

Two copper(II) complexes of curcumin derivatives: synthesis, crystal structure and in vitro antitumor activity

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Received: 12 February 2014 / Accepted: 28 April 2014
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Abstract Two Cu(II) complexes of curcumin derivatives, formulated as CuL_2^{a} (**1**) and CuL_2^{b} (**2**) [HL^{a} = 1,7-bis(4-ethyloxy-3-methoxy-phenyl)-1,6-heptadiene-3,5-dione and HL^{b} = 1,7-bis(4-butyloxy-3-methoxy-phenyl)-1,6-heptadiene-3,5-dione], have been synthesized and characterized by single-crystal X-ray diffraction, along with physico-chemical and spectroscopic methods. In both complexes, each Cu(II) center is surrounded by four oxygen atoms from two β -diketone ligands in a square planar geometry. Complex **1** forms a 2D layer structure through intermolecular π - π stacking interactions, as well as weak coordination interactions between the Cu and O atoms of the solvent 1,4-dioxane molecules. Complex **2** displays a 1D column structure stabilized by intermolecular π - π stacking interactions. MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assays were used to evaluate the cytotoxicities of these complexes against three human cancer cell lines. The results show that the Cu(II) complexes exhibit more potent inhibition tumor growth in comparison with the free ligands.

Introduction

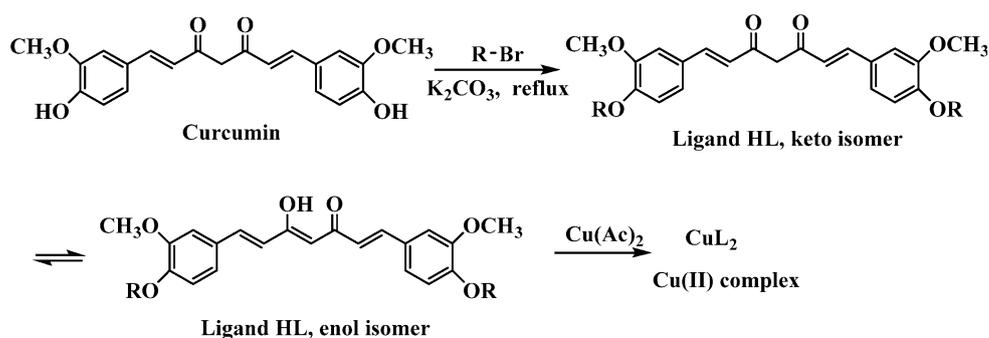
Curcumin, 1,7-bis(4-hydroxyl-3-methoxyphenyl)-1,6-heptadiene-3,5-dione, is a naturally occurring yellow pigment with a symmetric β -diketone unit extracted from the

rhizomes of turmeric (*Curcuma Longa Linn.*), which has attracted interest in the fields of biology, medicine and pharmacology over the past decades owing to its antitumor, anti-inflammatory, antiviral, antibacterial and antioxidant activities [1–4]. Ongoing clinical trials of curcumin, including studies in colon cancer (phase I/II), pancreatic cancer (phase II/III), cervical cancer (phase II/III), oral cancer (phase II/III), rectal cancer (phase II) and multiple myeloma, have shown that this compound exhibits anti-proliferative activity against a variety of tumors via diverse signaling pathways that affect tumor growth, involving nuclear factor- κ B (NF- κ B), PI3K/Akt, activator protein-1 (AP-1) and cyclooxygenase-2 (COX-2) [5–9]. Importantly, curcumin is very safe to humans even at 8,000 mg/day dosage orally for 3 months [10]. Although curcumin possesses wide-spectrum and low-toxicity antitumor activity, its clinical application in antitumor therapy has been greatly limited by its rapid metabolism, low absorption and poor stability in vivo [11]. In order to improve the bio-availability and antitumor activity of curcumin, a large number of curcumin derivatives and analogues have been designed and synthesized through structural modifications such as variation of the aromatic rings and their substituents and/or replacing the heptadiendione bridge chain of curcumin with other linkers [12–17]. Because the poor stability of curcumin is reported to mainly result from the phenolic hydroxyl groups [18], our group has synthesized a series of dialkoxyl substituted curcumin derivatives, such as 1,7-bis(4-ethyloxy-3-methoxy-phenyl)-1,6-heptadiene-3,5-dione (HL^{a}) and 1,7-bis(4-butyloxy-3-methoxy-phenyl)-1,6-heptadiene-3,5-dione (HL^{b}), using the Williamson ether reaction between halohydrocarbons and the phenolic hydroxyl group of curcumin (Fig. 1). The introduction of R groups to the hydroxyl positions not only increases the stability of the derivatives but also retains the β -diketone

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Fig. 1 Synthesis route of curcumin derivatives and their Cu(II) complexes (HL^a: R =CH₂CH₃; HL^b: R =CH₂CH₂CH₂CH₃)



fragment which can act as a good O-donor chelating ligand. It has been demonstrated that the coordination of curcumin with various metals can enhance the antitumor activity [19–25]. Metal complexes have attracted considerable attention as new anticancer therapeutic candidates in recent years because metal ions are essential in many natural biological processes. Moreover, metal complexes can offer mechanisms of action that are unavailable to organic compounds. However, to the best of our knowledge, the coordination chemistry of curcumin derivatives has been little explored [26, 27]. Considering that Cu(II) is an essential micronutrient participating in many biological processes such as mitochondrial respiratory reaction, energy generation, cellular stress response, antioxidant and antitumor defense [28], in this paper, we have used our curcumin derivatives HL^a and HL^b as ligands to prepare two new Cu(II) complexes, formulated as CuL₂^a (**1**) and CuL₂^b (**2**), respectively (Fig. 1). It is hoped that the conjugation of Cu(II) with these curcumin derivatives may enhance their antitumor activity. Herein, we report the synthesis, crystal structures and in vitro antitumor activities of the two complexes.

Experimental

Materials and instruments

All reagents were purchased commercially and used directly without further purification. The curcumin derivatives, HL^a and HL^b, were prepared by the method we reported in Ref. [29]. Elemental analyses were obtained on a Perkin-Elmer 240 analyzer. Infrared spectra were recorded on a Nicolet FT-IR 170 SX spectrophotometer with KBr pellets in the 4,000–400 cm⁻¹ region.

Synthesis of complex 1

CuL₂^a (**1**) was prepared as follows. HL^a (0.85 g, 2.0 mmol) was dissolved in ethanol (30 mL), then solid sodium hydroxide (0.09 g, 2.2 mmol) and a solution of Cu(OAc)₂

(0.18 g, 1.0 mmol) in ethanol (10 mL) were added into the solution sequentially at room temperature. The reaction mixture was stirred at 80 °C for 6 h. After cooling to room temperature, the complex was filtered off, washed thoroughly with water and ethanol and then dried under vacuum. Yellow green crystals; Yield 0.65 g (65 %). FTIR (KBr, cm⁻¹): 2,979 (CH₃), 1,623 (C=O), 1,510 (C=C), 1,598, 1,577, 1,477 (Ar), 1,257 (=CH), 543 (Cu–O); Anal. Calcd. for C₅₄H₆₂CuO₁₄: C 65.0, H 6.2, Cu 6.4 %; Found: C 64.5, H 6.0, Cu 6.2 %; ¹H-NMR (d₆-DMSO, 400 MHz) δ: 1.63 (*t*, 12H, *J* = 8 Hz, CH₃), 4.0 (*s*, 12H, –OCH₃), 4.14–4.21 (*q*, 16H, *J* = 8 Hz, OCH₂), 5.96 (*s*, 2H, O=C–CH–C=O), 6.61–6.70 (*d*, 4H, *J* = 16 Hz, =CH), 7.07–7.31 (*m*, 12H, Ar–H), 7.57–7.63 (*d*, 4H, *J* = 16 Hz, =CH–Ar).

Synthesis of complex 2

CuL₂^b (**2**) was synthesized by a similar procedure to complex **1**, except using ligand HL^b instead of HL^a. Yellow crystals; Yield 0.60 g (59 %). FTIR (KBr, cm⁻¹): 2,957 (CH₃), 1,624 (C=O), 1,510 (C=C), 1,598, 1,579, 1,466 (Ar), 1,262 (=CH), 533 (Cu–O); Anal. Calcd. for C₅₈H₇₀CuO₁₂: C 68.1, H 6.9, Cu 6.2 %; Found: C 68.4, H 6.6, Cu 6.5 %; ¹H-NMR (d₆-DMSO, 400 MHz) δ: 1.32 (*t*, 12H, *J* = 8 Hz, CH₃), 3.15–3.34 (*m*, 16H, –CH₂), 3.87 (*s*, 12H, –OCH₃), 4.10–4.19 (*m*, 8H, OCH₂), 5.88 (*s*, 2H, O=C–CH–C=O), 6.56–6.64 (*m*, 4H, =CH), 6.82–7.16 (*m*, 12H, Ar–H), 7.42–7.53 (*m*, 4H, =CH–Ar).

Crystal structure determination

Single crystals, suitable for X-ray analyses, were obtained by slow evaporation of ethyl acetate/1,4-dioxane solutions at room temperature. Single-crystal X-ray diffraction data were collected on a Bruker SMART APEX CCD diffractometer equipped with graphite monochromated Mo Kα radiation (λ = 0.71073 Å) at ambient temperature. All independent reflections were collected in a range of 0.94–25.00° for complex **1** and 1.52–24.97° for complex **2** (determined in the subsequent refinement). The structures were solved by direct methods with SHELXS-97 and

Table 1 Crystal data and structure refinement parameters for **1** and **2**

Complex	1	2
Empirical formula	C ₅₄ H ₆₂ CuO ₁₄	C ₅₈ H ₇₀ CuO ₁₂
Formula weight	998.58	1,022.68
Temp (K)	296 (2)	298 (2)
λ (Å)	0.71073	0.71073
Crystal system	Monoclinic	Monoclinic
Space group	P2 ₁ /c	C2/m
<i>a</i> (Å)	21.86 (2)	15.556 (5)
<i>b</i> (Å)	16.453 (16)	34.764 (5)
<i>c</i> (Å)	7.184 (7)	8.546 (5)
β (°)	97.643 (11)	110.882 (5)
<i>V</i> (Å ³)	2,561 (4)	4,318 (3)
<i>Z</i>	2	2
<i>D_c</i> (g cm ⁻³)	1.295	0.787
μ (mm ⁻¹)	0.492	0.291
<i>F</i> (000)	1,054	1,086
Crystal size (mm)	0.30 × 0.20 × 0.20	0.30 × 0.20 × 0.20
θ range (°)	0.94–25.00	1.52–24.97
Limiting indices	–25 ≤ <i>h</i> ≤ 25 –18 ≤ <i>k</i> ≤ 19 –8 ≤ <i>l</i> ≤ 8	–18 ≤ <i>h</i> ≤ 18 –41 ≤ <i>k</i> ≤ 40 –10 ≤ <i>l</i> ≤ 8
Reflections collected	17,861	13,453
Reflections unique (<i>R_{int}</i>)	4,510 (0.0247)	3,783 (0.0967)
Parameters	317	165
Goodness of fit on <i>F</i> ²	1.095	0.934
Final <i>R</i> indices [<i>I</i> > 2 σ (<i>I</i>)] ^a	<i>R</i> ₁ = 0.0503, <i>wR</i> ₂ = 0.1420	<i>R</i> ₁ = 0.0836, <i>wR</i> ₂ = 0.2191
<i>R</i> indices (all data) ^a	<i>R</i> ₁ = 0.0629, <i>wR</i> ₂ = 0.1557	<i>R</i> ₁ = 0.1524, <i>wR</i> ₂ = 0.2523
Largest diff. peak and hole (e [–] Å ⁻³)	0.354 and –0.718	0.305 and –0.365

^a $\omega = 1 / [\sigma^2(F_0^2) + (0.0881P)^2 + 1.2757P]$ for **1** and $\omega = 1 / [\sigma^2(F_0^2) + (0.1216P)^2 + 0.0000P]$ for **2**, where $P = (F_0^2 + 2F_c^2)/3$

Table 2 Selected bond lengths (Å) and angles (°) for **1** and **2**^a

	1		2
Cu1–O3	1.929 (2)	Cu1–O3	1.988 (3)
Cu1–O4	1.903 (2)	O3–Cu1–O3#2	93.28 (18)
O3–Cu1–O4	93.68 (11)	O3–Cu1–O3#3	86.72 (18)
O3–Cu1–O4#2	86.32 (11)		

^a Symmetry transformations used to generate equivalent atoms: complex **1** #2 –*x* + 2, –*y* + 2, –*z* + 1; complex **2** #2 –*x* + 2, *y*, –*z* + 3 and #3 *x*, –*y* + 2, *z*

refined by full-matrix least-squares methods on *F*² with SHELXL-97 software. All non-hydrogen atoms were refined anisotropically, while hydrogen atoms were added at the calculated positions and refined isotropically using a

riding model. In the final difference map, the residual maximum and minimum were 0.354 and –0.718 e Å⁻³ for **1**, 0.305 and –0.365 e Å⁻³ for **2**, respectively. Crystallographic data and experimental details for complexes **1** and **2** are summarized in Table 1. Selected bond lengths and angles are listed in Table 2.

Cell lines and culture conditions

Three different human cancer cell lines were used: ASPC-1 (pancreatic carcinoma), MCF-7 (breast cancer) and HeLa (cervical cancer). All cell lines were purchased from the Cell Culture Center of China Pharmaceutical University. The cell lines were cultured in RPMI 1,640 medium supplemented with 10 % (v/v) fetal bovine serum (FBS) and antibiotics (100 U/mL of penicillin and 100 µg/mL of streptomycin) in cell culture incubators at 37 °C in a humidified atmosphere containing 5 % (mass fraction) CO₂.

MTT cytotoxicity assay

The effects of the free ligands and their complexes on cell proliferation were determined using the MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assay. The cells were seeded in 96-well plates at a concentration of 2 × 10⁴ cells/well. Subsequently, the cells were treated with graded concentrations of reagents in triplicate. After 48 h of incubation, 40 µL of MTT (5 mg/mL) was added to each well, which was then incubated for an additional 4 h. After discarding the medium, the cells were dissolved in DMSO (150 µL/well) and the absorbance was recorded at 570 nm. The cell lines without any drug treatment were used as a blank control in all experiments. The cell survival rates were calculated as a percentage by comparison of the treated cells with the untreated control. The maximal inhibitory concentration at 50 % cell survival rate of each reagent (IC₅₀) was determined from the curve or regression equation of the survival rate versus drug concentration.

Results and discussion

Crystal structures of the complexes

As shown in Fig. 2a, complex **1** has a neutral mononuclear structure, composed of one copper(II) center and two anionic (L^a)[–] ligands, being formulated as CuL₂^a. The central Cu atom is four-coordinated by two oxygen atoms of the carbonyl groups and two oxygen atoms of enolic hydroxyl groups from two β-diketone ligands, which adopt the chelating coordination modes in their enol forms. The

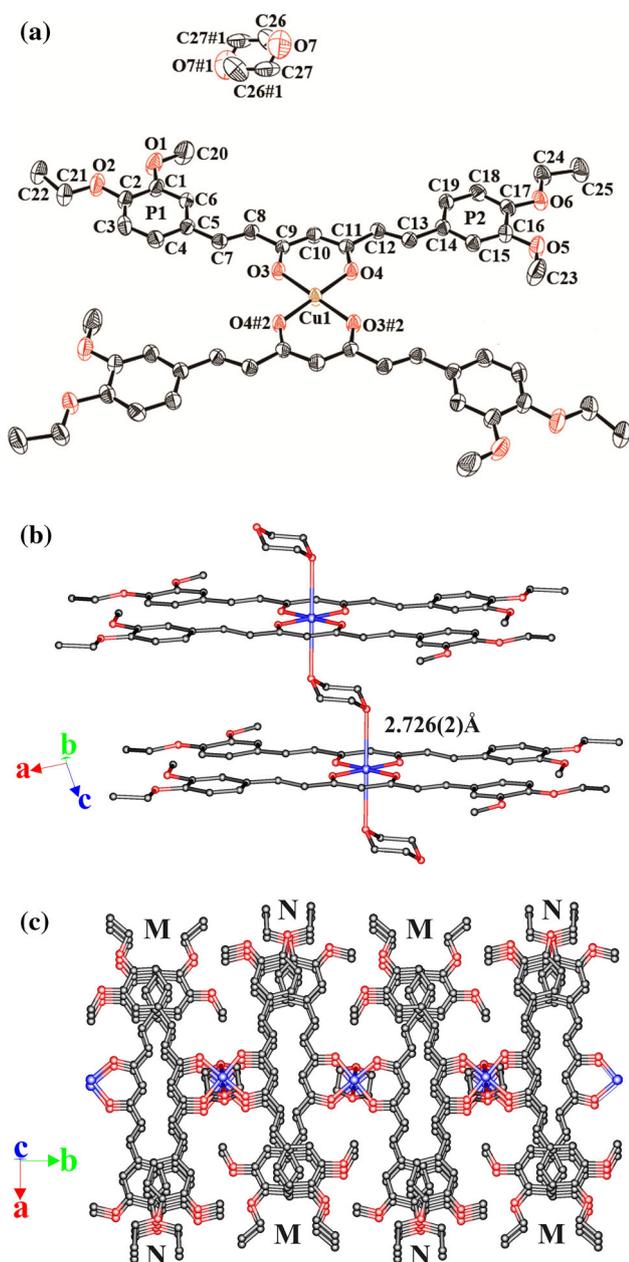


Fig. 2 **a** ORTEP drawing of the molecular structure of **1** with atomic labeling scheme (50 % thermal ellipsoids probability, symmetry code: #1 $-x + 2, -y, -z$; #2 $-x + 2, -y + 2, -z + 1$). **b** View showing the 1D column structure of **1** along c -axis formed through weak coordination interactions between the Cu and the O atoms of the 1,4-dioxane molecules. **c** View showing the 2D layer structure of **1** on the ab plane stabilized by π - π stacking interactions (Hydrogen atoms are omitted for clarity)

Cu–O bond lengths are 1.929 (2) and 1.903 (2) Å, which are a little shorter than those in other copper diketonate complexes [30, 31]. The O–Cu–O bond angles are 93.68 (11) and 86.32 (11)°, and the sum of them is 180°, indicating the Cu(II) center adopts a square planar configuration. The dihedral angle between the two benzene rings P1 and P2 is 4.09°, while the dihedral angles they form with

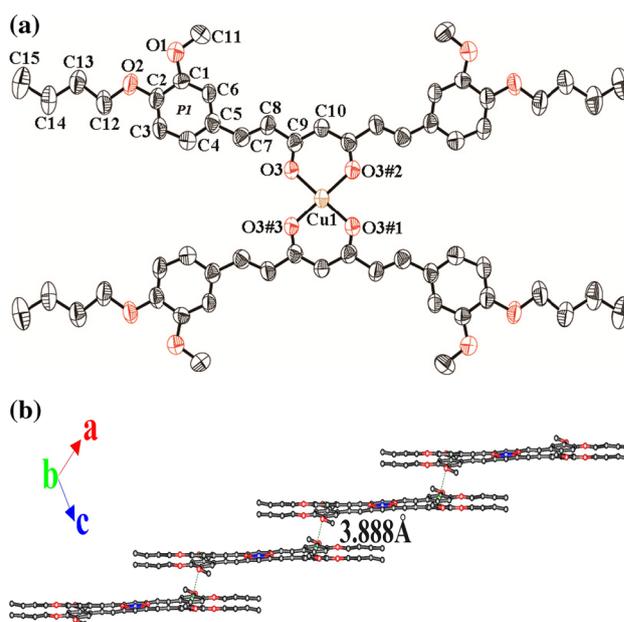


Fig. 3 **a** ORTEP drawing of the molecular structure of **2** with atomic labeling scheme (50 % thermal ellipsoids probability, symmetry code: #1 $-x + 2, -y + 2, -z + 3$; #2 $-x + 2, y, -z + 3$; #3 $x, -y + 2, z$). **b** View showing the 1D column structure of **2** formed by π - π stacking interactions (Hydrogen atoms are omitted for clarity)

the coordination plane composed of one Cu(II) atom and four oxygen atoms are 3.33° and 1.42°, respectively, showing that they are nearly coplanar. In complex **1**, an infinite 1D column arrangement along the c -axis is observed. This 1D column structure is formed by weak coordination interactions between the Cu and the O atoms of the 1,4-dioxane molecules, with a distance of 2.726 (2) Å (Fig. 2b). The Cu...Cu separation across the 1,4-dioxane bridge is 7.184 Å, and all Cu(II) atoms are in a straight line. The 1D columns are further extended into a 2D layer structure extending along the ab plane through two kinds of alternant π - π interactions between almost parallel benzene rings P1 and P1 (centroid–centroid separation 3.846 Å with a dihedral angle of 5.62°, described as M in Fig. 2c), and between P2 and P2 rings (centroid–centroid separation 3.897 Å with a dihedral angle of 8.13°, described as N in Fig. 2c) from different mononuclear units.

The crystal structure of complex **2** is shown in Fig. 3a. The asymmetric unit contains one Cu(II) atom and two L^b ligands, the complex being formulated as CuL_2^b . The Cu atom is located on an inversion center and four-coordinated by four crystallographically equivalent oxygen atoms from two chelating β -diketonate ligands in their enol form. The Cu–O bond lengths are all 1.988 (3) Å. The O–Cu–O bond angles are 93.28 (18) and 86.72 (18)°, respectively, and their sum is also 180°, revealing the Cu(II) center adopts a square planar geometry in this complex also. The dihedral

Table 3 IC₅₀ values of the free ligands and their Cu(II) complexes against three cancer cell lines

Compound	IC ₅₀ (μM)		
	ASPC-1	MCF-7	HeLa
CuL ₂ ^a (1)	6.52	3.22	11.48
CuL ₂ ^b (2)	7.45	4.43	13.34
HL ^a	22.90	14.96	40.69
HL ^b	24.68	18.06	43.26

angle between benzene ring P1 and the coordination plane composed of one Cu(II) atom and four oxygen atoms is 6.01°, and the dihedral angle between two benzene rings in each ligand is 9.23°, which are both slightly larger than those in complex **1**, indicating that complex **2** is twisted slightly to reduce steric hindrance. The structure is extended into a 1D column by π - π stacking interactions between the benzene rings, with a centroid-centroid separation of 3.888 Å and dihedral angle of 9.23° (Fig. 3b).

In vitro antitumor activities

We evaluated the cytotoxicities of the free ligands and their Cu(II) complexes against three cell lines, ASPC-1, MCF-7 and HeLa. MTT assays were performed to detect cell survival rates following exposure to the tested compounds during 48 h of incubation. Table 3 summarizes IC₅₀ values from a series of individual experiments. It can be observed that both complexes significantly enhanced the inhibition activity of tumor growth in contrast to the two ligands. The cytotoxicities of complexes **1** and **2** were about three–fivefold higher than the corresponding free ligands against every cancer cell tested. The increased growth-suppressive activity of the complexes may be associated with the DNA-binding interactions of copper ions [32–35]. The IC₅₀ levels of complex **1** against all three cancer cells were a little lower than those of complex **2**, and the same phenomenon was also observed for the free ligands, which may be a result of a spacer effect. When HL^b was used in place of HL^a, the increased steric hindrance of the R groups would influence the efficient transmembrane transportation and penetration of the compounds into tumor cells, resulting in the decrease in cytotoxicity.

Conclusions

In this work, we have synthesized two new Cu(II) complexes of curcumin derivatives. The difference in R groups, which were introduced to the phenolic hydroxyl sites of curcumin, not only leads to different crystal structures for the two complexes, but also influences their antitumor

activities. Complex **1** exhibits higher antiproliferative activity against three human cancer cell lines tested than complex **2**, suggesting that increased steric hindrance in the R groups will bring down the antitumor activity. Both complexes show significantly enhanced antitumor activity in comparison with the free ligands, which suggests that exploring the coordination chemistry of curcumin derivatives is an effective method to improve their antitumor activity. Further studies on the in vivo antitumor activities of these complexes are in progress.

Supplementary material

CCDC reference numbers 972326 (CuL₂^a) and 937711 (CuL₂^b) contain the supplementary crystallographic data for this paper. These data can be obtained free of charge at www.ccdc.cam.ac.uk/conts/retrieving.html or from the Cambridge Crystallographic Data Centre, 12, Union Road, Cambridge CB2 1EZ, UK; fax: +44 1223/336 033; e-mail: deposit@ccdc.cam.ac.uk.

Acknowledgments This research was financially supported by the Opening Foundation of Anhui Province Key Laboratory of Environment-friendly Polymer Materials (No. 2013KF001), the National Natural Science Foundation of China (No. 21071001), the Natural Science Foundation of Anhui Province (No. 1208085MH173), and the Youth Foundation of Anhui University of Chinese Medicine (No. 2011qn027).

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