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Study on extraction methods of polysaccharides from a processed product of *Aconitum carmichaeli* Debx.

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Traditional Chinese medicine PaoTianXiong (PTX) is a processed product of *Aconitum carmichaeli* Debx. with polysaccharide as the main ingredient. The properties of PTX polysaccharide (PTXP) may be affected by different extraction methods. To develop and utilize PTXP better, it is of great significance to study the extraction methods of PTXP. Thus, we extracted PTXPs with dilute alkaline water extraction, ultrasound-assisted extraction, cellulase-assisted extraction, and hot water extraction (HWE), respectively. The characterizations of PTXPs extracted by different methods were analyzed based on purity determination, infrared analysis, molecular weight and monosaccharide composition. And antioxidant experiments of PTXPs were conducted. The results showed that PTXPs extracted by the four extraction methods were all glucan. After purification, the PTXPs showed similar antioxidant activity *in vitro*. The molecular weight of polysaccharides extracted by the cellulase-assisted method was different from that extracted by the other three methods. Our results showed that not only the yield but also the effect of extraction methods on the properties of PTXP should be considered when selecting the best extraction method. Therefore, HWE was considered to be the best extraction method of PTXP. The yield and purity of purified PTXP were 24.5% and 97.1%, respectively. The optimized extraction conditions were: an extraction temperature of 90 °C, extraction time of 2.17 h, solid–liquid ratio of 1 : 29 (g mL⁻¹), and number of extractions of 2.

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1 Introduction

Traditional Chinese medicine PaoTianXiong (PTX), one of the processed products of the tuber of *Aconitum carmichaeli* Debx., is characterized by main active alkaloids and polysaccharides. Processing, an ancient Chinese pharmaceutical technique (including roasting, baking, stir-frying, *etc.*), aims to reduce the content of toxic alkaloids in PTX.^{1,2} However, the complex processing process also results in different properties of polysaccharide of PTX (PTXP) extracted from the tuber of *Aconitum carmichaeli* Debx. In recent years, polysaccharides have attracted much attention because of their various activities, including anti-inflammatory, antioxidant, anti-tumor, immunomodulatory, and other activities.^{3–7} However, polysaccharides are difficult to synthesize artificially. A reliable extraction method is very important for the development, research and quality control of PTXP.

The extraction methods of polysaccharides mainly include hot water extraction (HWE),⁸ cellulase-assisted extraction

(CAE),⁹ ultrasonic-assisted extraction (UAE),¹⁰ dilute alkaline extraction (DAE),¹¹ *etc.* HWE works by letting water into plant cells to dissolve the polysaccharides which are then extracted by diffusing from a higher concentration to a lower concentration. UAE uses ultrasound to break down plant cell walls, thus accelerating the dissolution and diffusion of the polysaccharides. Different from the UAE, CAE uses enzymes to break down plant cell walls and speeds up the dissolution of polysaccharides. DAE had better extraction efficiency for acidic polysaccharides. Each method has its special advantages and disadvantages in terms of extraction efficiency, economic cost and environmental impact. Besides, according to the research on the structural modification of polysaccharides, the structure of polysaccharides can be changed,¹² which is commonly characterized by monosaccharide composition, infrared analysis and molecular weight.^{13,14} Based on different extraction principles, PTXP extracted by different extraction methods may have different properties. To the best of our knowledge, there is no information on the effect of different extraction methods on the properties of PTXP. In addition, the current research on the extraction technology of polysaccharides mainly takes the extraction yield of polysaccharides as the index but ignores the purity of polysaccharides.^{10,15,16} We held that the reported polysaccharide activity is not convincing without considering the purity of the polysaccharide. In order to better study,

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develop and use PTXP, it is very important to research and contrast the extraction methods of PTXP.

Therefore, in the present study, we extracted crude PTPs with CAE, HWE, DAE and UAE methods, respectively. To improve the purity of PTXP, anion exchange chromatography was used in the screening process of PTXP extraction methods for the first time. Then the characterizations of PTPs extracted by different methods were analyzed based on infrared analysis, molecular weight and monosaccharide composition. Except for information about the preliminary structural feature, we also conducted *in vitro* antioxidant experiments to compare the effects of different extraction methods on the activity of PTXP. Based on the above physical and chemical analysis methods, we finally screened the best extraction method of PTXP and then optimized it.

2 Materials and methods

2.1 Chemical and reagents

Cellulase was supported by Dalian Meilun Biological Technology Co., Ltd (Dalian, China). DEAE Sepharose™ fast flow was purchased from GE Healthcare Life Science (Piscataway, NJ, USA). Phenol, potassium persulfate, 1,1-diphenyl-2-picrylhydrazyl (DPPH) were purchased from Shanghai Macklin Biochemical Co., Ltd (Shanghai, China). Sodium hydroxide, ethanol, chloroform, *N*-butanol, sodium chloride and sulfuric acid were of analytical grade and were obtained from Guangzhou Reagent Co., Ltd (Guangzhou, China). 2,2'-Azino-bis(3-ethylbenzothiazoline-6-sulfonic acid)diammonium salt (ABTS) was purchased from Sigma Life Science (St. Louis, MO, USA). Rhamnose, L-arabinose, L-fucose, DL-xylose, D-mannose, D-glucose, and D-galactose were purchased from Shanghai Yuanye Bio-Technology Co. Ltd (Shanghai, China). All other chemicals and reagents used were of analytical grade.

2.2 Plant material

PTX (lot number YPA8C0001) was purchased from Guangzhou medicine company, Chinese medicine Yinpian factory. The preparation method of PTX adopted the unique preparation method of traditional Chinese medicine. Briefly, the tuber of *Aconitum carmichaeli* Debx. was soaked in sodium chloride solution and then was dried in the sun. This process was repeated until sodium chloride crystals appeared on the surface of the tuber of *Aconitum carmichaeli* Debx. Subsequently, 1200 mL water was added to 1000 g of the product, and refreshed the water twice a day for a total of five days. Then the product was soaked in 1000 mL water extract of ginger and steamed in a pressure cooker for 1.5 h. After drying at 80 °C, it was stir-fried with hot sand at 210–230 °C until it become brown. The obtained PTX was powdered and sifted through no. 65 mesh before use.

2.3 Extraction and purification of PTPs

2.3.1 Hot water extraction. HWE was performed based on the reported method with some modifications.¹⁷ 10 g of PTX powder was extracted with hot water at 90 °C for 3 times, 2 h each time, according to the solid to liquid ratio of 1 : 15 (g mL⁻¹).

2.3.2 Ultrasonic-assisted extraction. UAE was based on our previous research.¹⁸ In brief, 10 g of PTX powder was firstly extracted with the help of ultrasound and then bathed in water at 80 °C for 3 h. The extracted solution was filtered and stored at 4 °C for later use. The conditions of ultrasonic-assisted extraction were as follows: the power of ultrasonics was 270 W, extraction time was 13 min, and the solid–liquid ratio was 1 : 25 (g mL⁻¹).

2.3.3 Cellulase-assist extraction. CAE of PTXP was conducted according to the reported method with some modifications.⁹ 10 g of PTX was added with pure water at the solid–liquid ratio of 1 : 25 (g : mL), and then pH was adjusted to 4.5 with hydrochloric acid. 150 mg of cellulase was added at the dosage of 6000 U g⁻¹ and extracted at 55 °C for 3 h.

2.3.4 Dilute alkaline extraction. DAE of PTXP was referred to the literature with some modifications.¹⁹ 10 g of PTX powder was added into 0.09 mol L⁻¹ NaOH solution according to the solid–liquid ratio of 1 : 21.2 (g : mL), and then reflux extraction was conducted at 90 °C for 0.5 h.

The PTXP solution extracted by above methods were concentrated to 100 mL, centrifuged at 3000 rpm for 10 min, then anhydrous ethanol was added until the ethanol content was 80%, and the precipitation of PTXP was collected by centrifugation. The precipitated PTXP was dissolved in water and precipitated again with ethanol for another 2 times, and then the protein was removed by the Sevage method. After dialysis and freeze-drying, we obtained four kinds of crude PTXP, named HWE-PTXP, UAE-PTXP, CAE-PTXP and DAE-PTXP.

2.3.5 Purification of crude PTPs. The crude polysaccharide was applied to a DEAE Fast Flow anion exchange column (50 cm × 5.5 cm) at a concentration of 3% (w/v) and then was eluted with 0.4 mol L⁻¹ NaCl solution. The eluent was collected by automatic collector and detected by phenol-sulfuric acid method. (The elution of 0.04 mol L⁻¹ NaCl solution was determined by the preliminary experiment, and the concentration could elute all polysaccharides without eluting impurities). The eluent was concentrated, dialyzed with distilled water for 48 hours and then freeze-dried. Therefore, we obtained purified PTPs, named PHWE-PTXP, PUAE-PTXP, PCAE-PTXP and PDAE-PTXP.

2.4 Yield and purity determination

The extraction yield (W_1) and purity (W_3) of crude PTXP were calculated respectively. After purification by anion exchange chromatography, we calculated the extraction yield (W_2) and purity (W_4) of purified PTPs. W_4 was determined by the phenol-sulfate method.¹⁶ The standard curve equation is $Y = 0.0052X + 0.1939$ ($R^2 = 0.990$). Where X is the concentration of PTXP samples and Y is the absorbance. The equations of W_1 , W_2 and W_3 were as follows:

$$\begin{aligned} W_1 &= \frac{m_1}{m_2} \times 100\%, & W_2 &= \frac{m_1}{m_2} \times \frac{m_3}{m_4} \times 100\%, & W_3 \\ &= \frac{m_3}{m_4} \times 100\%, & W_4 &= \frac{C_1}{C_0} \times 0.9 \times 100\% \end{aligned}$$



where m_1 is the mass of crude PTXP obtained, m_2 is the mass of extracted PTX, m_3 is the mass of purified PTXP, m_4 is the mass of crude PTXP used for purification, C_0 is the configured sample concentration of purified PTXP, and C_1 is the concentration of PTXP calculated by the standard curve. In the formula of W_4 , 0.9 is the conversion factor.

2.5 Determination of molecular weight

The molecular weight of PTPs was determined by high performance gel permeation chromatography (HPGPC) using a Waters instrument equipped with tandem Ultrahydrogel 1000 column (300 mm \times 7.8 mm i.d., 12 μ m) and Ultrahydrogel 500 column (300 mm \times 7.8 mm i.d., 10 μ m). The samples were dissolved in the mobile phase at 2 mg mL⁻¹, respectively. The mobile phase was 0.02 mol L⁻¹ potassium dihydrogen phosphate at the flow rate of 0.8 mL min⁻¹. The column temperature was 35 °C and the injection volume was 20 μ L.

2.6 FTIR-ATR analysis

The FTIR-ATR spectra of purified PTPs were recorded in the range of 600–4000 cm⁻¹ on a Fourier transform spectrometer in combination with a KBr beam splitter. Measurements have been performed in dry atmosphere to avoid dirty contributions.²⁰ The experimental parameters setting of the resolution was 4 cm⁻¹ and the number of the scans was 16 times.

2.7 Monosaccharide composition

Under the protection of N₂, 10 mg of purified PTPs were hydrolyzed with 4 mL of 2 mol L⁻¹ trifluoroacetic acid at 120 °C, respectively. Acetylation was then carried out with 10 mg of hydroxylamine hydrochloride and 1 mL of pyridine for 40 min at 90 °C. When the reaction solution was cooled, 1 mL of acetic anhydride was added and heated continuously for 40 min. The acetate derivative was analyzed by GC/MS (Agilent 7890A/5975C, USA) method with a HP-5 capillary column (30 nm \times 0.32 mm i.d., film thickness 0.25 mm). The parameters of the instrument were as follows: the initial column temperature was 120 °C (maintained for 3 min), and the temperature was programmed to 220 °C at a rate of 5 °C min⁻¹. Auxiliary heater temperature was 240 °C; the carrier gas was HE, the flow rate was 1 mL min⁻¹, the injection volume was 1 μ L, the split ratio was 10 : 1, and the scanning range was 50–550 amu. D-Glucose, D-galactose, D-mannose, D-xylose, rhamnose, L-fucose and L-arabinose were used as the monosaccharide standards and myoinositol was used as the interior standard.²¹

2.8 Antioxidant activity analysis *in vitro*

2.8.1 DPPH scavenging activity. The alcohol solution of DPPH radicals is purple and produces a colorless substance when in contact with reducing agents. Therefore, the scavenging rate of DPPH free radical can be used to evaluate the antioxidant activity of samples. The DPPH radical scavenging activity was conducted according to the reported method with modifications.²² Briefly, each polysaccharide sample was added with 0.2 mmol L⁻¹ freshly prepared DPPH-ethanol solution in

the ratio of 1 : 1, and then vibrated uniformly and reacted in the dark for 15 min. After the reaction, the absorbance at the wavelength of 517 nm was measured. The sample concentration ranges from 200 to 1000 μ g mL⁻¹. The DPPH radical scavenging activity was calculated as follows:

$$\text{Scavenging effect} = \frac{A_0 - (A_1 - A_2)}{A_0} \times 100\%,$$

where A_0 , A_1 , and A_2 are the absorbance of the water with DPPH solution, the sample was mixed with the DPPH solution, and the sample in anhydrous ethanol, respectively.

2.8.2 ABTS radical scavenging activity. A solution of ABTS radical is green, which can be reduced by the addition of a reductive substance. Therefore, ABTS free radical scavenging rate is also a commonly used index to evaluate the antioxidant activity of samples. The ABTS radical scavenging activity was also conducted according to the reported method with modifications.²² Briefly, 7 mmol L⁻¹ of ABTS solution was mixed with 2.5 mmol L⁻¹ of potassium persulfate and was stored in the dark at room temperature for 16 h to prepare ABTS radical. The ABTS free radical solution was then diluted with distilled water to an absorbance of about 0.7 at 734 nm. Sample solutions of different concentrations (200–1000 μ g mL⁻¹) were mixed in ABTS free radical solutions in a ratio of 1 : 1. The absorbance was measured at 734 nm after the reaction at room temperature for 15 min. ABTS radical scavenging activity was calculated by the following formula.

$$\text{Scavenging effect} = \frac{B_0 - (B_1 - B_2)}{B_0} \times 100\%,$$

where B_0 , B_1 , and B_2 are the absorbance of the water with ABTS solution, the sample mixed with the ABTS solution, and the sample in distilled water, respectively.

2.9 Optimization of the extraction process by response surface method

HWE method was considered to be the best extraction method of PTXP (analysis process was in results section). The extraction conditions were optimized with the extraction yield of PHWE-PTXP as an index. In this experiment, the standard curve built in Section 2.4 was still used. The calculation formula was as follows:

$$\text{Purity} = \frac{C \times V \times N \times 0.9}{m} \times 100\%,$$

where C is the concentration of PHWE-PTXP measured by the standard curve, V is the volume of the solution of PHWE-PTXP, N is the dilution factor of PHWE-PTXP solution, m is the mass weight of the extracted PHWE-PTXP, and 0.9 is the conversion factor.

2.9.1 Single factor experiments. After extraction by HWE, the content of PTXP was determined directly by standard curve. In the single factor experiment, the effects of solid–liquid ratio (1 : 10, 1 : 15, 1 : 20, 1 : 25, 1 : 30 g : mL), extraction temperature (50, 60, 70, 80, 90 °C), extraction time (1, 1.5, 2, 2.5, 3 h), number of extraction (1, 2, 3, 4, 5 times) on the yield of PHWE-PTXP were investigated to screen out the best factor level.



2.9.2 Box–Behnken design. On the basis of single factor experiments, the extraction conditions of HWE were further optimized by the response surface method. A BBD was performed with four independent variables at three levels.

2.9.3 Statistical analysis. Data of yield and purity determination from at least three experiment was represented as mean \pm standard deviation. The statistical difference among the compared groups was calculated by the Two-way analysis of variance using SPSS 17.0 software (International Business Machines Corporation, New York, USA), and P value < 0.05 was considered statistically significant.

3 Results and discussion

3.1 Yield and purity determination of PTXPs

The yield and purity of PTXPs were shown in Table 1. The extraction yield (W_1) order of crude PTXPs was as follows: HWE $>$ DAE $>$ UAE $>$ CAE. After purification by anion exchange chromatography, the extraction yield (W_2) order of purified PTXPs was changed to HWE $>$ CAE $>$ UAE $>$ DAE. By comparing W_1 and W_2 , we found that the extraction yield of crude PTXPs decreased significantly after the purification by anion exchange chromatography. Therefore, the extraction yield order of purified PTXPs also changed, which was caused by the difference of the purity of crude PTXPs (W_3). After purification, the purity (W_4) of purified PTXPs were all above 95%, which laid a foundation for the subsequent experiments. For both crude PTXPs and purified PTXPs, the extraction yield of the HWE method was the highest.

The activities and extraction yield of crude polysaccharides were often used to screen different extraction methods.^{23–25} Our results indicated that only the extraction yield or activities of crude polysaccharides were taken as evaluation indicators might be inaccurate in screening different extraction methods. The purity of crude polysaccharides might vary greatly, and crude polysaccharides still contained many small non-polysaccharides molecules, which might interfere with the activities of polysaccharides. Therefore, in our experiment, we used anion exchange chromatography to purify the crude PTXPs. After purification, the PTXPs had high purity, so that the yield could accurately reflect the efficiency of different extraction methods, and the activities of PTXPs could be accurately measured without interference from small molecules.

Table 1 Yield and purity of PTXPs (%; $\bar{x} \pm s$, $n = 3$)^a

	W_1	W_2	W_3	W_4
DAE	29.4 \pm 2.9 ^{####}	15.3 \pm 2.5 [#]	52.1 \pm 5.1 ^{ΔΔΔ}	95.6 \pm 0.9
UAE	28.4 \pm 1.9 ^{*#}	18.1 \pm 4.0	63.8 \pm 14.3 ^{ΔΔΔ}	96.5 \pm 3.0
CAE	25.2 \pm 2.3 ^{**}	20.3 \pm 1.9	80.4 \pm 4.5 ^{∇∇}	96.7 \pm 0.9
HWE	38.7 \pm 2.0 ^{####}	24.5 \pm 4.7	63.3 \pm 9.0 ^{ΔΔΔ}	97.1 \pm 2.5

^a W_1 compared with W_2 : * $P < 0.05$, ** $P < 0.01$; DAE, UAE, CAE compared with HWE: # $P < 0.05$, ## $P < 0.01$, ### $P < 0.001$; W_3 compared with W_4 : Δ $P < 0.05$, ΔΔ $P < 0.01$, ΔΔΔ $P < 0.001$; DAE, UAE, HWE compared with CAE: ∇ $P < 0.05$, ∇∇ $P < 0.01$, ∇∇∇ $P < 0.001$.

3.2 Molecular weight of purified PTXPs

Since crude PTXPs were purified only by anion exchange chromatography, it was difficult to determine the molecular weight accurately. If the molecular weight of polysaccharides was to be determined accurately, they were usually purified by anion exchange chromatography and further separated by gel column chromatography.^{17,26} But with the help of HPGPC, we could know whether the molecular weight of PTXPs extracted by different extraction methods was different. HPGPC, a type of size exclusion chromatography that separates analytes on the basis of molecular size, is suitable for the analysis of the molecular weight of PTXPs.²⁷ The molecular weight of PTXPs could be compared through the distribution range of retention time and chromatographic peak shape. As shown in Fig. 1A–C, there was almost no difference in the retention time of DAE-PTXP, UAE-PTXP and HWE-PTXP. The chromatographic peak shape of CAE-PTXP was different from the other three (Fig. 1D). The result showed that the CAE method affected the molecular weight of CAE-PTXP in the process of extraction and purification under the conditions of this experiment.

3.3 FTIR-ATR analysis of purified PTXPs

Infrared spectrum scanning can identify the characteristic functional groups in polysaccharides. Taking Fig. 2A as an example, 3312 cm^{-1} corresponded to the OH stretching vibrations of hydroxyl groups, 2922 cm^{-1} corresponded to the C–H stretching vibrations of CH_2 groups, and 1151, 1077 and 1012 cm^{-1} indicated that DAE-PTXP was a pyranose. As shown in Fig. 2, the infrared spectrum of DAE-PTXP, UAE-PTXP and HWE-PTXP had no difference. And the infrared spectrum of CAE-PTXP has an additional absorption peak at 1105 cm^{-1} (Fig. 2D), which might correspond to C–O or C–N stretching vibrations. Thus, it could be seen that CAE did have an impact on the structure of CAE-PTXP.

3.4 Monosaccharide composition of purified PTXPs

The polysaccharides are complex carbohydrates. Different polysaccharides have different monosaccharide compositions.⁷ The analysis of monosaccharide composition is an important method to characterize the structure of polysaccharides. But as shown in Fig. 3, DAE-PTXP, UAE-PTXP, HWE-PTXP and CAE-PTXP were all composed of glucose compared to the GC chromatogram of the standards. According to the results of molecular weight determination and infrared spectrum scanning, the effect of CAE on the structure of PTXP was mainly on the molecular weight and some functional groups. In our experiment, the CAE method affected the structure of PTXP, but this does not mean that the CAE method cannot be used to extract PTXP. If the CAE method is to be used, the extraction conditions should be further studied and strictly controlled.

3.5 In Vitro antioxidant activity of PTXPs and purified PTXPs

DPPH and ABTS radicals were widely used to evaluate the antioxidant ability of polysaccharides.^{22,28,29} As shown in Fig. 4A, the scavenging activity of crude polysaccharides DAE-PTXP,



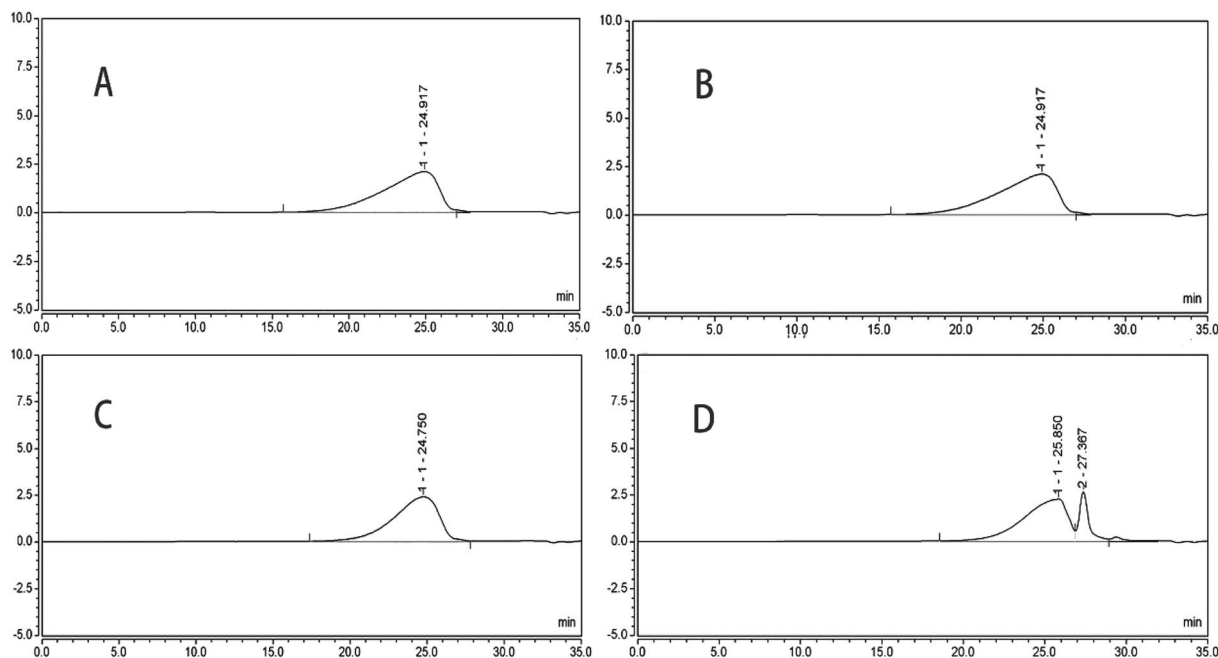


Fig. 1 Molecular weight of purified PTXPs. (A) Dilute alkaline extraction-polysaccharide of PaoTianXiong (DAE-PTXP); (B) ultrasonic-assisted extraction-polysaccharide of PaoTianXiong (UAE-PTXP); (C) hot water extraction-polysaccharide of PaoTianXiong (HWE-PTXP); (D) cellulase-assist extraction-polysaccharide of PaoTianXiong (CAE-PTXP).

CAE-PTXP and HWE-PTXP on DPPH increased gradually when their concentrations increased, and the scavenging rate of UAE-PTXP for DPPH did not change. In general, the scavenging activity order of PTXPs for DPPH free radical was as follows:

DAE-PTXP > CAE-PTXP > UAE-PTXP > HWE-PTXP. After purification, the DPPH radical scavenging activity of purified PTXPs was not in a dose-dependent manner (Fig. 4B). At this time, there was no significant difference in DPPH scavenging activity

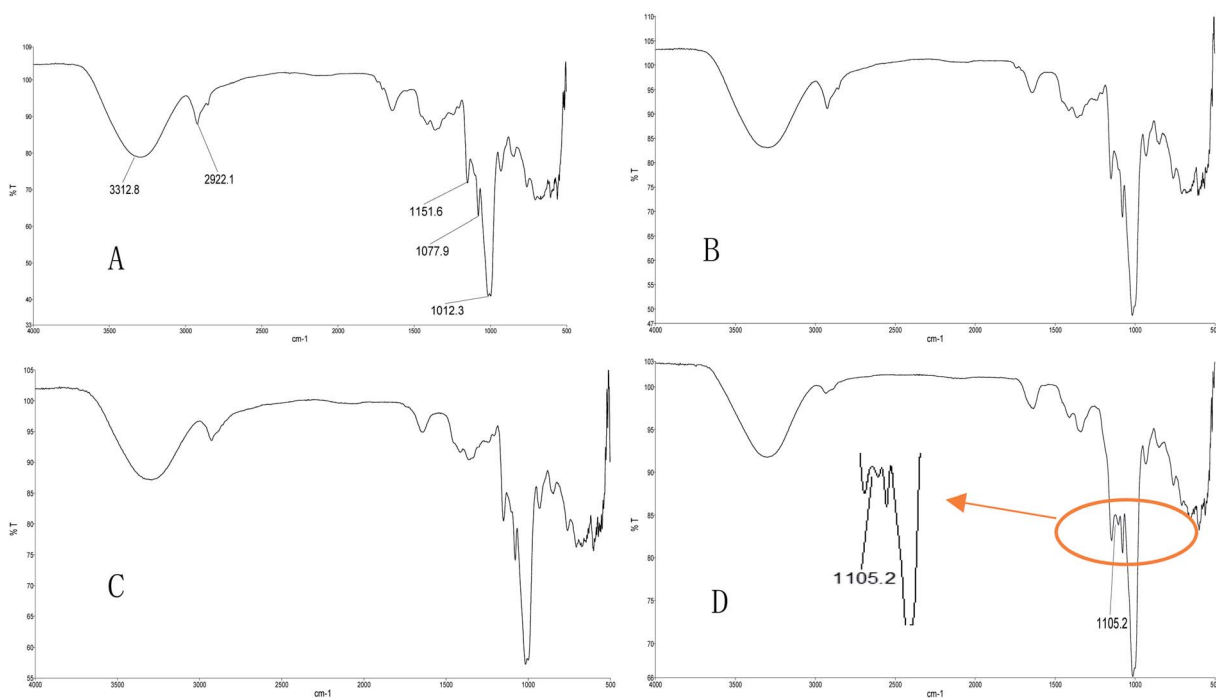


Fig. 2 The infrared spectrums of purified PTXPs. (A) Dilute alkaline extraction-polysaccharide of PaoTianXiong (DAE-PTXP); (B) ultrasonic-assisted extraction-polysaccharide of PaoTianXiong (UAE-PTXP); (C) hot water extraction-polysaccharide of PaoTianXiong (HWE-PTXP); (D) cellulase-assist extraction-polysaccharide of PaoTianXiong (CAE-PTXP).



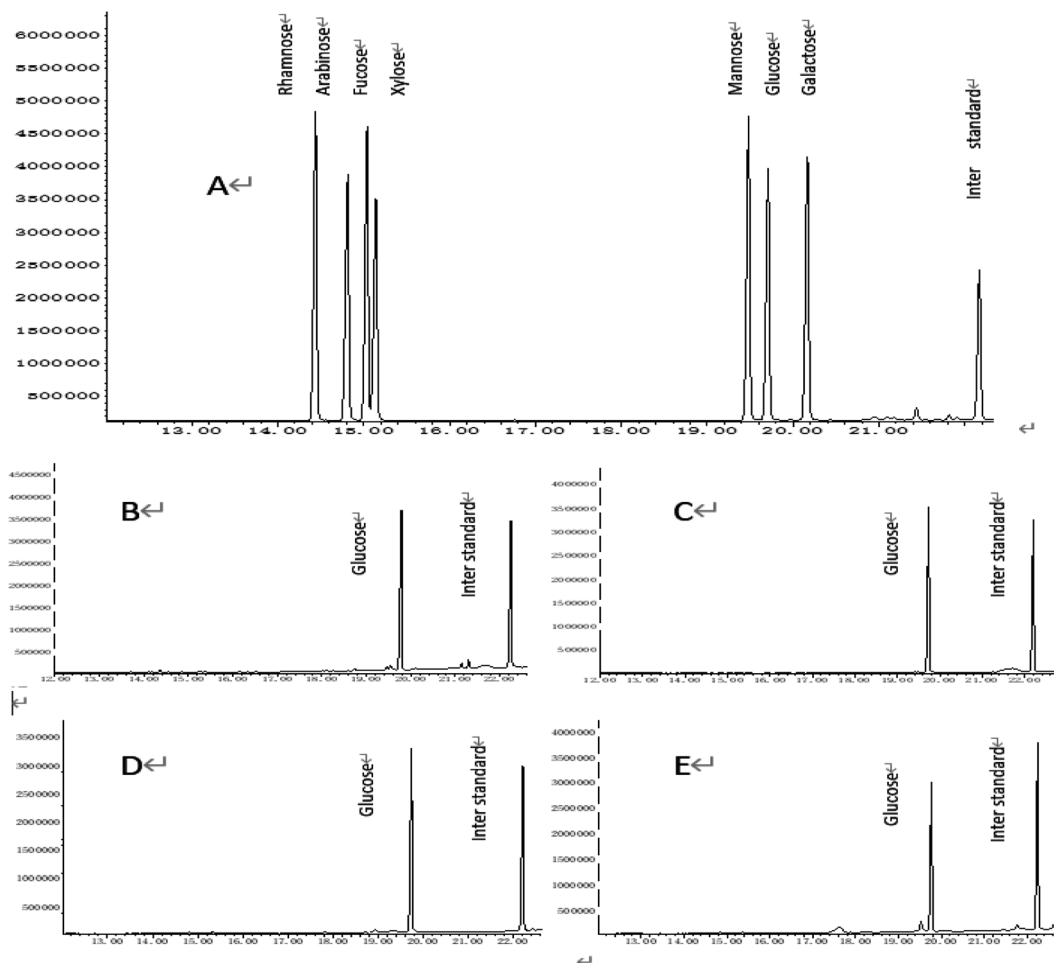


Fig. 3 Mass spectrogram of monosaccharide composition. (A) mixed standers; (B) purified dilute alkaline extraction-polysaccharide of PTX (PDAE-PTXP); (C) purified ultrasonic-assisted extraction-polysaccharide of PTX (PUAE-PTXP); (D) purified hot water extraction-polysaccharide of PTX (PHWE-PTXP); (E) purified cellulase-assist extraction-polysaccharide of PTX (PCAE-PTXP).

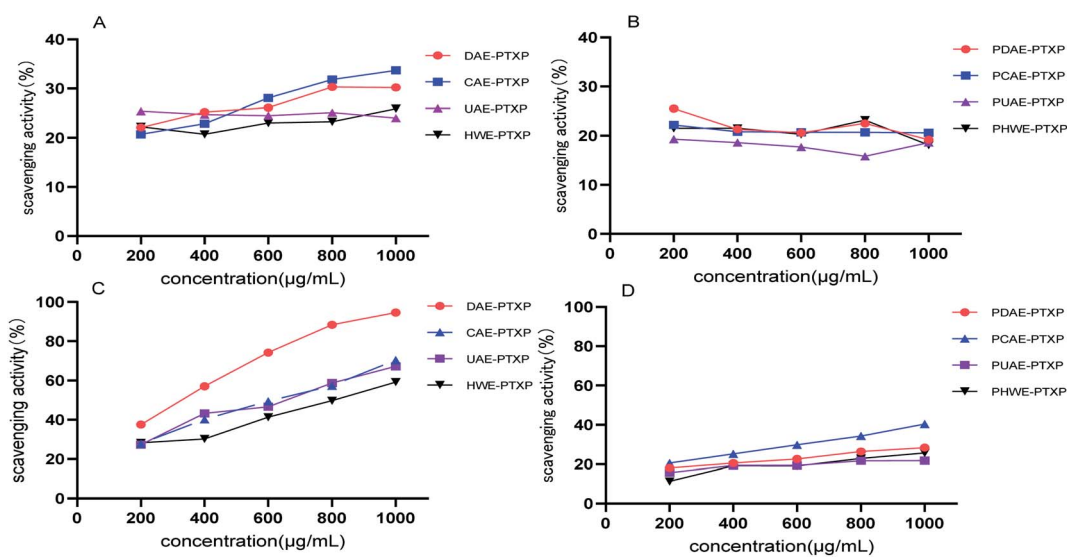


Fig. 4 DPPH and ABTS radical scavenging activity of crude PTXPs and purified PTXPs. (A) and (B) are about DPPH; (C) and (D) are about ABTS. PDAE-PTXP: purified dilute alkaline extraction-polysaccharide of PTX; PCAE-PTXP: purified cellulase-assist extraction-polysaccharide of PTX; PUAE-PTXP: purified ultrasonic-assisted extraction-polysaccharide of PTX; PHWE-PTXP: purified hot water extraction-polysaccharide of PTX.



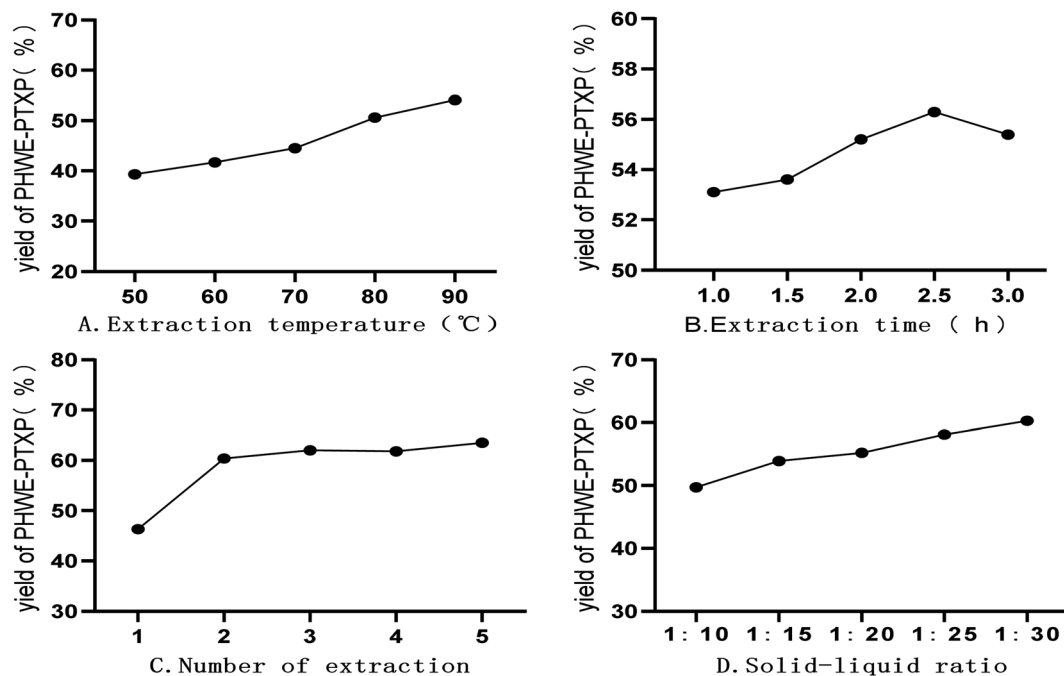


Fig. 5 Effects of different independent factors on extraction yield of purified hot water extraction-polysaccharide of PTX (PHWE-PTXP).

between PDAE-PTXP, PCAE-PTXP and PHWE-PTXP, and the activity was slightly better than that of PCAE-PTXP.

As shown in Fig. 4A, the crude PTXPs had good ABTS free radical scavenging activity in a dose-dependent manner, especially polysaccharides DAE-PTXP. The order of scavenging activity was as follows: DAE-PTXP > CAE-PTXP = UAE-PTXP > HWE-PTXP. But after purification, the scavenging activity of purified PTXPs on ABTS free radical was significantly reduced (Fig. 4D). In general, there was no significant difference in ABTS free radical scavenging activity between PDAE-PTXP, PUAE-PTXP and PHWE-PTXP, and their activity was slightly lower than that of PCAE-PTXP.

The results showed that the purity of PTXP had an effect on their antioxidant activities *in vitro*. If the scavenging activity of crude PTXPs on ABTS free radicals was used to screen the best extraction method among DAE, CAE, UAE and HWE, the DAE method had an obvious advantage. However, the purified PTXPs showed no significant difference in scavenging activity against ABTS and DPPH free radicals. This indicated that the purity of PTXP has a great influence on its antioxidant activity or other activities, and the use of anion-exchange chromatography for purification can be a good solution to this problem. Based on the

extraction yield, monosaccharide composition, molecular weight distribution and infrared spectroscopy results, we considered that HWE was the most suitable method for the extraction of PTXP. The structure of PHWE-PTXP was not significantly affected by the HWE method, and its activity was no worse than that of PTXP extracted by other methods. Therefore, the HWE method was optimized in the following experiment.

In the previous experiment, if only considering the extraction yield without detecting the structure of PTXP, we would not find the CAE method under the extraction conditions in this paper could influence the structure of PTXP. And if crude PTXPs were not purified, the DAE method would be mistaken as the best extraction method because of the best ABT-scavenging activity of DAE-PTXP. Our results suggested that the quality of polysaccharides should be controlled also by the structure and activity, rather than just considering the extraction yield before the study of PTXP in depth. Similarly, it is important to study and optimize extraction methods even for other polysaccharides.

3.6 Optimization of the HWE method by response surface method

3.6.1 Single factor experimental evaluation. Through the classical “one factor at a time” methodology, all factors remain unchanged during the single factor experiment except those under study.¹¹ The effects of solid-liquid ratio, extraction temperature, extraction time and the number of extraction on the extraction yield of PHWE-PTXP were shown in Fig. 5. The extraction yield of PHWE-PTXP increased when the temperature increase and the highest extraction yield was obtained at 90 °C (Fig. 5A). If the extraction temperature continues to increase to 100 °C, the yield of PHWE-PTXP may be higher, the energy

Table 2 Coding of Box–Behnken test design factors and levels

Independent variables	Factor levels		
	−1	0	1
Extraction temperature (°C)	70	80	90
Extraction time (h)	2	2.5	3
Solid-liquid ratio (mL : g)	1 : 20	1 : 25	1 : 30
Number of extraction	1	2	3



Table 3 Response surface experiment design and results

Run	Extraction temperature (°C)	Extraction time (h)	Solid-liquid ratio (g : mL)	Number of extraction	Extraction yield (%)
1	-1	0	0	-1	33.6
2	0	1	0	1	60.1
3	0	0	0	0	56.9
4	0	-1	0	1	60.4
5	0	-1	1	0	55.3
6	0	0	1	1	60.2
7	0	1	1	0	57.5
8	0	-1	-1	0	51.3
9	-1	0	0	1	56.1
10	-1	0	-1	0	53.8
11	0	1	-1	0	51.8
12	0	0	0	0	55.6
13	0	0	0	0	57.9
14	0	-1	0	-1	33.3
15	-1	1	0	0	52.4
16	1	0	0	1	62.3
17	-1	0	1	0	57.3
18	1	-1	0	0	60.4
19	-1	-1	0	0	51.9
20	1	0	-1	0	56.7
21	0	0	-1	1	58.4
22	1	1	0	0	54.7
23	1	0	0	-1	45.5
24	0	1	0	-1	39.0
25	0	0	1	-1	35.9
26	1	0	1	0	60.3
27	0	0	-1	-1	34.4

consumption will also be greatly increased. Given the high yield at 90 °C, we considered that 90 °C was a very suitable extraction temperature for PHWE-PTXP.

As the extraction time changed from 1 h to 3 h, the extraction yield of PHWE-PTXP increased firstly and then decreased, and

the extraction yield was the highest at 2.5 h (Fig. 5B). When the extraction time exceeded 2.5 h, the yield of PHWE-PTXP decreased, which might be related to the degradation of PHWE-PTXP. As shown in Fig. 5C, when PHWE-PTXP was extracted only once, the yield was very low. When the extraction

Table 4 F-test and ANOVA analysis of the response surface quadratic model

Source	Sum of squares	df	Mean square	F value	P-value prob > F	
Model	2081.269	14	148.662	27.340	<0.0001	Significant
X_1	100.920	1	100.920	18.560	0.0010	
X_2	0.701	1	0.701	0.129	0.7258	
X_3	33.668	1	33.668	6.192	0.0285	
X_4	1536.803	1	1536.803	282.627	<0.0001	
X_1X_2	9.610	1	9.610	1.767	0.2084	
X_1X_3	0.003	1	0.003	0.000	0.9832	
X_1X_4	8.122	1	8.122	1.494	0.2451	
X_2X_3	0.722	1	0.722	0.133	0.7218	
X_2X_4	9.000	1	9.000	1.655	0.2225	
X_3X_4	0.023	1	0.023	0.004	0.9498	
X_1^2	1.080	1	1.080	0.199	0.6638	
X_2^2	14.741	1	14.741	2.711	0.1256	
X_3^2	6.021	1	6.021	1.107	0.3134	
X_4^2	322.403	1	322.403	59.292	<0.0001	
Residual	65.251	12	5.438			
Lack of fit	62.591	10	6.259	4.706	0.1879	Not significant
Pure error	2.660	2	1.330			
Cor total	2146.520	26				
R^2	0.9696					



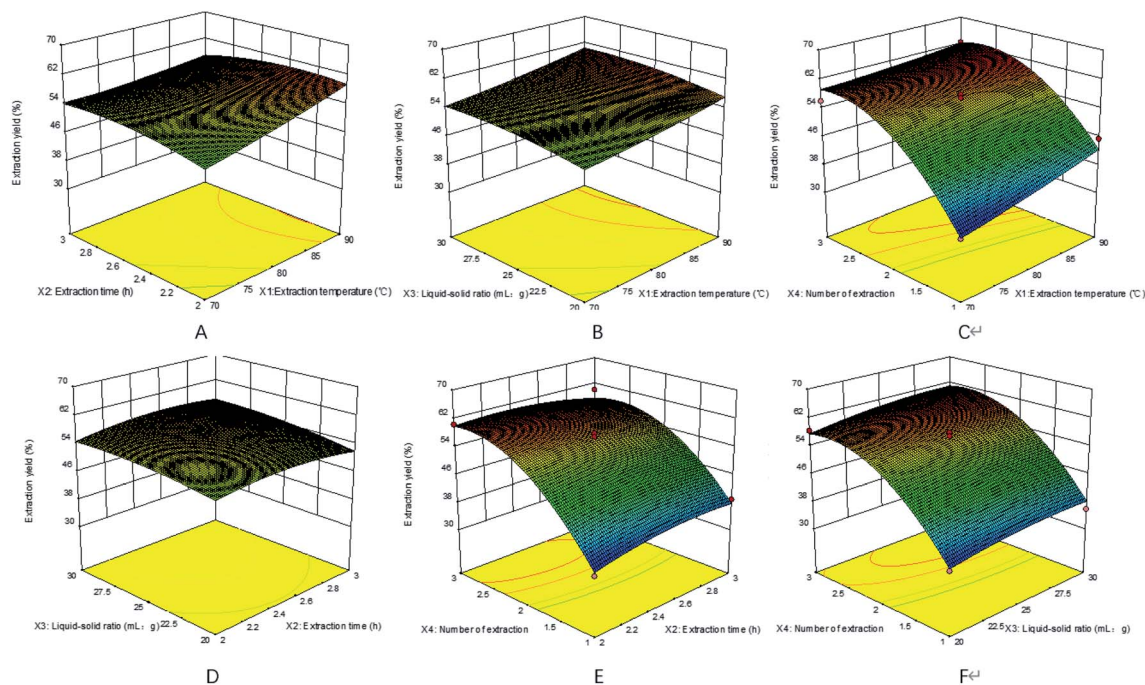


Fig. 6 Response surface (3D) showing the interactive effects of independent variables.

times reached more than 2 times, the yield remained at about 60%. At the same time, we could find that when the solid–liquid ratio changed from 1 : 10 to 1 : 30, the yield of HWE-PTXP kept increasing (Fig. 5D).

3.6.2 Optimization of extracting conditions by the Box–Behnken design. According to the single-factor experimental evaluation above, we chose four factors and three levels to optimize the extraction conditions of the HWE method. The range of the independent factors and levels were presented in Table 2. The response surface experiment design and results were presented in Table 3. By applying multiple regression analysis on the experimental data, the response variable and the test variables were related by the following equation:

$$Y(\%) = 56.8 + 2.9X_1 + 0.24X_2 + 1.68X_3 + 11.32X_4 - 1.55X_1X_2 + 0.025X_1X_3 - 1.43X_1X_4 + 0.43X_2X_3 - 1.5X_2X_4 + 0.075X_3X_4 + 0.45X_1^2 - 1.66X_2^2 - 1.06X_3^2 - 7.78X_4^2$$

where Y is the extraction yield of HWE-PTXP, X_1 is the extraction temperature, X_2 is the extraction time, and X_3 is solid–liquid ratio, and X_4 is number of extraction.

We used F -test and ANOVA analysis to estimate the significance and applicability of the regression model (Table 4). The regression model of PHWE-PTXP yield was extremely significant ($P < 0.0001$), the correlation coefficient (R^2) was 0.9696 and the lack of fit had no significant effect ($P > 0.05$). The results showed that the model represented the data satisfactorily. It could be seen that X_1 , X_3 , X_4 and X_4^2 had a significant effect on the extraction yield of HWE-PTXP.

3.6.3 Response surface analysis. To ensure the reproducibility of pharmacological experiments, some scholars established quality control methods for related drugs.^{30,31} For

polysaccharides, reliable extraction conditions are an important means of quality control. The three-dimensional apparent diagram of the interaction effects of extraction time, extraction temperature, number of extraction and solid–liquid ratio on the extraction yield of HWE-PTXP was drawn by Desk-Expert 10.0.7 software (Fig. 6). For the extraction yield of HWE-PTXP, it could be seen that the interaction between X_1 and X_2 , X_1 and X_4 , and X_2 and X_4 were not significant (Fig. 6A, B and D). The interactions of X_1 and X_4 , X_2 and X_4 , and X_3 and X_4 have significant effects on the yield of HWE-PTXP. Using the generated model to predict the optimal conditions, the optimal extraction conditions of the HWE method were as follows: the extraction temperature was 90 °C, the extraction time was 2.17 h, the solid–liquid ratio was 1 : 28.5, the number of extraction was 2.69, and the maximum predicted extraction yield was 64.61%. Considering the feasibility and convenience in the actual operation, the actual conditions were changed slightly: the extraction temperature was 90 °C, the extraction time was 2.17 h, the solid–liquid ratio was 1 : 29, the number of extraction was 2. Under these conditions, the yield of HWE-PTXP was 63.1%, which was only 1.5% different from 64.61%. The analysis result showed that the experimental values were in good agreement with the predicted values, and also suggested the established model was satisfactory and accurate.

4 Conclusion

In the present study, the effect of DAE, UAE, CAE and HWE on yield, characteristics, and antioxidant activities of PTPs were investigated to determine the optimal extraction method. The results showed that CAE-PTPX, HWE-PTXP, UAE-PTXP and DAE-PTXP had the same monosaccharide composition. They all



have the typical characteristics of polysaccharides based on infrared analysis but CAE-PTXP was slightly different from the other three. The molecular weight of CAE-PTXP was also different from HWE-PTXP, UAE-PTXP and DAE-PTXP. In addition, the purity and antioxidant activity of crude PTPPs were significantly different from that of purified PTPPs. These results indicated that the properties of PTPPs extracted by different extraction methods were different. In the process of extracting PTPP, not only the extraction yield but also the effects of different extraction methods on the properties should be considered. When the activity and structure of polysaccharides were determined, the purity of PTPPs should not be ignored, which supported our previous points. The HWE had the highest extraction yield and had little effect on the properties of PTPP, so it was considered to be the best extraction method of PTPP. Next, with the help of the optimized HWE, we will conduct an in-depth study on the biological activity of PTPP.

Author contributions

Kuncheng Qiu: conceptualization, methodology, investigation, writing-original draft. Zunjiang Li: methodology, investigation, formal analysis. Yingxin Long: investigation, validation, visualization. Zhongyu Lu: software, validation. Wei Zhu: conceptualization, funding acquisition, supervision, writing-review & editing, project administration.

Conflicts of interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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References

- 1 X. Wu, S. Wang, J. Lu, Y. Jing, M. Li, J. Cao, B. Bian and C. Hu, *Chin. Med.*, 2018, **13**, 4.
- 2 E. Nyirimigabo, Y. Xu, Y. Li, Y. Wang, K. Agyemang and Y. Zhang, *J. Pharm. Pharmacol.*, 2015, **67**, 1–19.
- 3 S. G. Sousa, L. A. Oliveira, D. de Aguiar Magalhaes, T. V. de Brito, J. A. Batista, C. M. C. Pereira, M. de Souza Costa, J. C. R. Mazulo, M. de Carvalho Filgueiras, D. F. P. Vasconcelos, D. A. da Silva, F. C. N. Barros, V. G. Sombra, A. L. P. Freitas, R. C. M. de Paula, J. P. de Andrade Feitosa and A. L. Dos Reis Barbosa, *Carbohydr. Polym.*, 2018, **197**, 515–523.
- 4 G. Huang, X. Mei and J. Hu, *Curr. Drug Targets*, 2017, **18**, 1296–1300.
- 5 Y. Fu, L. Shi and K. Ding, *Int. J. Biol. Macromol.*, 2019, **141**, 693–699.
- 6 C. Y. Shen, J. J. Jiang, M. Q. Li, C. Y. Zheng and W. Zhu, *J. Funct. Foods*, 2017, **35**, 352–362.
- 7 C. Y. Shen, W. L. Zhang and J. G. Jiang, *J. Funct. Foods*, 2017, **34**, 118–129.
- 8 H. Feng and L. Tian, *Molecules*, 2021, **26**, 2359.
- 9 Z. G. Qian, *Carbohydr. Polym.*, 2014, **101**, 432–434.
- 10 J. Gu, Q. Li, J. Liu, Z. Ye, T. Feng, G. Wang, W. Wang and Y. Zhang, *Int. J. Biol. Macromol.*, 2021, **167**, 423–433.
- 11 Z. He, Y. Zhu, X. Bao, L. Zhang, N. Li, G. Jiang and Q. Peng, *Molecules*, 2019, **24**, 2221.
- 12 F. Chen and G. Huang, *Int. J. Biol. Macromol.*, 2018, **112**, 211–216.
- 13 Y. Han, Y. Wu, G. Li, M. Li, R. Yan, Z. Xu, H. Lei, Y. Sun, X. Duan, L. Hu and R. Huang, *Food Chem.*, 2021, **342**, 128537.
- 14 H. Wang, J. Chen, P. Ren, Y. Zhang and S. Omondi Onyango, *Ultrason. Sonochem.*, 2021, **70**, 105355.
- 15 T. Wang, H. Zhao, Y. Bi and X. Fan, *J. Food Sci.*, 2021, **86**, 977–986.
- 16 Y. B. Ji and F. L. Wang, *Bioengineered*, 2021, **12**, 382–391.
- 17 X. Yang, Y. Wu, C. Zhang, S. Fu, J. Zhang and C. Fu, *Int. J. Biol. Macromol.*, 2020, **143**, 314–324.
- 18 W. L. Chen, L. H. Xiao, W. Zhu and J. Qin, *J. Northwestern Pharm.*, 2018, **33**, 584–588.
- 19 C. Zhao, M. Li, Y. Luo and W. Wu, *Carbohydr. Res.*, 2006, **341**, 485–491.
- 20 R. Stancanelli, R. Ficarra, C. Cannava, M. Guardo, M. L. Calabro, P. Ficarra, R. Ottana, R. Maccari, V. Crupi, D. Majolino and V. Venuti, *J. Pharm. Biomed. Anal.*, 2008, **47**, 704–709.
- 21 Y. Chen, M. Y. Xie, Y. X. Wang, S. P. Nie and C. Li, *Phytochem. Anal.*, 2009, **20**, 503–510.
- 22 P. Rozi, A. Abuduwaili, P. Mutailifu, Y. Gao, R. Rakhmanberdieva, H. A. Aisa and A. Yili, *Int. J. Biol. Macromol.*, 2019, **131**, 97–106.
- 23 G. Chen, C. Fang, C. Ran, Y. Tan, Q. Yu and J. Kan, *Int. J. Biol. Macromol.*, 2019, **130**, 903–914.
- 24 H. Dong, S. Lin, Q. Zhang, H. Chen, W. Lan, H. Li, J. He and W. Qin, *Int. J. Biol. Macromol.*, 2016, **93**, 179–185.
- 25 S. Jia, F. Li, Y. Liu, H. Ren, G. Gong, Y. Wang and S. Wu, *Int. J. Biol. Macromol.*, 2013, **62**, 66–69.
- 26 J. Song, Y. Wu, X. Ma, L. Feng, Z. Wang, G. Jiang and H. Tong, *Carbohydr. Polym.*, 2020, **230**, 115586.
- 27 J. Xu, S. L. Li, R. Q. Yue, C. H. Ko, J. M. Hu, J. Liu, H. M. Ho, T. Yi, Z. Z. Zhao, J. Zhou, P. C. Leung, H. B. Chen and Q. B. Han, *Anal. Bioanal. Chem.*, 2014, **406**, 6409–6417.
- 28 S. N. Li, S. H. Tang, Q. He, J. X. Hu and J. Zheng, *J. Dairy Sci.*, 2020, **103**, 1120–1130.
- 29 T. L. Gunathilaka, K. W. Samarakoon, P. Ranasinghe and L. D. C. Peiris, *Molecules*, 2019, **24**, 3708.
- 30 K. Qiu, Z. Li, C. Li, H. Huang and W. Zhu, *J. Funct. Foods*, 2020, **71**, 104036.
- 31 Z. Y. Zhang, D. D. Gui, M. Sha, J. Liu and H. Y. Wang, *J. Dairy Sci.*, 2019, **102**, 68–76.

