased MKP-1 protein. These results are consistent with the previous reports that miR-101 is a negative regulator of MKP-1 expression [37].

To understand how genkwanin suppressed inflammation via miR-101, we transfected miR-101 inhibitor into RAW264.7 macrophages. In the resulting miR-101-deficient cells, the TNF-α production in response to LPS was significantly decreased as compared with the cells transfected with ssRNA negative control (NC) (Figure 5C). Genkwanin potently decreased LPS-induced supernatant TNF-α with an inhibition rate (IR) value of ~38% \[\text{IR\%} = \left(\frac{\text{TNF-α}_{\text{LPS}} - \text{TNF-α}_{\text{LPS+Genkwanin}}}{\text{TNF-α}_{\text{LPS}}}\right) \times 100\] in ssRNA NC cells. However, in miR-101-deficient cells, genkwanin only slightly decreased LPS-induced supernatant TNF-α with an IR value of ~10%, indicating that miR-101 played a predominant role in the anti-inflammatory activity of genkwanin.

Next, we transfected miR-101 mimic into RAW264.7 macrophages. In the resulting miR-101-abundant cells, the TNF-α production in response to LPS was significantly increased as
compared with the cells transfected with dsRNA NC (Figure 5D). Genkwanin significantly decreased LPS-induced TNF-α with an IR value of ~36% in dsRNA NC cells, while in miR-101-abundant cells, the IR value dropped to ~7%, indicating that genkwanin was not effective against exogenous miR-101. Similar effects of miR-101 inhibitor and mimic on supernatant NO, IL-1β and IL-6 were also observed (Figure S2). Next, we evaluated the effects of genkwanin on the phosphorylation level of Akt in LPS-activated RAW264.7 macrophages. As shown in Figure 5E, LPS could induce the phosphorylation of Akt, but genkwanin did not appreciably affect the level of phospho-Akt (p-Akt).

**Discussion**

Genkwanin (4',5-dihydroxy-7-methoxyflavone), as one of the major bioactive components in Genkwa Flos, is used as a representative index for the quality control of this herb and are included in the State Pharmacopoeia Commission of the People’s Republic of China [1]. Many of its structurally similar analogues,
such as apigenin [38], acacetin [39], chrysin [40], baicalein [41], wogonin [42], luteolin [43] and velutin [44] (Figure 6), were reported to suppress proinflammatory mediators in LPS-stimulated macrophages. Analysis of the structure-activity relationships of flavones showed that the anti-inflammatory effects could be enhanced by the methoxylation of the 5- or 7-hydroxyl groups on the A-ring or non-methoxylation of the 3′-hydroxy groups on the B-ring [45]. Coincidentally, the chemical structure of genkwanin includes these dispositions, such as the 5-OCH$_3$ and 3′-OH. Indeed, our results show that genkwanin (12.5 μM - 50 μM) potently decreases LPS-induced proinflammatory mediators, such as iNOS, TNF-α, IL-1β and IL-6 at the transcriptional and translational levels in RAW264.7 macrophages without cytotoxicity (Figures 1 and 2), indicating genkwanin’s excellent anti-inflammatory potency.

Of course, due to the minor structural differences, the anti-inflammatory mechanisms of the analogues are diverse. For example, apigenin [46], baicalein [47] and luteolin [43] exert their anti-inflammatory effects mainly by inactivating NF-κB, while wogonin [48] can block JNK phosphorylation. Specially, acacetin [39] and velutin [44] can inactivate both NF-κB and MAPK. In this study, after carrying out Western blot assays by using phosphorylation antibodies, we found that genkwanin decreased the LPS-induced phospho-p38 (p-p38) and phospho-JNK (p-JNK) levels, but had no effect on phospho-ERK1/2 (p-ERK1/2) (Figure 4A).

Reversible activation MAPK requires phosphorylation on threonine and tyrosine residues of the activation domain of p38, JNK and ERK1/2. They are negatively regulated by a family of dual-specificity (threonine/tyrosine) phosphatases known as the MAPK phosphatases (MKPs) [49]. MKP-1, a stress-responsive MKP, localizes to the nucleus through its N terminus [50] and preferentially dephosphorylates activated p38 and JNK relative to ERK1/2. In LPS-stimulated mouse macrophages, MKP-1 shows a transient expression pattern with rapid induction, followed by a quick return to basal levels [51]. It is a critical negative regulator of macrophage signaling in response to inflammatory stimuli and is responsible for switching off the production of proinflammatory cytokines [51–53]. Therefore, the differential regulations of genkwanin on MAPK phosphorylation strongly suggest that the upstream regulator may be MKP-1. As expected, our results show that genkwanin increases the MKP-1 expression at the posttranscriptional level (Figure 4B–C).

MicroRNAs (miRNAs), the short (~22 nucleotides) non-coding RNAs, play a central role in the regulation of gene expression at the posttranscriptional level via an RNA interference mechanism [54]. Recently, it was found that miR-101, a tumor-related miRNA, repress MKP-1 expression by binding to the 3′ untranslated region of MKP-1 in a direct and sequence-specific manner [37]. In our study, we also found the negative regulatory effect of miR-101 on MKP-1 protein (Figure 5B). Moreover, in response to LPS, the supernatant TNF-α, NO, IL-1β and IL-6 levels of miR-101-deficient cells is decreased (Figures 5C and S2), while these levels of miR-101-abundant cells is increased (Figures 5D and S2). Based on these slight effects of genkwanin on LPS-induced TNF-α, NO, IL-1β and IL-6 in exogenous miR-101-abundant cells (Figure 5D), we infer that genkwanin up-regulates MKP-1 protein may be attributed to the decrease of miR-101 production. In LPS-stimulated miR-101-deficient cells, genkwanin still somewhat suppresses supernatant TNF-α, NO, IL-

![Figure 7. Proposed mechanism by which genkwanin exerts anti-inflammatory activity in LPS-activated RAW264.7 macrophages.](https://www.plosone.org/static/content/9/5/96741_g007.png)

The gray color indicates the targets of genkwanin.
doi:10.1371/journal.pone.0096741.g007
1β and IL-6 (Figures SC and S2), which suggests another mechanism, not depending on miR-101, remains possible. As predicted, genkwanin not only can decrease the phosphorylation level of JNK (Figure 4A), but also can directly inhibit the activity of p-JNK (Figure S3).

PI3K/Akt is known to regulate proinflammatory cytokine expression, but its exact role (positive versus negative) is controversial. Some studies have demonstrated that the PI3K/Akt pathway negatively regulates TLR-induced MAPK activation and proinflammatory cytokine production [35–58]. Other reports, however, have displayed that the PI3K/Akt implicated as a positive regulator of TLR-induced inflammatory response [59–61]. Zhu and his colleagues [37] proposed that PI3K/Akt negatively regulated the expression of MKP-1 through the induction of miR-101. However, our results indicated that genkwanin could not significantly affect the level of p-Akt (Figure 5E), suggesting that the phosphorylation of Akt may be not responsible for the effect of genkwanin on miR-101 production.

Taken together, our results demonstrate that the anti-inflammatory effect of genkwanin may be mainly attributed to the down-regulation of the LPS-induced miR-101, thus increasing the protein expression of MKP-1, which dephosphorylates p38 and JNK in RAW264.7 macrophages (Figure 7). To our knowledge, genkwanin is the first compound derived from plant source shown to exert its anti-inflammatory activities mainly through the decrease of miR-101 production.

Supporting Information
Figure S1 Effect of genkwanin on cell viability. RAW264.7 macrophages were incubated with genkwanin for 24 h and the cell viability were evaluated by MTT assay. Data represent the mean ± SD of three independent experiments. (TIF)

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Figure S2 Effects of genkwanin on supernatant NO, IL-1β and IL-6 in LPS-stimulated RAW264.7 macrophages which have been transfected with shRNA inhibitor or dsRNA mimic for mmu-miR-101a. RAW264.7 macrophages were transfected with miR-101 inhibitor or mimic or their negative controls, and then stimulated with LPS (10 ng/mL) for 24 h in the presence or absence of genkwanin (50 μM). Supernatant NO (A-B), IL-1β (C-D) and IL-6 (E-F) were measured. ##p<0.01 vs. resting cells transfected with miR-101 inhibitor or mimic; *p<0.05 vs. LPS-treated cells transfected with miR-101 inhibitor or mimic; Δp<0.01 vs. resting cells transfected with negative controls; @p<0.01 vs. LPS-treated cells transfected with negative controls. (TIF)

Figure S3 Effect of genkwanin on p-JNK activity. RAW264.7 macrophages were treated with or without LPS (10 ng/mL) for 1 h. The intracellular p-JNK was extracted and purified by immunoprecipitation. The obtained p-JNK was assayed by Western blot analysis and represented as the blots of p-c-Jun. All of the extraction and purification of p-JNK and the kinase activity assay were performed according to the manufacturer’s instructions of KinaseSTAR JNK Activity Assay Kit (BioVision, Inc., San Francisco, California, USA). **p<0.01 vs. normal control group; ***p<0.01 vs. LPS alone. Bars represent mean ±SD of three independent experiments. (TIF)

Author Contributions
Conceived and designed the experiments: YG YQ. Performed the experiments: YG FL LF RG CZ. Analyzed the data: YG YQ. Contributed reagents/materials/analysis tools: RC. Wrote the paper: YG YQ.
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