Discovery and evaluation of novel anti-inflammatory derivatives of natural bioactive curcumin

Yali Zhang1,*, Xin Jiang2,*, Kesong Peng1, Chengwei Chen3, Lili Fu1, Zhe Wang1, Jianpeng Feng1, Zhiguo Liu1,4, Huajie Zhang1, Guang Liang1, Zheer Pan3

1 Chemical Biology Research Center, School of Pharmaceutical Sciences, Wenzhou Medical University, Wenzhou, Zhejiang, People’s Republic of China; 2 The Children’s Hospital, Zhejiang University School of Medicine, Hangzhou, Zhejiang, People’s Republic of China; 3 Department of Orthopedic Surgery, The First Affiliated Hospital, Wenzhou Medical University, Wenzhou, Zhejiang, People’s Republic of China; 4 Wenzhou Undersun Biotechnology Co. Ltd., Wenzhou, Zhejiang, People’s Republic of China

*These authors contributed equally to this work

Abstract: Curcumin is a natural active product that has various pharmacological activities such as anti-inflammatory effects. Here, we report the synthesis and evaluation of 34 monocarbonyl curcumin analogs as novel anti-inflammatory agents. Among the analogs, the symmetrical heterocyclic type displayed the strongest inhibition of lipopolysaccharide (LPS)-stimulated expression of pro-inflammatory cytokines in macrophages. Analogs S1–S5 and AS29 reduced tumor necrosis factor-α (TNF-α) and interleukin-6 (IL-6) production in a dose-dependent manner and also displayed excellent stability and low cytotoxicity in vitro. In addition, analog S1 dose-dependently inhibited LPS-induced extracellular signal-regulated kinase (ERK) phosphorylation. Furthermore, analogs S1 and S4 displayed a significant protective effect on LPS-induced septic death in mouse models, with 40% and 50% survival rates, respectively. These data demonstrate that the heterocyclic monocarbonyl curcumin analogs have potential therapeutic effects in acute inflammatory diseases.

Keywords: curcumin, anti-inflammation, sepsis, macrophage, cytokine

Introduction

Inflammation is part of a complex biological response that is induced by harmful stimulation, such as burns, pathogens, and chemical irritants. Over-expressed inflammatory cytokines are involved in the pathological processes of a number of diseases.1,2 In particular, sepsis, a systemic inflammatory response syndrome characterized by cytokine storm and multi-organ failure that occurs during severe infection, is believed to be related to an exacerbated release of pro-inflammatory cytokines, such as tumor necrosis factor-α (TNF-α), interleukin-1 (IL-1), and interleukin-6 (IL-6).3,4 Excessive or sustained release of inflammatory cytokines leads to serious consequences, including septic shock.5 Cytokines also play a vital role in the development of sepsis.6–9 The development of novel therapeutic agents that block pro-inflammatory pathways or inhibit the over-expression of cytokines has been the focus of researchers investigating the treatment of inflammatory disease.10–12

Curcumin is a natural yellow product derived from the turmeric rhizome of the herb Curcuma Longa (Figure 1). Curcumin has shown to be non-toxic and exhibits various biological activities such as anti-oxidant, anti-inflammatory, anti-carcinogenic, anti-diabetic, anti-bacterial, anti-fungal, anti-viral, anti-fibrotic and anti-ulcer effects.13–17 Curcumin treatment has been observed to have both preventive and therapeutic anti-inflammatory effects in various animal models.18 In addition, more than 65 clinical trials of curcumin have finished and curcumin is proven to be safe.19 Despite the effective activity of curcumin on many cellular targets linked to a variety of diseases, low bioavailability and
weak stability have significantly limited its clinical application.\textsuperscript{20} To find new analogs with increased bioavailability and enhanced pharmacological activity, researchers have attempted the chemical modification of curcumin.\textsuperscript{21–23} Among these analogs, monocarbonyl analogs have received much attention as potential curcumin analogs, and the beta-diketone moiety, which is believed to be responsible for the weak metabolism, has been removed.\textsuperscript{24,25} Our group has adopted the strategy of synthesis of monocarbonyl curcumin analogs to overcome these limitations (Figure 1). Based on the aforementioned studies, and as a part of our ongoing research, 34 novel monocarbonyl analogs of curcumin were synthesized, and their cytokine-inhibitory and anti-sepsis effects were studied.

Materials and methods

Chemical experimental procedures

The synthesis of enamide 2 and 7

To a stirred solution of cyclopentanone or piperidine or pyrone (20 mM) and p-TSA (200 mg) in cyclohexane (20 mL) was added morpholine (30 mM), and the mixture reaction was refluxed for four hours at 90°C. Subsequently, the solvent was removed under reduced pressure to give various activated enamine intermediates 2 or 7 as yellow viscous oils, and they were used directly without further purification.

General procedure for the synthesis of AS1–AS15, AS28, and AS29

The stirred solution of 3 or 9 (2 mM) in EtOH (25 mL) was added to the various aldehydes (2 mM) at room temperature. Subsequently, HCl (gas) was bubbled into the mixture for 30 minutes, and the mixture was stirred for 3 to 5 hours at 50°C–70°C. The mixture was then poured into ice water and extracted with EtOAc (3×20 mL). The combined organic layers were washed with 10% sodium bicarbonate (NaHCO\textsubscript{3}), dried over anhydrous magnesium sulfate (MgSO\textsubscript{4}), filtered, and concentrated in vacuo. Flash chromatography of the residue over silica gel using EtOAc/petroleum as the eluent gave AS1–AS15, AS28 and AS29.

General procedure for the synthesis of AS16–AS27

Similar to the procedure described for the preparation of AS1–AS15 with the exception of using a different aldehyde, all the phenolic hydroxyls of the resulting products (1 mM) were estered by dropping the propionyl chloride (10 mM) in the presence of Et\textsubscript{3}N (0.25 mL) at 0°C, using THF (10 mL) as the solvent. The reaction mixture was stirred at room temperature overnight. Thin layer chromatography demonstrated that the reaction had been completed. After removing the solvent, the resultant residue was chromatographed over silica gel using EtOAc/petroleum, as the eluent generated the desired products AS16–AS27.

Cell line and reagents

Mouse macrophages and human normal hepatic HL-7702 cells were obtained from the American Type Culture
Collection, Manassas, VA, USA. Cell culture reagents were obtained from Gibco Life Technologies, Grand Island, NY, USA. Fetal bovine serum (obtained from HyClone, Logan, UT, USA) was heat-inactivated for thirty minutes at 65°C. Lipopolysaccharide (LPS) and chemical reagents were purchased from Sigma–Aldrich, St Louis, MO, USA. Anti-p-ERK and anti-ERK antibody was obtained from Santa Cruz Biotechnology, Santa Cruz, CA, USA; anti-p-p38, anti-p38, anti-p-JNK and anti-JNK were from Cell Signaling Technology, Danvers, MA, USA. Curcumin and compounds S1–S5, and AS1–AS29 were dissolved in dimethyl sulfoxide (DMSO) for in vitro experiments.

Animals
Male C57BL/6 mice weighing 18–22 g were obtained from the Animal Center of Wenzhou Medical University (Wenzhou, People’s Republic of China). Animals were housed at constant room temperature with a 12:12 hour light:dark cycle, fed with a standard rodent diet, and provided with sufficient water. The animals were allowed to adapt to the laboratory environment for at least 7 days before their use in experiments. Protocols involving the use of animals were approved by the Wenzhou Medical University Animal Policy and Welfare Committee.

Enzyme-linked immunosorbent assay
Mouse macrophages were incubated in Dulbecco’s Modified Eagle’s Medium (DMEM) (Gibco, Eggenstein, Germany) and replenished with 10% fetal bovine serum (FBS), 100 U/mL penicillin, and 100 mg/mL streptomycin at 37°C with 5% CO₂. Cells were pretreated with compounds for 30 minutes, then treated with LPS (0.5 μg/mL) for 24 hours. After treatment, the culture media and cells were collected separately. The levels of tumor necrosis factor alpha (TNF-α) and interleukin-6 (IL-6) in the media were determined by enzyme-linked immunosorbent assay (ELISA) using mouse TNF-α and mouse IL-6 ELISA kits (eBioScience, San Diego, CA). The total quantity of the inflammatory factor in the media was standardized to the total protein amount of the viable cell pellets.

Western blot
Macrophages were treated with 0.5 μg/mL LPS for 20 minutes before incubation with or without compounds for 30 minutes. For the preparation of total cell proteins, macrophages were washed with PBS and lysed. The proteins were separated by sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE), and were electrophoretically transferred onto nitrocellulose membrane. The membranes were first incubated with the primary antibodies, then incubated with the horseradish peroxidase (HRP)-coupled secondary antibodies. Detection was performed with enhanced chemiluminesence reagents, and the protein bands were quantified by densitometry using ImageJ image processing program (US National Institutes of Health, Washington, DC, USA).

Methyl thiazolyl tetrazolium assay
HL-7702 cells were seeded into 96-well plates at a density of 5,000 cells per well in Roswell Park Memorial Institute 1640 medium, and replenished with 5% heat-inactivated serum, 100 U/mL penicillin, and 100 μg/mL streptomycin. Cells were incubated at 37°C in 5% CO₂ for 24 hours. Cells were cultured with DMSO or 10 μM tested compounds for 24 hours before the methyl thiazolyl tetrazolium (MTT) assay. A fresh solution of MTT (5 mg/mL) prepared in phosphate buffer solution was added to each single well. The plate was then incubated in a CO₂ incubator for 4 hours, cells dissolved with 150 μL DMSO, and read for optical density at 490 nm. Viability was defined as the ratio (expressed as a percentage) of absorbance of treated cells to DMSO treated cells.

In vivo study method
Compounds were firstly dissolved with macrogol 15 hydroxystearate (a nonionic solubilizer [BASF, Ludwigshafen, Germany]) for injection with or without medium chain triglycerides (MCT), from BASF, in a water bath at 37°C. The concentration of compounds was 2 mg/mL. The concentration of solubilizer ranged from 5%–10%, and MCT 0.5%–2% in the final solution. For the vehicle, the mixture of solubilizer and MCT was prepared at 10% and 2%, respectively. Male C57BL/6 mice weighing 18–22 g were pretreated with compound S1 or S4 (10 mg/kg) in a water solution by intravenous injection 15 minutes before the intraperitoneal injection of LPS (15 mg/kg). Control animals received a similar volume (200 μL) of vehicle. Body weight change and mortality were recorded for 7 days.

The stability experiment of active analogs
Compounds were dissolved in DMSO, and the concentration was 10 mM. The stock solution was diluted 10 times, and then added 1 μL diluent into an Eppendorf Tube which was assembled with 99 μL phosphate buffered saline (PBS). After that we could obtain the miscible liquid with the concentration of 10 μM and add it into the cuvette which was read for optical density at 250–600 nm. The absorbance was read at set intervals (0, 5, 10, 15, 20, and 25 minutes).
Statistical analysis
The results are presented as means ± standard error of the means (SEMs). Student’s t-test was employed to analyze the differences between sets of data. Statistics were performed using GraphPad Prism 5.0 (GraphPad Software, La Jolla, CA, USA). P-values less than 0.05 (P<0.05) were considered indicative of significance. All experiments were repeated at least three times.

Results and discussion
Chemistry
The synthetic routes of the symmetrical and asymmetrical monocarbonyl analogs of curcumin (MACs) are shown in Figure 2. The synthesis of the symmetrical curcumin analogs S1–S5 began with the key Claisene–Schmidt condensation reaction between vanillin and corresponding heterocyclic ketone, including piperidone, pyrone and thiopyrone. All of the phenolic hydroxyls of the products were then esterified with propionyl or isobutyryl chloride in the presence of Et₃N. The asymmetrical monocarbonyl curcumin derivatives (AS1–AS29) were synthesized also through the aldol condensation strategy. Commercially available cyclopentanone 3 or heterocyclic ketone 4 were activated with morpholine, and the resulting enamines 2 and 7 were condensed with vanillin and 2-chlorobenzaldehyde, respectively, to afford the desired unilateral unsaturated ketones. Subsequently, AS1–AS15, AS28, and AS29 were obtained by constructing the second unsaturated ketone in an acidic environment. Finally, propionylation of the phenolic hydroxyls was carried out in the presence of Et₃N to complete the

![Chemical structure diagram](Image)

**Figure 2** The synthetic pathway for monocarbonyl curcumin analogs S1–S5, and AS1–AS29.

**Note:** Reagents and conditions were a) morpholine, p-TSA, cyclohexane, reflux, 50%; b) vanillin, EtOH, reflux, HCl, 30%; c) differently aldehydes, HCl gas, EtOH, 50°C–70°C, 10%–70%; d) THF, propionyl chloride, Et₃N, rt, 5%–54%; e) piperidone, pyrone, or thiopyrone, HCl gas, EtOH, rt, 40%–50%; f) THF, propionyl, or isobutyryl chloride, Et₃N, rt, 8%–31%; g) piperidone or pyrone, p-TSA, cyclohexane, reflux, 40%–50%; h) 2-chlorodebenzaldehyde, EtOH, reflux, HCl, 30%–35%; i) 3,4-dihydroxybenzaldehyde or 3,4,5-trimethoxybenzaldehyde, HCl gas, EtOH, rt, 10%–15%.

**Abbreviations:** AS, asymmetrical monocarbonyl curcumin derivative; EtOH, ethanol; Et₃N, triethylamine; Me, methyl; rt, room temperature; THF, tetrahydrofuran; TSA, toluenesulfonic acid; HCl, hydrochloric acid.
synthesis of AS16–AS27. All compounds were purified by recrystallization or column chromatography. Their structures, determined by spectral data from electrospray mass spectrometry (ESI-MS) and proton nuclear magnetic resonance (1H-NMR) spectroscopy, are shown in Figure 2. Spectral data can be found in the Supplementary material. Before biological experiments were carried out, the purity of all compounds was detected by high-performance liquid chromatography (HPLC), as shown in the Supplementary material.

Compounds inhibited LPS-induced TNF-α and IL-6 production

TNF-α and IL-6 are two well-known pro-inflammatory cytokines. It has been well demonstrated that they play important roles in the pathological development of many inflammatory diseases.26,27 To investigate whether the 34 synthetic analogs have anti-inflammatory activity, their ability to reduce LPS-stimulated TNF-α and IL-6 production was determined in macrophages. After pretreatment with compounds for 30 minutes, macrophages were treated with LPS for 24 hours. The levels of cytokines in the culture media were determined by ELISA. The results of the anti-inflammatory assay of two series of monocarbonyl curcumin analogs are shown in Figure 3. Initial screening showed that most of the analogs attenuated the LPS-induced TNF-α and IL-6 secretion at a dosage of 10 μM. Several of the synthetic analogs exhibited a more potent inhibitory ability than curcumin. In particular, the heterocyclic analogs (S1–S5, and AS29) showed stronger inhibition than the others, with a range of 59.5%–83.4%. Compound S5, which has a thiopyrone skeleton, displayed the greatest activity, with an inhibition ratio of 98.7%.

![Graph A: TNF-α inhibition](image)

![Graph B: IL-6 inhibition](image)

**Figure 3** Curcumin and curcumin analog inhibition of LPS-induced secretion of (A) TNF-α and (B) IL-6 in mouse macrophages. Notes: Macrophages were seeded on a plate with the density of 4.0×10^5/mL overnight at 37°C and 5% CO₂. Macrophages were pre-incubated with the vehicles, curcumin or curcumin analogs (10 μM) for 2 hours, followed by stimulation with LPS (0.5 μg/mL) for an additional 22 hours. The levels of TNF-α and IL-6 in culture medium were detected by ELISA and were standardized by the total protein. The results were indicated as the proportion of LPS control. Each bar expresses mean ± SEM of three experiments. Statistical difference with regard to LPS was demonstrated: *P<0.05, **P<0.01.

Abbreviations: LPS, lipopolysaccharide; IL-6, interleukin-6; DMSO, dimethyl sulfoxide; CUR, curcumin; TNF-α, tumor necrosis factor-α; ELISA, enzyme-linked immunosorbent assay; SEM, standard error of the mean.
for the cyclopenta-containing monocarbonyl analogs, compound AS12, which has a heterocyclic-methylthiophen R₂ substitution-group, also showed an inhibition ratio over 50% on IL-6 production. Of the asymmetrical analogs, only the 4-piperidone derivative AS29 exhibited a strong inhibitory effect. Therefore, the design of novel monocarbonyl curcumin analogs with a symmetric and heterocyclic scaffold may improve the anti-inflammatory activity of this group of compounds.

Quantitative structure–activity relationship study

A quantitative structure–activity relationship (QSAR) study was carried out to demonstrate the effects of various substituents on the anti-inflammatory activity of the analogs. To obtain a QSAR model, the molecular structures of all the curcumin analogs were built with Maestro Version 9.1 (Schrödinger, New York, NY, USA) and represented by a series of molecular descriptors. All calculations were based on the semi-empirical parameterized model 6 (PM6) method. A detailed description of the QSAR study is presented in the Supplementary material. Scatter plots showing predicted activities versus experimental values are displayed in Figure 4. Statistically significant models, Equation 1 (Eq1) and Equation 2 (Eq2), containing three different variables, were obtained for the anti-TNF-α and anti-IL-6 activities of the compounds, respectively. Both models possessed relatively high regression coefficients: $R^2=0.86$ for TNF-α inhibition and $R^2=0.89$ for IL-6 inhibition. To further estimate the QSAR model, leave-one-out (LOO) cross-validation was performed. The Q-squares from LOO validation were 0.85 and 0.88 for TNF-α inhibition model and IL-6 inhibition model, respectively. LOO cross-validation manifested the final models as robust and satisfactory. The high-quality QSAR results indicated that the anti-inflammatory activities of curcumin analogs are highly correlated with their chemical structures. Comprehensive consideration of the structure–activity relationship (SAR) and QSAR results showed that molecular symmetry and electronegativity may play a crucial role in the anti-inflammatory activity of these monocarbonyl curcumin analogs.

Active compounds dose-dependently inhibited LPS-induced cytokines production

The dose-dependent inhibitory effects of six of the above-mentioned compounds, S1–S5 and AS29, on LPS-induced TNF-α and IL-6 production were studied. After pretreatment with analogs at a series of concentrations (1.0, 2.5 and 5 μM) for 30 minutes, cells were treated with LPS (0.5 μg/mL) for 24 hours. An ELISA assay was used to determine the amount of TNF-α and IL-6 released. As shown in Figure 5, all six compounds exhibited good, dose-dependent reduction of LPS-induced TNF-α and IL-6 production. This result further demonstrates the potential of these analogs as anti-inflammatory agents.

**Figure 4** Plots of predicted activity against the corresponding experimental activity on (A) TNF-α and (B) IL-6 inhibition.

**Note:** Numbers in parentheses mean the standard deviation of the coefficients.

**Abbreviations:** PER, predicted activity; EXP, experimental activity; TNF-α, tumor necrosis factor-α; IL-6, interleukin-6; Eq, equation; IR, inhibition rate; EEVA, electronic eigenvalue descriptors; RDF, radial distribution function; EVA, eigenvalue descriptor; SAS, solvent-accessible surface; N, the number of compounds taken into account in the regression; $R^2$, the multiple correlation coefficient; $R^2_{adj}$, adjusted multiple correlation coefficient; s, residual standard error.
**S1 attenuated LPS-induced ERK phosphorylation**

In the lipopolysaccharide–toll-like receptor 4 (LPS–TLR4) inflammatory pathway, NF-κB, a nuclear transcriptional factor, and mitogen-activated protein kinases (MAPKs) play key roles in regulating inflammatory cytokine gene expression. Curcumin has been demonstrated to reduce a variety of gene transcriptions via suppressing the activation of NF-κB and MAPKs. To identify the possible signaling pathways involved in the anti-inflammatory effects of compounds S1–S5 and AS29, we first examined the effect of these six compounds on LPS-activated NF-κB and MAPKs. Three types of MAPKs have been identified: extracellular signal-regulated kinase (ERK), c-Jun N-terminal kinase (JNK), and p38 MAPK. Under resting conditions, NF-κB is bound to inhibitory κB proteins such as IκB-α, which sequester NF-κB in cytoplasm as an inactive complex. In response to LPS, IκB-α is subjected to ubiquitylation, followed by proteasome-mediated degradation, after which NF-κB moves into the nucleus to promote the transcription of inflammatory genes. Thus, we used a Western blot to detect IκB degradation and the phosphorylation of ERK, JNK, and p38 in LPS-stimulated macrophages with or without the pretreatment of these compounds. As illustrated in Figure 6A, the addition of LPS for 20 minutes resulted in obvious phosphorylation (ERK, P38, and JNK) and degradation of IκBα. Pretreatment with the anti-inflammatory active analogs S1–S5 and AS29 at 10 μM did not reduce the MAPK phosphorylation and IκB-α degradation, with the exception of S1, which significantly inhibited the LPS-induced phosphorylation.

![Figure 5](https://www.dovepress.com/)

*Figure 5* Active compounds dose-dependent inhibition of the production of (A) TNF-α and (B) IL-6 induced by LPS in mouse macrophages.

**Notes:** (A and B) Cells were seeded on a plate with the density of 4.0×10⁵/mL overnight at 37°C and 5% CO₂. Macrophages were pre-incubated with vehicle (DMSO) or analogs at 1, 2.5, and 5.0 μM for 2 hours, followed by stimulation with LPS (0.5 μg/mL) for an additional 22 hours. The levels of TNF-α and IL-6 in culture medium were detected by ELISA and were standardized by the total protein. The results were indicated as the proportion of LPS control. Each bar expresses mean ± SEM of 3–6 independent experiments. Statistical difference with regard to LPS was demonstrated: *P<0.05, **P<0.01.

**Abbreviations:** LPS, lipopolysaccharide; IL-6, interleukin-6; DMSO, dimethyl sulfoxide; TNF-α, tumor necrosis factor-α; ELISA, enzyme linked immunosorbent assay; SEM, standard error of the mean.
of ERK. To determine the dose-dependent effect of S1, macrophages were pretreated with doses of 2.5, 5, or 10 μM for 30 minutes. LPS was then added, and the macrophages were incubated for an additional 20 minutes. As shown in Figure 6B, S1 dose-dependently attenuated LPS-induced ERK phosphorylation. These data confirm that S1 can inhibit LPS-induced ERK phosphorylation and may therefore be a potential ERK inhibitor. These results indicate that there may be a mechanistic difference among these curcumin analogs, although they are derived from the same structural lead.

The toxicity and stability of active compounds
As a natural active product, curcumin exhibits no toxicity and is very safe in clinical trials. Before the in vivo experiment, an MTT assay was used to test the cytotoxicity of six active analogs, S1–S5 and AS29, in the human normal hepatic cell line HL-7702. As shown in Figure 7A, all six analogs showed low toxicity at a concentration of 10 μM, indicating that they are relatively safe. However, the clinical applications of curcumin have been significantly limited by its poor chemical stability. The monocarbonyl analogs of curcumin are designed to improve chemical stability by removing the central β-diketone moiety. Thus, the stability of the new synthetic analogs in phosphate buffer (pH 7.4) containing 5% DMSO were tested here using the ultraviolet-visible (UV-visible) absorption spectra at different times (0, 5, 10, 15, 20 and 25 minutes). The optical density (OD) values of the maximal absorption peak of curcumin decreased quickly, and analogs S1, S3, S4, S5, and AS29 showed much higher stability than curcumin, while S2 slightly decomposed during the incubation (Figure 7B–H). Meanwhile, the half-life of the most active compounds S1 and S4 were detected by HPLC in vitro. As shown in Figure S1, the half-lives of S1 (>120 minutes) and S4 (>240 minutes) are longer than the compound curcumin (<90 minutes). These results indicate that the active compounds are chemically more stable than curcumin in vitro.

Anti-inflammatory effect of S1 and S4 in vivo
LPS has been implicated as an important pathogenic factor for the induction of sepsis, which is characterized by an inflammatory cytokine storm. Curcumin and its analogs have been reported to have therapeutic effects on sepsis and septic shock. We have illustrated the inhibitory effects of these analogs on LPS-induced pro-inflammatory cytokine production. We further determined whether two representative compounds, S1 and S4, were able to relieve septic shock through inhibition of LPS-induced inflammatory response in mice. The water-soluble formulations of S1 and S4 were prepared for intravenous injection. Mice were injected intraperitoneally (ip) with LPS at the dosage of 15 mg/kg 15 minutes after the intravenous (iv) injection of 10 mg/kg S1 or S4, and their survival rates were monitored for 7 days.

**Figure 6** Compound S1 inhibition of ERK phosphorylation.
**Notes:** (A) Macrophages were cultured with or without S1–S5 or AS29 for 0.5 hour and stimulated with LPS (0.5 μg/mL) for a further 20 minutes. Cells were harvested and the total protein was extracted. The protein levels of p-ERK, ERK, p-JNK, JNK, p-P38 and P38 were determined by Western blot. (B) S1 dose-dependently decreased ERK phosphorylation. Representative blots of three independent experiments in each study are shown. Statistical significance with regard to LPS was demonstrated: *P*<0.05, **P**<0.01, and ***P***<0.001, respectively.

**Abbreviations:** Comp, compound; LPS, lipopolysaccharide; ERK, extracellular regulated protein kinase; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; JNK, c-Jun N-terminal kinase; P, phosphorylation; SEM, standard error of the mean.
Figure 7 Six active analogs exhibited non-cytotoxicity and chemical stability.

Notes: (A) Viability of HL-7702 cells was detected through an MTT assay after treatment with active analogs for 24 hours. (B–H) Compound stability was described by the curves, which represent absorbance at various optical densities (250–600 nm) and intervals (0, 5, 10, 15, 20, and 25 minutes).

Abbreviations: HL-7702, human normal hepatic cell line 7702; MTT, methyl thiazolyl tetrazolium; OD, optical density; DMSO, dimethyl sulfoxide; Cur, curcumin; min, minute; nm, nanometer.

Figure 8 Protection of (A) S1 and (B) S4 on the mortality of mice caused by LPS.

Notes: Mice were injected with 10 mg/kg compound S1 or S4 15 minutes prior to treatment with 15 mg/kg LPS. After the LPS injection, survival rates were recorded daily for 7 days. Results from the record of experiments are displayed. (A) depicts three groups of nine animals each. (B) depicts three groups of ten animals each.

Abbreviations: LPS, lipopolysaccharide; n, number.
The results in Figure 8 show that 100% of the animals injected with LPS alone died within 3–4 days as a result of septic shock. In septic animals administrated with S1 and S4, the survival rates were increased significantly compared with the LPS-alone group (40% in the S1-treated group and 50% in the S4-treated group). Thus, these data suggest that S1 and S4 exhibit anti-inflammatory activity in vivo.

Conclusion

In summary, we synthesized a series of symmetrical and asymmetrical analogs of curcumin and estimated their anti-inflammatory activities against LPS-induced TNF-α and IL-6 release in mouse macrophages. The majority of analogs effectively inhibited the LPS-induced production of TNF-α and IL-6. Together with the SAR and QSAR analysis, we found that the monocarbonyl curcumin analogs with symmetric and heterocyclic structures have stronger anti-inflammatory activity. Six compounds with potent anti-inflammatory activity also exhibited excellent chemical stability and low toxicity in vitro. Mechanistically, compound S1 significantly inhibited LPS- induced phosphorylation of ERK. Furthermore, S1 and S4 were selected for in vivo anti-sepsis testing and were found to markedly decrease LPS- induced lethality in septic mice. All of these results indicate that these novel monocarbonyl curcumin analogs may serve as potential agents for the treatment of various inflammatory diseases.

Acknowledgments

Financial support was provided by the National Natural Science Fund of China (Grants 21272179, 81272462, and 81202462); Project of Zhejiang Provincial Key Constructive Subject (Traditional Chinese Medicine, Grant 2012-XK-A28); Qianjiang Talent Project of Zhejiang Province (Grant 2013R10020), and the Zhejiang Key Group Project in Scientific Innovation (Grant 2010R50042).

Disclosure

The authors report no conflicts of interest in this work.

References


