

Natural Small-Molecule-Based Carrier-Free Self-Assembly Library Originated from Traditional Chinese Herbal Medicine

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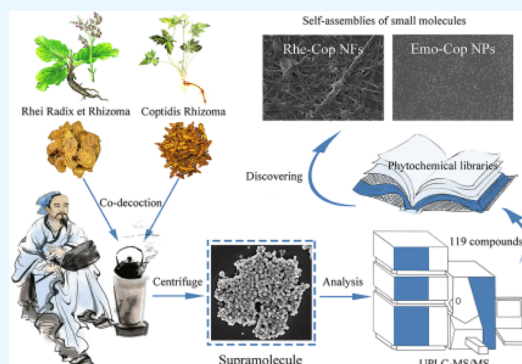
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ABSTRACT: The carrier-free self-assembly of small molecules opens a new window for the development of nanomaterials. This study is dedicated to developing binary small-molecular self-assemblies derived from phytochemicals in traditional Chinese herbal medicine. Among them, *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* are a common pair used in clinics for thousands of years. Here, we found that there were numerous spherical supramolecular nanoparticles (NPs) originated from *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* decoction. Ultra-performance liquid chromatography/tandem mass spectrometry (UPLC-MS/MS) was used to analyze the composition of the supramolecules, and a total of 119 phytochemicals were identified (23 anthraquinones, 31 alkaloids, 24 organic acids, 8 tannins, and other components). Isothermal titration calorimetry (ITC) showed that the interaction between *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* was a spontaneous exothermic reaction, indicating that their phytochemicals had the property of self-assembly and interacted to form supramolecules in the decocting process. Furthermore, scanning electron microscopy (SEM), UV, IR, NMR, and ITC were used to verify that rhein and coptisine could self-assemble into nanofibers (Rhe-Cop NFs), while emodin and coptisine could self-assemble into nanoparticles (Emo-Cop NPs). The formation mechanism analysis of the self-assemblies revealed that they were induced by electrostatic attraction, hydrogen bonding, and π - π stacking, forming nanospheres of about 50 nm and nanofibers. The current study not only provides an idea of discovering carrier-free self-assemblies from traditional herbal medicine decoction but also supplies a reference for the design of binary self-assembly of small molecules in the future.



1. INTRODUCTION

Supramolecular refers to the association of two or more kinds of molecules with intermolecular interactions to form a complex and orderly whole with certain integrity, microstructure, and macroscopic characteristics.¹ At present, supramolecular self-assembly has shown great prospects in the research and development of nanomaterials.² It has great potential in the fields of anticancer,^{3,4} disease diagnosis,⁵ targeted drug delivery,^{6,7} and other nanomedicines.⁸ Self-assembly behavior refers to the process of an orderly arrangement of molecules in a system under the action of noncovalent bonds and finally forming a structure with a certain geometric appearance.⁹ Numerous self-assembly phenomena from small molecules have been found,^{10,11} like puerarin hydrogel,^{12,13} gallic acid gels,¹⁴ pomolic acid gels,¹⁵ and rhein hydrogel,^{16,17} and triterpenoids can also self-assemble under suitable solvent conditions,^{18–20} which exhibit different morphologies and functions, such as oleanolic acid, glycyrrhetic acid,²¹ and ursolic acid.²² Our previous studies and other researchers have also found binary assemblers: berberine can self-assemble with baicalin,²³ rhein,²⁴ aristolochic acid,²⁵ cinnamic acid,²⁶ curcumin,²⁷ and sennoside A²⁸ to

form nanoparticles (NPs). There are also several small-molecule compounds self-assembling with metal ions: luteolin-Fe³⁺/Ca²⁺/Mg²⁺/Zn²⁺/Cu²⁺,^{29,30,29,30} tannic acid/catecholamines-Cu²⁺,³¹ and baicalin-Al³⁺.³² Although some progress has been made in this field, the self-assembly of natural small molecules is still at the stage of random, serendipity discovery. And the previous self-assembly system often involves a complex preparation process, high cost, and so on.^{33,34} The preparation of a self-assembly system is easily affected by external physical and chemical factors (such as solvent,³⁵ pH^{36,37}), resulting in the emergence of a self-assembly system with different structures and functions. Therefore, a goal-oriented design to obtain natural small-molecule self-assembly with a simple strategy is a hot research topic at present (Scheme 1).

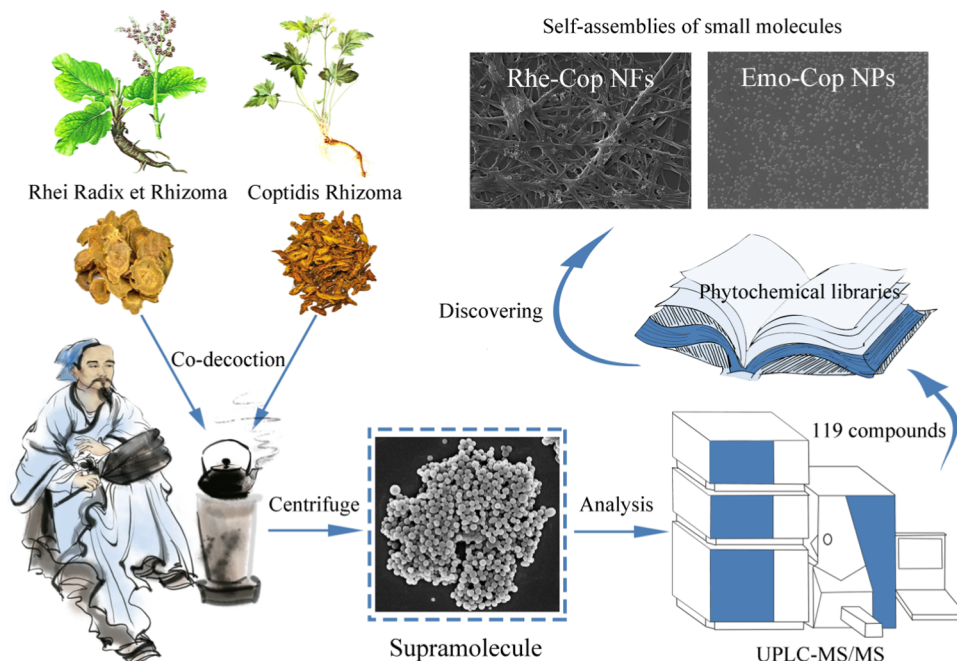
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Scheme 1. Natural Small-Molecule-Based Carrier-Free Self-Assembly Library Originated from Traditional Chinese Herbal Medicine



The composition of traditional Chinese medicine (TCM) decoction is complex and diverse, which is a collection of phytochemicals' libraries.³⁸ The self-assemblies of small molecules in TCM decoction are widespread,^{39,40} and so it could provide valuable ideas for the accurate discovery of supramolecular components in plants. For example, nanoparticles have been found in Ma-Xing-Shi-Gan-Tang decoction,⁴¹ Xue-Fu-Zhu-Yu-Tang decoction,⁴² Bai-Hu-Tang decoction,⁴³ Ge-Gen-Qin-Lian-Tang decoction,⁴⁴ San-Huang-Xie-Xin-Tang decoction,⁴⁵ and others,^{46,47} which showed fascinating properties and activities. Nanoparticles were not only found in TCM decoction but also in single-flavor decoctions such as Fen-Ge decoction,^{48,49} Huang-Lian decoction,⁵⁰ and so on. Most of the nanoparticles in the above decoction had a particle size of 100–600 nm, which exhibited good stability and high bioavailability. Taking Sini decoction⁵¹ and Huang-Lian-Jie-Du decoction⁵² as examples, it was found that the nanoparticles in the decoction were formed by the interaction of many small molecular bioactive components. In addition to the self-assembly through small-molecule interactions in the decoction, some researchers found that polysaccharide^{53,54} and protein^{55,56} components in the decoction could also self-assemble and showed good biological activity. Meanwhile, many chemical components in TCM have unique structures and many modification sites, which facilitate self-recognition and self-assembly.¹⁰ Therefore, TCM decoction can be used as a carrier-free self-assembly library of small molecules to reduce blindness, contingency, and a green strategy in supramolecular design and construction.

Rhei Radix et Rhizoma and *Coptidis Rhizoma* are commonly used as herbal pair medicine in TCM. There is obvious turbidity in the decocting process. As one of the rich compound libraries of decoction, Rhein-Ber nanoparticles were found in our previous study.²⁴ Therefore, starting from the nanoaggregate in *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* co-decoctions, a total of 119 compounds were identified (23 anthraquinones, 31 alkaloids, 24 organic acids,

8 tannins, and other components). Interestingly, we found that the herbal pair's main components rhein and coptisine could self-assemble into nanofibers (Rhe-Cop NFs), while emodin and coptisine could self-assemble into nanoparticles (Emo-Cop NPs). Combined with multiple technologies, this study took Emo-Cop NPs and Rhe-Cop NFs as examples to describe the formation mechanism and exploration process of discovering self-assemblies of small molecules without a carrier from Chinese herbal compound decoction.

2. MATERIALS AND METHODS

2.1. Reagents and Materials. *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* were purchased from Beijing Tongrentang (batch number: 20201125, 200301001). Rhein ($C_{15}H_8O_6$, Rhe, 98%), emodin ($C_{15}H_{10}O_5$, Emo, 98%), coptisine chloride ($C_{19}H_{14}ClNO_4$, Cop, 98%), and DMSO- d_6 were all purchased from Beijing Inokai Technology Co., Ltd. Sodium hydroxide (NaOH), dimethyl sulfoxide (DMSO, C_2H_6OS), and methanol (CH_4O) were all purchased from Beijing Chemical Plant. Ordinary dialysis bag (MWCO: 3000 Da, Shanghai Yuanye Biotechnology Co., Ltd.). Phosphate buffer (Beijing Bairdi Biotechnology Co., Ltd.)

2.2. Sample Preparation. According to the ratio of *Rhei Radix et Rhizoma*/*Coptidis Rhizoma* = 1:1, the corresponding weight of herbs was put into a nonwoven bag and 8 times the amount of water was added and boiled together for 40 min. The co-decoction of *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* was obtained after filtering out the residue. The co-decoction was centrifuged at 80,000 rpm for 30 min, and the supernatant and supramolecular sites were collected, respectively. All samples were lyophilized.

Coptisine chloride and emodin were precisely weighed at 0.1 mmol and dissolved by adding methanol and DMSO, respectively. After mixing and stirring the two solutions, sodium hydroxide solution was added to adjust the pH to 7.0–7.5. The above solution mixture was added to 60 °C

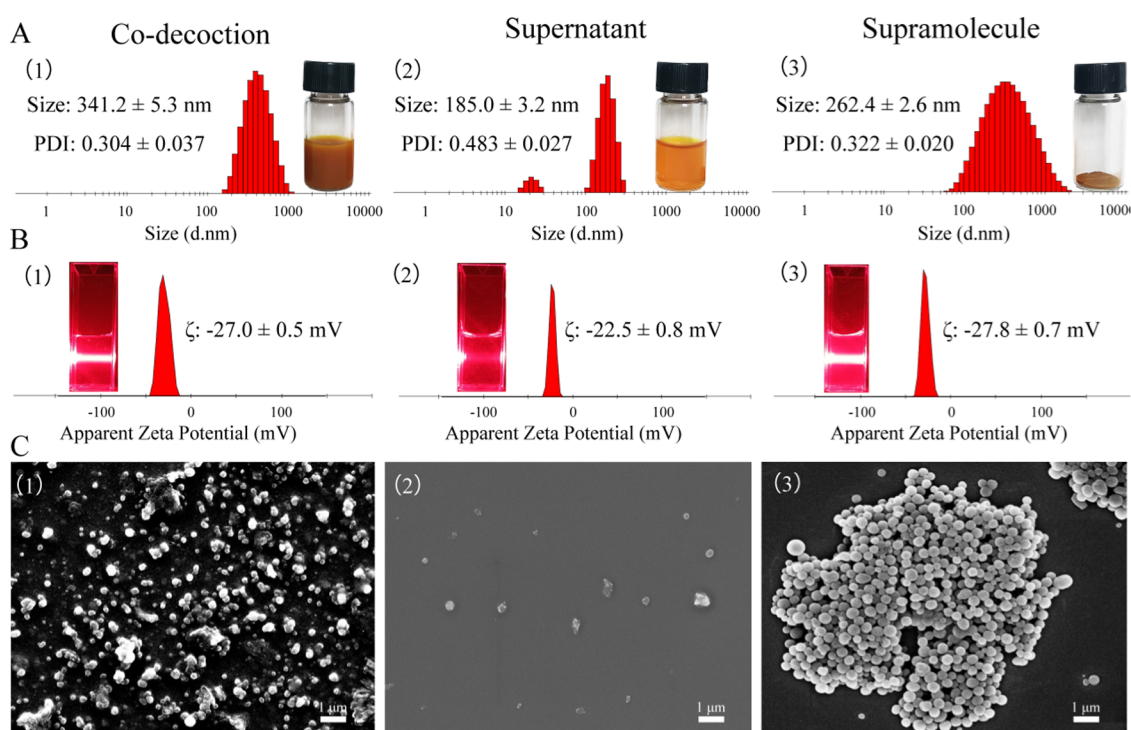


Figure 1. Morphological characterization of *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* co-decoction (1), supernatant (2), and supramolecular portion (3). (A) Particle size and PDI characterization. (B) ζ-Potential and Tyndall effect characterization. (C) Scanning electron microscopy (SEM) characterization.

phosphate-buffered saline (PBS) drop by drop. After thorough stirring, water bath ultrasonication was performed for 1 h. The reaction solution was transferred to a dialysis bag for 12 h to remove unreacted raw materials and organic solvents. The dialysate fluid outside should be replaced every 3 h. Emodin-Coptisine nanoparticles (Emo-Cop NPs) were obtained by freeze-drying its dialysate fluid inside. Similarly, Rhein-Coptisine nanofibers (Rhe-Cop NFs) assembly could be prepared.

2.3. Characterization by Dynamic Light Scattering (DLS). Each sample was dispersed in deionized water and transferred to a colorimetric dish. Using a Malvern particle size analyzer, the mean particle size and polydispersity index (PDI) of the samples were measured in size mode and the ζ-potential of the samples was measured in ζ-mode. The results were repeated three times and averaged.

2.4. Field Emission Scanning Electron Microscopy Observation. The lyophilized powder samples were dispersed in deionized water and 2 μL was transferred to silicon wafers for drying naturally at room temperature. After spraying gold on the surface, a working voltage of 15.0 kV (ZEISS-SUPRA55, ZEISS, Germany) was set to observe the morphology and particle size of each sample.

2.5. Ultra-Performance Liquid Chromatography/Tandem Mass Spectrometry (UPLC-MS/MS). TC-C18 column (4.6 mm × 250 mm, 5 μm, Agilent) was used. The mobile phase consisted of 0.1% (v/v) aqueous formic acid solution (A) and acetonitrile (B). The gradient elution conditions were 0–30 min, 4–98% B. The flow rate was 0.3 mL/min and the injection volume was 5 μL. The ion source adopted ESI to collect information in positive (ESI⁺) and negative (ESI⁻) electrospray ionization mode (EM ACE600, Leica Technology Co., Ltd.). The atomized pressure was 45 psi. The dry gas was

nitrogen gas. The capillary and auxiliary gas heater temperatures were both set to 350 °C.

2.6. UV–Visible Absorption Spectrometric Determination. Two lyophilized powders of self-assembly and three monomer components were fully dissolved in methanol separately. The detection wavelength of the UV–visible spectrophotometer (UH5300, HITACHI, Japan) was set at 190–600 nm and methanol was used as a blank solution for full-wavelength scanning.

2.7. Fourier Transform Infrared Spectroscopy Determination. Two lyophilized powders of self-assembly and three monomer components were put into a Fourier transform infrared spectrometer (Tensor 27, Bruker). Later, the infrared spectra of each sample were measured in the range of 400–4000 cm⁻¹ with air as the background.

2.8. Nuclear Magnetic Resonance Hydrogen Spectrometry (¹H NMR) Determination. Two lyophilized powders of self-assembly and three monomer components were placed in the nuclear magnetic tube. Then, deuterium dimethyl sulfoxide was added to dissolve thoroughly, and tetramethylsilane (TMS) was used as an internal standard to determine the ¹H NMR spectra of each sample in NMR apparatus (Bruker Avance 400 MHz, Bruker, Germany).

2.9. Isothermal Titration Calorimetry (ITC) Determination. All reserve fluid must be filtered with a 0.45 μm microporous membrane before use. The titration threshold was explored through the preliminary experiment, and the titration condition was determined as follows: 10 mg/mL single decoction of *Coptidis Rhizoma*-titrated 20 mg/mL single decoction of *Rhei Radix et Rhizoma*; 2.5 mmol/L coptisine chloride-titrated 0.5 mmol/L rhein; 5 mmol/L emodin-titrated 0.5 mmol/L coptisine chloride. The instrument parameters were set as follows: a stirring rate of 250 r/min, titration

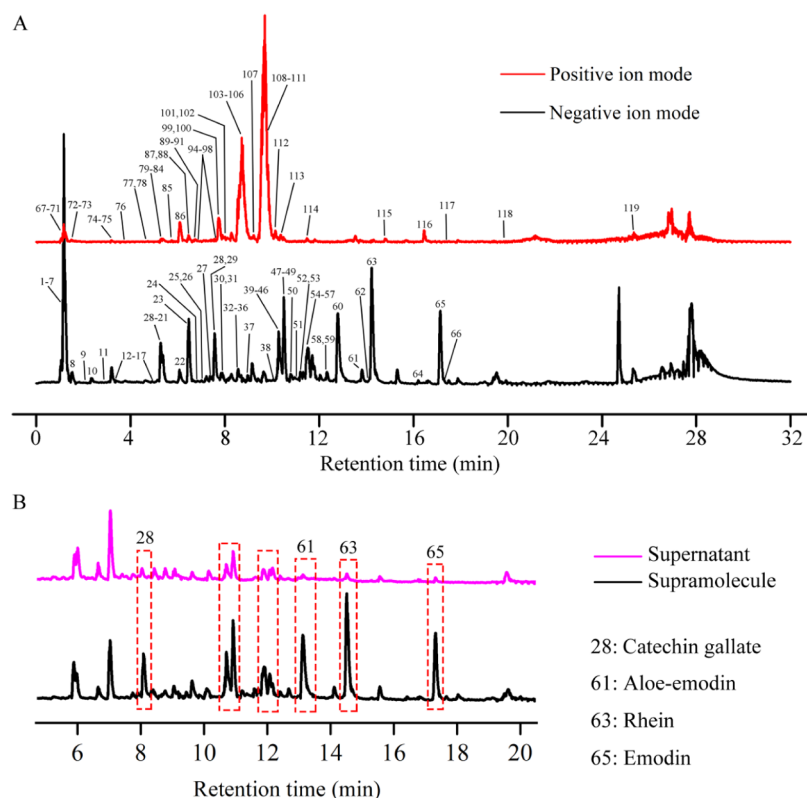


Figure 2. UPLC-MS/MS characterization. (A) Total ion current chromatograms of the supramolecular site in the positive and negative ion modes. (B) Negative ion current chromatograms of supramolecular and supernatant sites.

temperature of 25 °C, and titration interval 180 s (Nano ITC, TA).

3. RESULTS AND DISCUSSION

3.1. Morphological Characterization. As shown in Figure 1A,B, the co-decoction of *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* was turbid. Dynamic light scattering characterization showed that the co-decoction was a homogeneous and stable system with a size of 341.2 nm (PDI = 0.304) and a ζ -potential of -27.0 mV. The Tyndall effect also had a distinct pathway. The co-decoction was centrifuged at 80,000 rpm for 30 min, and then the supernatant and supramolecular site were collected, respectively. After centrifuging, no significant Tyndall effect was observed in the supernatant and its particle size was about 185 nm with uneven dispersion (PDI = 0.483). The collected supramolecular samples were redissolved in water, and it was found that the solution also had a significant Tyndall effect. The particle size of the supramolecular site was mainly 262 nm (PDI = 0.322), and the ζ -potential value was -27.8 mV.

SEM characterization is shown in Figure 1C; it was observed that the co-decoction mainly contained spherical nanoparticles of 300–500 nm. Rare nanoparticles were observed in the supernatant and their scale distribution was relatively discrete, which was consistent with the DLS characterization. Meanwhile, the supramolecular portion comprised nanospheres of about 300 nm with uniform size and uniform dispersion. SEM results showed that the nanospheres in the co-decoction mainly existed in the supramolecular part.

3.2. Identification of the Constituents in the Supramolecule. Totally 119 compounds had been tentatively identified in the supramolecular portion (Figure 2A). A total of

23 anthraquinones were identified, including 10 free anthraquinones, 11 bound anthraquinones, and 2 bound anthraones. They were all characteristic constituents of *Rhei Radix et Rhizoma* and were more likely to produce fragment ions such as $[M-H-CO]^-$ and $[M-H-CO_2]^-$ in negative ion mode. Compounds 63 and 65 were characterized to be rhein and emodin, respectively. Compound 63 (rhein) exhibited a molecular ion at m/z 283.0237 and its fragment ions were $[M-H-CO_2]^-$ (m/z 239.0347), $[M-H-C_2O_3]^-$ (m/z 211.0394), and $[M-H-C_3O_4]^-$ (m/z 183.0443). Likewise, emodin was produced as a precursor ion at m/z 269.0444 ($[M-H]^-$). The fragment ion at m/z 241.0502 ($[M-H-CO]^-$), m/z 225.0555 ($[M-H-CO_2]^-$), m/z 197.0601 ($[M-H-C_2O_3]^-$), and m/z 181.0648 ($[M-H-C_2O_4]^-$) originated from the loss of CO moiety and CO_2 moiety (Figure 3).

Alkaloids were the main medicinal ingredients of *Coptidis Rhizoma*. In this test, 31 alkaloids and their glycoside compounds were detected. For example, Compound 106 (coptisine) produced a precursor ion at m/z 320.0917 ($[M]^+$), yielding the most intense fragment ion at m/z 292.0966 ($[M-CO]^+$). Compound 111 was proposed to be berberine, showing a deprotonated molecule at m/z 336.1230 ($[M]^+$). The major fragment ion at m/z 320.0917 ($[M-CH_4]^+$) and m/z 306.0764 ($[M-CH_4-CH_2]^+$) is attributed to the continuous loss of a CH_4 moiety and a CH_2 moiety. Other fragments at m/z 321.0997 ($[M-CH_3]^+$) and m/z 292.0969 ($[M-C_2H_4O]^+$) could also be observed in the product ion spectrum. In addition to anthraquinones and alkaloids, there were 24 organic acids, 8 tannins, 8 flavonoids, 5 amino acids, and 7 monosaccharides and polysaccharides in the supramolecular site. Furthermore, there were 13 other components, including phenylbutanone, stilbene glycosides, naphthalene glycosides,

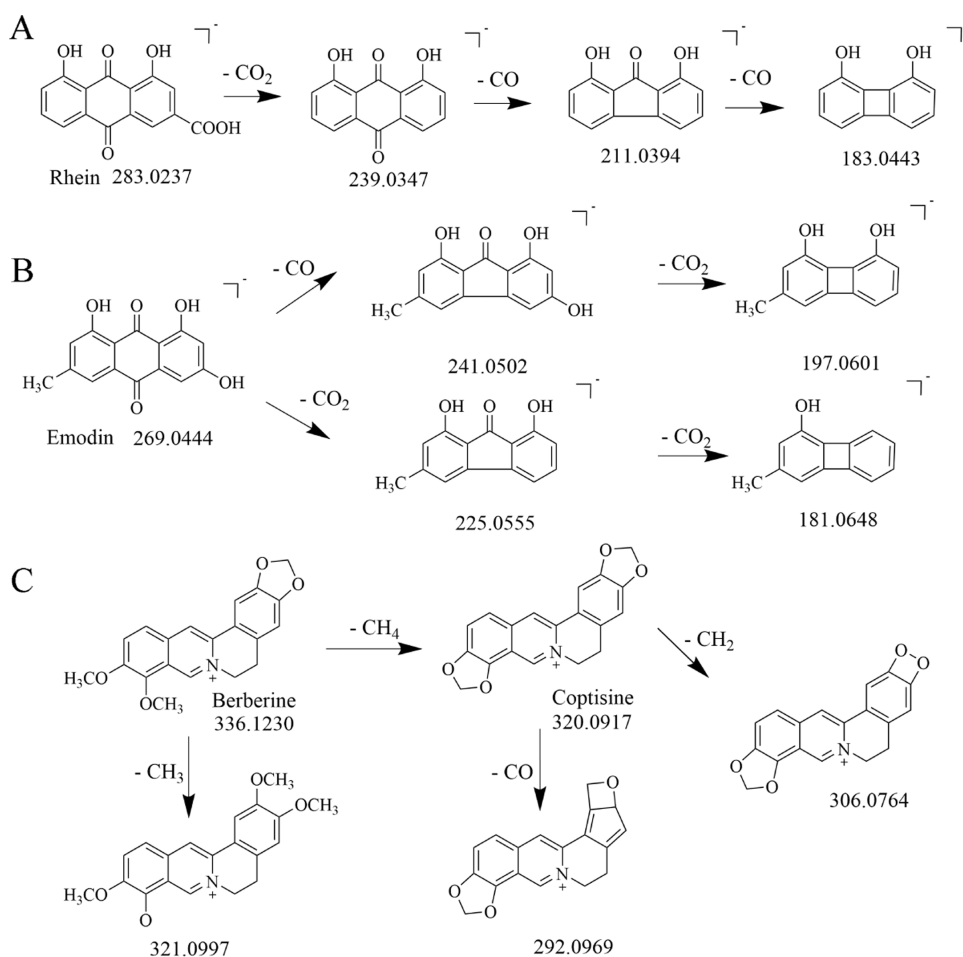


Figure 3. Cracking law of (A) rhein, (B) emodin, and (C) berberine and coptisine.

and other compounds. The fragment ions and cleavage pathways of some compounds are shown in Table 1, Table S1, and Figure S1.

The contents of anthraquinone and tannin in the supra-molecular fraction were much higher than those in the supernatant fraction (Figure 2B). Among them, the content of catechin gallate, 3-methyl-rhein, rhein, and emodin in the supramolecular site accounted for 80 to 90 percent (Table 2), which was much higher than that in the supernatant. It was further speculated that the components of *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* interacted with each other during the decocting process, and the products of interaction mainly existed in the supramolecular site but not in the supernatant site.

3.3. Isothermal Titration Calorimetry Analysis between the Herbal Pair Medicine. Binding constant (K_a), dissociation constant (K_b), Gibbs free energy (ΔG), enthalpy (ΔH), and entropy (ΔS) could be directly and accurately determined from a single experiment.^{57,58} ITC has been used by some researchers to analyze the thermodynamic parameters of interaction during co-decocting of medicinal materials.^{59,60} As shown in Figure 4, the upward peaks of *Coptidis Rhizoma*-titrated *Rhei Radix et Rhizoma* indicated an exothermic reaction. In the control group, *Coptidis Rhizoma*-titrated deionized water presented a downward endothermic characteristic peak (dilution process). The fitted curve showed an approximate S-shape, which was a typical characteristic peak trend of chemical reaction. The results showed that the

reaction between the two drugs was thorough, and the effect was strong. Their binding constant K_a was 4.885×10^3 .

Herein, thermodynamic parameters of titration between the herbal pair medicine were $\Delta H = -52.64 \text{ kJ/mol} < 0$, $\Delta S = -105.9 \text{ J/mol} < 0$. According to the second law of thermodynamics, $|\Delta H| > |-T\Delta S|$, $\Delta G = -23.713 \text{ kJ/mol} < 0$. It was suggested that the reaction was a spontaneous reaction driven by enthalpy, and the entropy change was unfavorable to the reaction. The results showed that the active components of *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* were bonded by noncovalent bonds, such as hydrogen bonds and electrostatic attraction, rather than aggregation driven by physical precipitation.

3.4. Morphological Characterization of Emo-Cop NPs and Rhe-Cop NFs. Based on the results shown in Figure 2B, emodin, rhein, and coptidine were selected as the research objects to further reveal the carrier-free self-assembly phenomenon in this drug pair. The prepared Emo-Cop NPs and Rhe-Cop NFs were characterized by SEM (Figure 5). It was found that the morphology of the assembly formed by different components was significantly different. The Emo-Cop NPs were a nanosphere with a particle size between 50 and 100 nm and evenly dispersed in the field of vision. Some scholars have found that protonated rhein tends to aggregate, and so spherical, columnar, and membrane-shaped aggregated micelles could be observed.⁶¹ In contrast, the Rhe-Cop NFs presented a fibrous surround state with a fiber width of about 100 nm. By comparing the morphology of the two kinds of

Table 1. Identification of Compounds at Supramolecular Sites

no.	t_R (min)	compound	formula	identity	precursor ion		Δppm	fragment ion (m/z)
					theoretical (m/z)	experimental (m/z)		
9	2.01	galloylglucose	C ₁₃ H ₁₆ O ₁₀	[M-H] ⁻	331.0659	331.0674	4.53	169.0132 [M-H-C ₆ H ₁₀ O ₅] ⁻ , 125.0232 [M-H-C ₇ H ₁₀ O ₇] ⁻
10	2.34	gallic acid	C ₇ H ₆ O ₅	[M-H] ⁻	169.0131	169.0133	1.18	125.0231 [M-H-CO ₂] ⁻
27	7.37	rhein glucoside	C ₂₁ H ₁₈ O ₁₁	[M-H] ⁻	445.0765	445.0775	2.25	283.0250 [M-H-C ₆ H ₁₀ O ₅] ⁻ , 239.0348 [M-H-C ₇ H ₁₀ O ₇] ⁻ , 211.0396 [M-H-C ₈ H ₁₀ O ₈] ⁻
29	7.65	sennoside C/D	C ₄₂ H ₄₀ O ₁₉	[M-H] ⁻	847.2080	847.2097	2.00	685.1560 [M-H-C ₆ H ₁₀ O ₅] ⁻
31	7.85	sennoside A/B	C ₄₂ H ₃₈ O ₂₀	[M-H] ⁻	861.1872	861.1890	2.09	699.1338 [M-H-C ₆ H ₁₀ O ₅] ⁻ , 386.0994 [M-H-C ₂₂ H ₁₉ O ₁₂] ⁻
35	8.70	aloesin	C ₁₉ H ₂₂ O ₉	[M-H] ⁻	393.1180	393.1196	4.07	231.0658 [M-H-C ₆ H ₁₀ O ₅] ⁻ , 203.0709 [M-H-C ₇ H ₁₀ O ₆] ⁻
36	8.86	aloe emodin glucoside	C ₂₁ H ₂₀ O ₁₁	[M-H] ⁻	431.0972	431.0984	2.78	269.0456 [M-H-C ₆ H ₁₀ O ₅] ⁻
44	10.30	emodin glucoside	C ₂₁ H ₂₀ O ₁₁	[M-H] ⁻	431.0972	431.0984	2.78	269.0456 [M-H-C ₆ H ₁₀ O ₅] ⁻
48	10.50	chrysophanol	C ₁₅ H ₁₀ O ₄	[M-H] ⁻	253.0495	253.0506	4.35	225.0555 [M-H-CO] ⁻
54	11.42	physcion	C ₁₆ H ₁₂ O ₅	[M-H] ⁻	283.0600	283.0612	4.24	268.0377 [M-H-CH ₃] ⁻ , 240.0425 [M-H-C ₂ H ₃ O] ⁻
55	11.47	physcion glucoside	C ₂₂ H ₂₂ O ₁₀	[M-H] ⁻	445.1129	445.1139	2.25	283.0612 [M-H-C ₆ H ₁₀ O ₅] ⁻ , 268.0377 [M-H-C ₇ H ₁₃ O ₃] ⁻ , 240.0425 [M-H-C ₈ H ₁₃ O ₆] ⁻
60	12.80	3-methyl-rhein	C ₁₆ H ₁₀ O ₆	[M-H] ⁻	297.0393	297.0406	4.37	253.0505 [M-H-CO ₂] ⁻ , 225.0553 [M-H-C ₂ O ₃] ⁻
61	13.82	aloe-emodin	C ₁₃ H ₁₀ O ₅	[M-H] ⁻	269.0444	269.0456	4.46	241.0498 [M-H-CO] ⁻ , 240.0426 [M-H-CHO] ⁻ , 223.0397 [M-H-CH ₂ O ₂] ⁻ , 197.0597 [M-H-C ₂ O ₃] ⁻ , 183.0449 [M-H-C ₃ H ₂ O ₃] ⁻
63	14.23	rhein	C ₁₅ H ₈ O ₆	[M-H] ⁻	283.0237	283.0249	4.24	239.0347 [M-H-CO ₂] ⁻ , 211.0394 [M-H-C ₂ O ₃] ⁻ , 183.0443 [M-H-C ₃ O ₄] ⁻ , 155.0492 [M-H-C ₄ O ₅] ⁻
65	17.14	emodin	C ₁₅ H ₁₀ O ₅	[M-H] ⁻	269.0444	269.0456	4.46	241.0502 [M-H-CO] ⁻ , 225.0552 [M-H-CO ₂] ⁻ , 197.0601 [M-H-C ₂ O ₃] ⁻ , 181.0648 [M-H-C ₂ O ₄] ⁻
86	6.08	magnoflorine	C ₂₀ H ₂₄ NO ₄ ⁺	[M] ⁺	342.1699	342.1700	0.29	297.1122 [M-C ₂ H ₄ N] ⁺ , 282.0886 [M-C ₃ H ₁₀ N] ⁺ , 265.0860 [M-C ₃ H ₁₁ NO] ⁺ , 237.0909 [M-C ₄ H ₁₁ NO ₂] ⁺
97	7.58	berberrubine	C ₁₉ H ₁₆ NO ₄ ⁺	[M] ⁺	322.1073	322.1073	0.00	307.0839 [M-CH ₃] ⁺ , 294.1124 [M-CO] ⁺
103	8.55	columbamine	C ₂₀ H ₂₀ NO ₄ ⁺	[M] ⁺	338.1386	338.1387	0.29	323.1352 [M-CH ₃] ⁺ , 322.1074 [M-CH ₄] ⁺ , 308.0917 [M-C ₂ H ₆] ⁺ , 306.1130 [M-C ₂ H ₈] ⁺ , 294.1124 [M-C ₂ H ₄ O] ⁺
104	8.64	epiberberine	C ₂₀ H ₁₈ NO ₄ ⁺	[M] ⁺	336.1230	336.1229	-0.30	321.0991 [M-CH ₃] ⁺ , 320.0918 [M-CH ₄] ⁺ , 306.0753 [M-C ₂ H ₆] ⁺ , 292.0966 [M-C ₂ H ₄ O] ⁺
105	8.69	jatrorrhizine	C ₂₀ H ₂₀ NO ₄ ⁺	[M] ⁺	338.1386	338.1385	-0.30	323.1352 [M-CH ₃] ⁺ , 322.1073 [M-CH ₄] ⁺ , 308.0916 [M-C ₂ H ₆] ⁺ , 306.1127 [M-C ₂ H ₈] ⁺ , 294.1124 [M-C ₂ H ₄ O] ⁺
106	8.75	coptisine	C ₁₉ H ₁₄ NO ₄ ⁺	[M] ⁺	320.0917	320.0918	0.31	292.0966 [M-CO] ⁺
109	9.44	worenine	C ₂₀ H ₁₆ NO ₄ ⁺	[M] ⁺	334.1073	334.1073	0.00	319.0843 [M-CH ₃] ⁺ , 306.1122 [M-CH ₄] ⁺
110	9.55	palmatine	C ₂₁ H ₂₂ NO ₄ ⁺	[M] ⁺	352.1543	352.1542	-0.28	337.1307 [M-CH ₃] ⁺ , 336.1229 [M-CH ₄] ⁺ , 322.1074 [M-C ₂ H ₆] ⁺ , 308.1281 [M-C ₃ H ₈] ⁺
111	9.67	berberine	C ₂₀ H ₁₈ NO ₄ ⁺	[M] ⁺	336.1230	336.1230	0.00	321.0997 [M-CH ₃] ⁺ , 320.0919 [M-CH ₄] ⁺ , 306.0764 [M-C ₂ H ₆] ⁺ , 292.0969 [M-C ₂ H ₄ O] ⁺ , 278.0805 [M-C ₃ H ₆ O] ⁺

Table 2. Proportion of Partial Components in Supramolecular and Supernatant Sites

no.	compound	peak area (mAU)		proportion (%)	
		supernatant	supramolecule	supernatant	supramolecule
28	catechin gallate	150697588	710382109	17.50	82.50
61	3-methyl-rhein	104630117	1060477996	8.98	91.02
63	rhein	126177173	1781611708	6.62	93.38
65	emodin	84962980	1080546999	7.29	92.71
103–106	alkaloids	24427908772	32315248803	43.05	56.95
108–111		49331489593	68356178573	41.92	58.08

self-assembly, the Tyndall Effect and aggregations in the decoction of *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* were confirmed.

3.5. Isothermal Titration Calorimetry Analysis. To further reveal the self-assembly mechanism of the herbal pair medicine, we found that the titration process of emodin and coptisine, rhein and coptisine was an exothermic reaction. As shown in Figure 4B,C, Gibbs free energy was -22.61 and

-27.59 kJ/mol, respectively. Both of them displayed $|\Delta H| > |T\Delta S|$. These results indicated that the reaction of emodin and coptisine, rhein and coptisine was a spontaneous chemical reaction driven by enthalpy, rather than a simple physical process. Current evidence further proved that the assembly process was driven by weak interactions, which was in accordance with the reaction of the herbal pair medicine (Figure 4A).

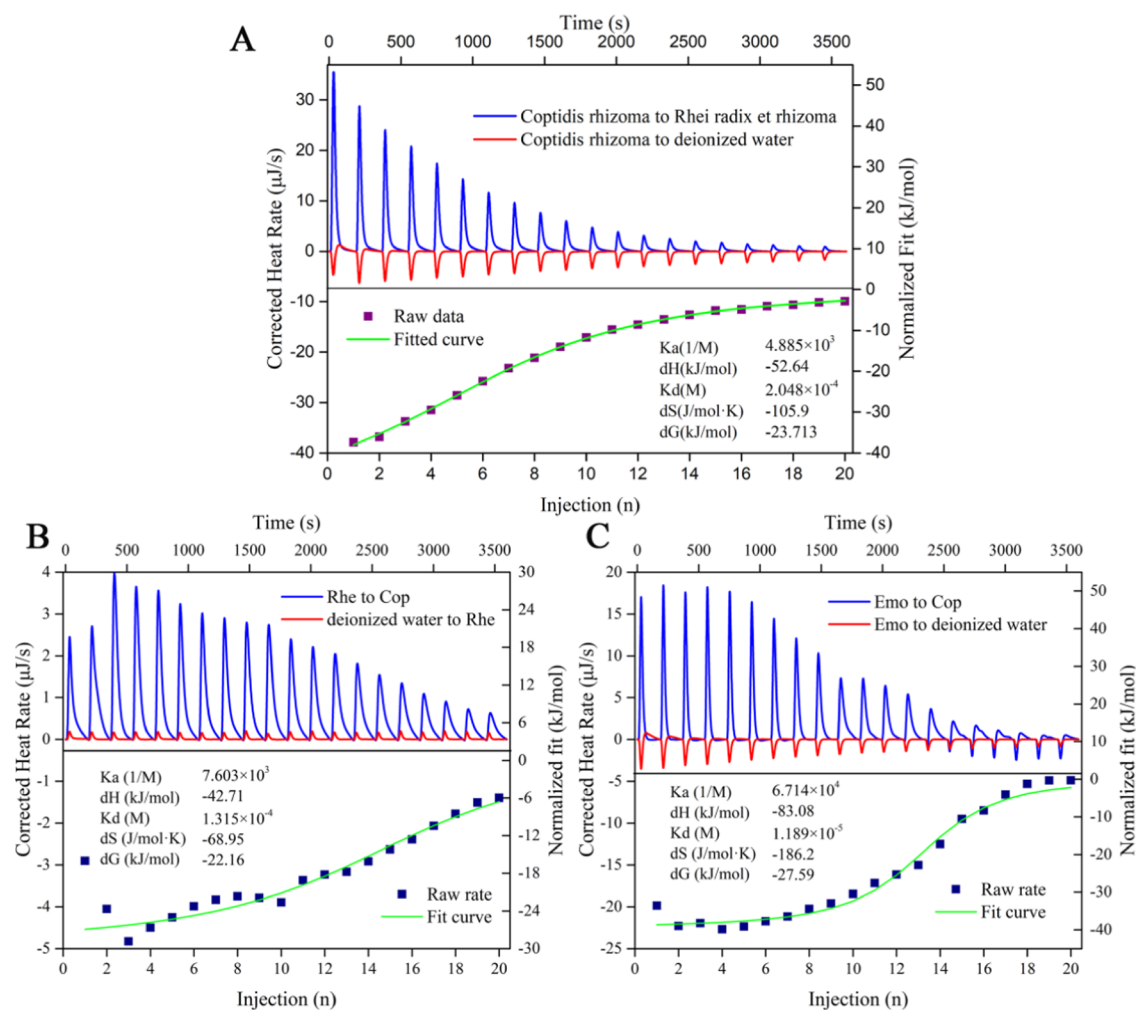


Figure 4. ITC characterization: (A) *Coptidis Rhizoma*-titrated *Rhei Radix et Rhizoma*. (B) Rhe-titrated Cop. (C) Emo-titrated Cop.

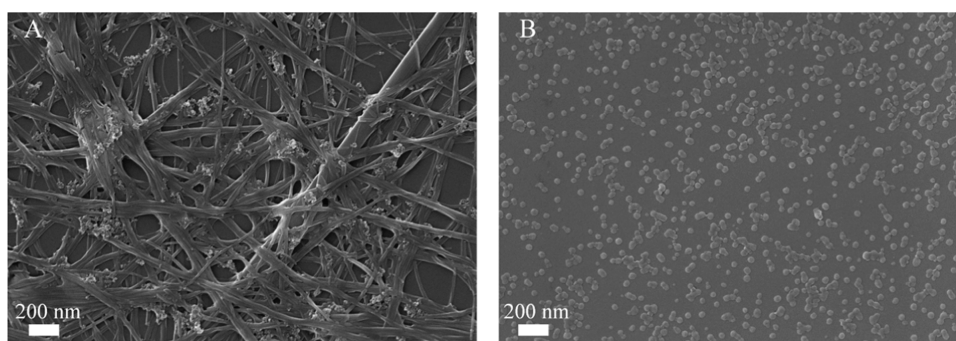


Figure 5. Scanning electron microscopy characterization: (A) Rhe-Cop NFs and (B) Emo-Cop NPs.

3.6. UV–Vis Absorption Spectral Analysis. UV–vis absorption spectra reflected the characteristics of some groups in molecules.⁶² After molecular interaction, the change in the benzene-conjugated system resulted in the shift of the maximum absorption wavelength. Therefore, the law of molecular interaction or the process of self-assembly could be inferred from the changes in UV–vis spectra.^{63,64}

There were five characteristic absorption peaks of coptisine at 226, 242, 266, 360, and 460 nm. The characteristic absorption peaks of rhein were at 204, 230, 258 nm ($\pi \rightarrow \pi^*$ transitions), and 430 nm ($n \rightarrow \sigma^*$ transitions). The characteristic absorption peaks of emodin were at 222 nm (n

$\rightarrow \sigma^*$ transitions), 266, 286 nm ($\pi \rightarrow \pi^*$ transitions), and 442 nm ($n \rightarrow \sigma^*$ transitions). Both Rhe-Cop NFs and Emo-Cop NPs had the characteristic absorption peak of coptisine at 360 nm. Rhe-Cop NFs had the characteristic absorption peaks of rhein at 204, 228, and 268 nm. The characteristic absorption peak at 434 nm of Rhe-Cop NFs showed a blue shift compared with coptisine (from 462 to 434 nm). Compared with emodin, Emo-Cop NPs had characteristic absorption peaks at 244 and 264 nm, while the characteristic absorption peaks of emodin were at 266 and 286 nm (Figure 6A,B).

Emodin, rhein (anthraquinones), and coptisine (benzyl isoquinoline alkaloid) all had large cyclic conjugated structures.

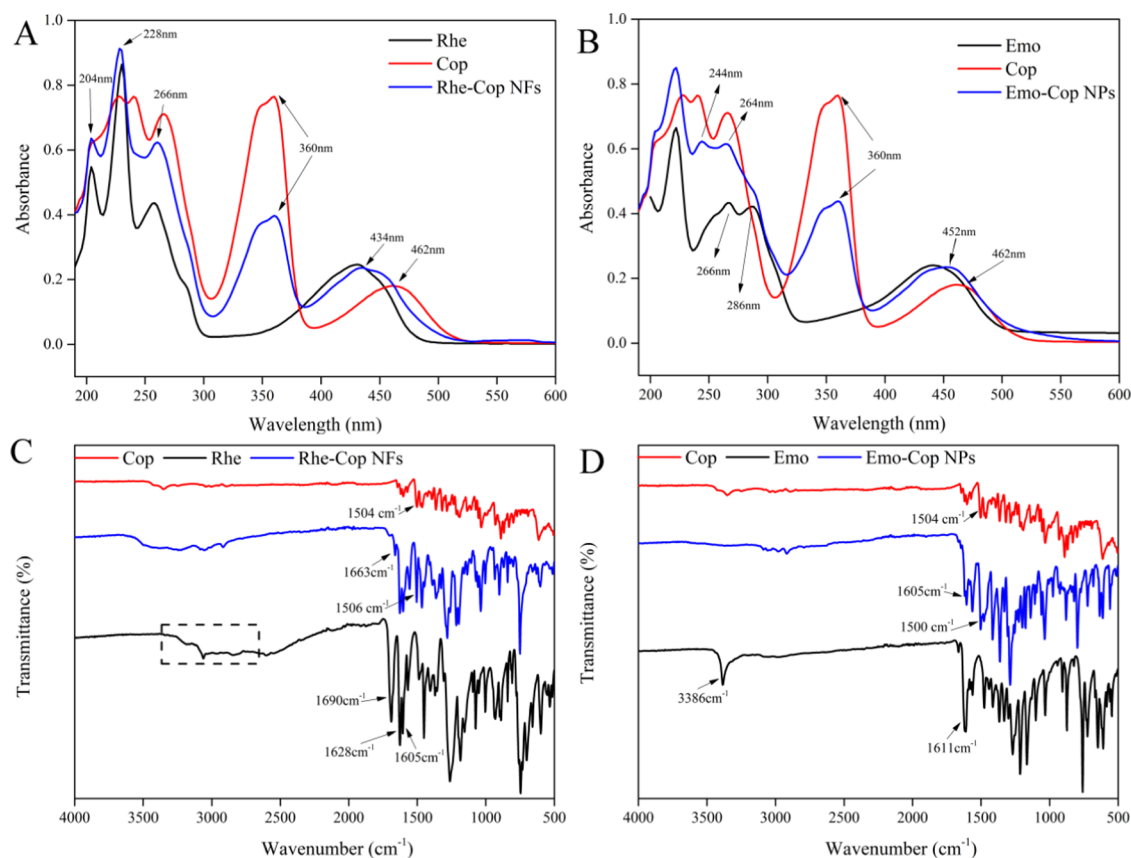


Figure 6. (A) UV–visible spectra of Rhe-Cop NFs. (B) UV–visible spectra of Emo-Cop NPs. (C) Fourier infrared spectra of Rhe-Cop NFs. (D) Fourier infrared spectra of Emo-Cop NPs.

The hydroxyl group and methylene dioxy group of benzene ring as the chromophore made the two kinds of components to appear with strong characteristic absorption in the UV–vis spectrum. These results indicated that the assemblies had characteristic absorption peaks of monomer components and the characteristic peak shift of the assemblies mostly occurred at 260 and 430 nm. It was speculated that the weak intermolecular interaction affected the $n-\pi^*$ and $\pi-\pi^*$ transitions of intramolecular electrons.^{65,66}

3.7. Fourier Transform Infrared Spectroscopy (FTIR) Characterization Analysis. FTIR was often dominated by the overlap of the doubling frequency and merging frequency of the hydrogen group X–H (X = C, N, O). Because the absorption frequency of different functional groups was different, they were in different positions in the spectrum.^{67,68}

In the infrared spectrum for Rhe-Cop NFs, shown in Figure 6C, the OH stretching vibration band of the COOH of Rhe-Cop NFs appeared at a high wavenumber compared with that in rhein (3000 cm⁻¹). The C=O group vibration band of the COOH group of rhein shifted from 1690 to 1663 cm⁻¹ and the peak intensity decreased significantly. The band at 1504 cm⁻¹ in coptisine shifted to a higher wavenumber of 1506 cm⁻¹ and the peak band intensity increased. The bands 1628 and 1605 cm⁻¹ in the assembly were attributed to the stretching vibration band of the C=O group on rhein's quinone ring. In the infrared spectrum for Emo-Cop NPs, shown in Figure 6D, the 3386 cm⁻¹ stretching vibration absorption band of 3-OH of emodin disappeared. The C=O group stretching vibration band on the quinone ring of emodin shifted from 1611 to 1605 cm⁻¹ and the peak intensity decreased. The 1504

cm⁻¹ infrared characteristic band of coptisine was an aromatic ring skeleton vibration band, which shifted to a low wavenumber of 1500 cm⁻¹ and the peak intensity increased.

These results indicated that the 3-COOH band on the anthraquinone ring of emodin, the 3-OH band on the anthraquinone ring of emodin, and the aromatic ring skeleton of coptisine significantly changed after assembly, suggesting that there were hydrogen bonding or $\pi-\pi$ stacking between rhein and coptisine, emodin and coptisine, respectively.⁶⁹ Therefore, it was inferred that the interaction between 3-COOH of rhein, 3-OH of emodin, and the quaternary ammonium nitrogen atom of coptisine led to electron delocalization, which reduced the density of the electron cloud and showed that the C=O bond energy of anthraquinone components become weaker and the stretching vibration frequency becomes lower than their single molecules.

3.8. ¹H NMR Characterization Analysis. ¹H NMR spectra could be used to characterize weak interactions such as hydrogen bonding and $\pi-\pi$ stacking.⁷⁰ Through weak interactions, molecules were stacked and arranged regularly to form assemblies or supramolecules with a certain microscopic morphology. Changes in the spatial position of the molecule affected the density of the electron cloud on the hydrogen atom, leading to fluctuations in the hydrogen signal. Although the fluctuation was not as violent as the reaction of chemical synthesis, it could also accurately determine the site where the weak bond binds to each other.⁷¹ Anthracene had three linear benzene rings, which could generate $\pi-\pi$ stacking or charge transfer with other electron-rich or electron-poor molecules through π -conjugated groups.⁷² For example, rhein molecules

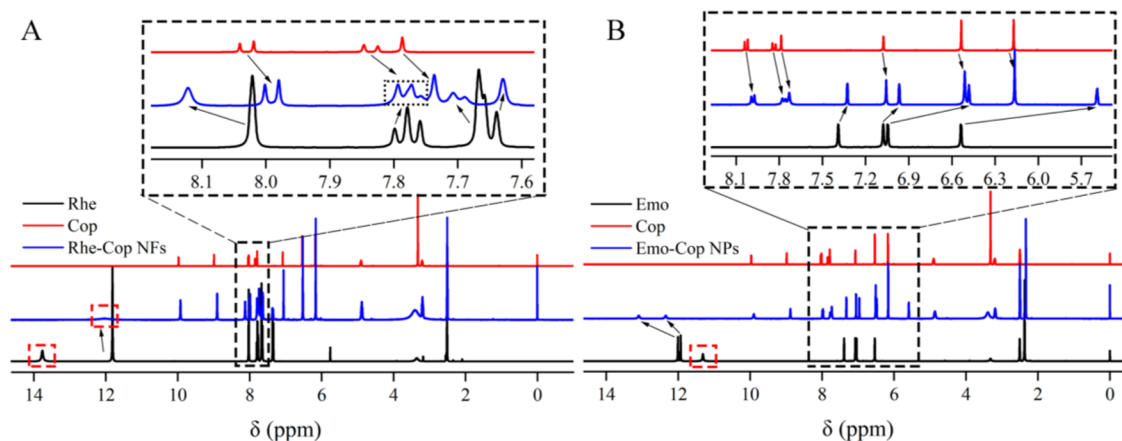


Figure 7. Nuclear magnetic hydrogen spectra: (A) Rhe-Cop NFs. (B) Emo-Cop NPs.

Table 3. Chemical Shifts of Emo-Cop NPs, Rhe-Cop NFs, and Their Anthraquinone Monomers

Rhe	Rhe-Cop NFs	$\Delta\delta$ (ppm)	Emo	Emo-Cop NPs	$\Delta\delta$ (ppm)
13.76 (3-COOH, s)	disappear		12.01 (1-OH, s)	13.10 (s)	+1.09
11.81 (1,8-OH, s)	12.02 (brs)	+0.11	11.93 (8-OH, s)	12.35 (s)	+0.42
8.02 (H-4, s)	8.12 (s)	+0.10	11.31 (3-OH, s)	disappear	
7.78 (H-6, t, $J = 8$ Hz)	7.77 (t, $J = 8$ Hz, $J = 4$ Hz)	-0.01	7.39 (H-5, s)	7.33 (d, $J = 0.8$ Hz)	-0.06
7.67 (H-2, s)	7.63 (s)	-0.04	7.08 (H-7, s)	6.97 (m)	-0.11
7.65 (H-5, d, $J = 8$ Hz)	7.70 (d, $J = 4$ Hz)	+0.05	7.05 (H-4, d, $J = 4$ Hz)	6.48 (m)	-0.57
7.35 (H-7, d, $J = 8$ Hz)	7.35 (d, $J = 8$ Hz)		6.54 (H-2, d, $J = 4$ Hz)	5.59 (m)	-0.95
			2.32 (6-CH ₃ , s)	2.34 (s)	+0.02

Table 4. Chemical Shifts of Emo-Cop NFs, Rhe-Cop NFs, and Coptisine

Cop	Rhe-Cop NFs	$\Delta\delta$ (ppm)	Emo-Cop NPs	$\Delta\delta$ (ppm)
9.97 (H-8, s)	9.92 (s)	-0.05	9.90 (s)	-0.07
8.99 (H-13, s)	8.90 (s)	-0.09	8.88 (s)	-0.11
8.03 (H-11, d, $J = 8$ Hz)	7.99 (d, $J = 8$ Hz)	-0.04	7.98 (d, $J = 8$ Hz)	-0.05
7.84 (H-12, d, $J = 8$ Hz)	7.77 (t, $J = 8$ Hz, $J = 4$ Hz)	-0.07	7.76 (t, $J = 8$ Hz, $J = 12$ Hz)	-0.08
7.79 (H-1, s)	7.74 (s)	-0.05		-0.03
7.08 (H-4, s)	7.05 (s)	-0.03	7.05 (s)	-0.03
6.54 (15-CH ₂ , s)	6.52 (s)	-0.02	6.51 (s)	-0.03
6.17 (14-CH ₂ , s)	6.16 (s)	-0.01	6.16 (s)	-0.01
4.90 (6-CH ₂ , t, $J = 8$ Hz, $J = 4$ Hz)	4.88 (t, $J = 8$ Hz, $J = 4$ Hz)	-0.02	4.87 (s)	-0.03
3.20 (5-CH ₂ , t, $J = 8$ Hz, $J = 4$ Hz)	3.19 (t, $J = 8$ Hz, $J = 4$ Hz)	-0.01	3.19 (s)	-0.01

could self-assemble to form three-dimensional networks composed of nanofibers through noncovalent interactions such as π - π stacking and hydrogen bonding,¹⁶ while emodin molecules are stacked into a supramolecular layer through π - π interactions.⁷³

According to the data analysis in Figure 7 and Tables 3–1, the characteristic peaks of 3-COOH of rhein and 3-OH of emodin disappeared, which indicated that rhein and coptisine, emodin and coptisine approached each other under the induction of electrostatic attraction (anthraquinone's 3-COO⁻ and alkaloids' 7-N⁺, anthraquinone's 3-O⁻ and alkaloids' s 7-N⁺). It was also found that the 1,8-OH on both anthraquinone rings fluctuated to the low field direction with large chemical shift values. Therefore, it could be shown that there were obvious hydrogen bonds between rhein molecules and emodin molecules, which formed a layered skeleton structure under the action of intermolecular hydrogen bonds. The π - π stacking between rhein and coptisine was distributed between the anthraquinone layers, resulting in the H-8, H-11, H-12, and H-13 on the benzyl isoquinoline ring moving 0.04–0.11 ppm to

the high field. The action of emodin and coptidine was the same as that of rhein and coptisine. Because the skeletal structure of coptidine was similar to that of berberine, the crystal structure of rhein-berberine²⁴ and emodin-berberine⁷³ could be inferred: rhein and coptisine, emodin and coptisine were close to each other under electrostatic action and further formed binary assemblies under the action of π - π stacking.

4. CONCLUSIONS

In this study, spherical supramolecules were found in *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* decoction. A total of 119 compounds were identified from the supramolecules by the UPLC-MS/MS method (23 anthraquinones, 31 alkaloids, 24 organic acids, 8 tannins, and other components). Among these, the content of anthraquinones and tannins in the supramolecular was much higher than that in the supernatant, such as rhein and emodin. ITC analysis showed that the interaction between *Rhei Radix et Rhizoma* and *Coptidis Rhizoma* was a spontaneous chemical reaction, indicating that active components could form supramolecules through weak

bonds. Furthermore, we found that the ingredients of emodin and coptisine could be assembled into NPs of about 50 nm, while rhein and coptidine were assembled into nanofibers. Based on the current evidence, both self-assemblies were mainly induced by electrostatic attraction, hydrogen bonding, and π - π stacking. In this study, the self-assembled entities of small molecules were found based on the herbal pair medicine, which provides a reference for discovery of the natural carrier-free self-assembly from TCM. And it is also beneficial to the design of binary self-assembly of small molecules in the future.

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acsomega.2c04098>.

UPLC-MS/MS analysis of supramolecular sites and schematic diagram of the cleavage law of the ingredients (PDF)

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Notes

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