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1. Introduction

Celastrus orbiculatus Thunb. (COT), belonging to the family Celastraceae and the genus Celastrus L., is widely distributed throughout China.¹ The parts including root, stem and leaf all could been used as Chinese folk medicine to treat rheumatoid arthritis, vomiting, abdominal pain, and snakebites.^{2,3} The root of COT was reported to possess anti-tumor,⁴ antiviral,⁵ bacteriostatic⁶ and lipid-lowering¹ activities, and about 50 compounds, including triterpenes, sesquiterpenoids, steroids and organic acids, were isolated from the root or root bark.^{3,7-9} The stem was also reported to have anti-inflammation, 10 anticancer¹¹ and fatty liver amelioration¹² effects, and nearly 100 compounds including triterpenes, diterpenoids, steroids, flavonoids, phenolics and benzoquinone were isolated.¹³⁻¹⁶ For the leaf part, previous studies have found that the extract of it had insecticidal effect and hypoglycemic effect,¹⁷ while only a few flavonoids were isolated.¹⁸ It was revealed that there were significant variation for the contents of celastrol or total

Comprehensive metabolomics analysis based on UPLC-Q/TOF-MS^E and the anti-COPD effect of different parts of Celastrus orbiculatus Thunb.

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The root, stem and leaf of Celastrus orbiculatus Thunb. (COT) have all been used as Chinese folk medicine. Aiming at revealing the secondary metabolites and screening the anti-COPD effect of COT, the comprehensive phytochemical and bioassay studies were performed. Based on the ultra-high performance liquid chromatography combined with quadrupole time-of-flight mass spectrometry (UPLC-Q/TOF-MSE), the screening analysis of components in COT was conducted with the UNIFI platform, the metabolomics of the three parts were analyzed with multivariate statistical analysis. Cigarette smoke extract (CSE)-stimulated inflammatory model in A549 cells was used to investigate the biological effect of the three parts. A total of 120 compounds were identified or tentatively characterized from COT. Metabolomics analysis showed that the three parts of COT were differentiated, and there were 13, 8 and 5 potential chemical markers discovered from root, stem and leaf, respectively. Five robust chemical markers with high responses could be used for further quality control in different parts of COT. The root, stem and leaf of COT could evidently reduce the levels of pro-inflammatory factors in a dose-dependent way within a certain concentration range. The stem part had a stronger anti-COPD effect than root and leaf parts. This study clarified the structural diversity of secondary metabolites and the various patterns in different parts of COT, and provided a theoretical basis for further utilization and development of COT. **PAPER**
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alkaloids in different parts of COT.^{19,20} Along with continuous expansion of the folk and clinical application of COT, an indepth study on the chemical constituents in different parts of COT has attracted more and more attention. However, the comprehensive comparative study on the chemical composition between root, stem and leaf parts of COT has not been reported so far.

Recently, the UPLC-Q/TOF-MS method has been innovatively used for screening and identifying chemical components in herbal medicines and traditional Chinese medicine. And the global profiling of various metabolites were reported. As part of these research works, we reported this method to detect some natural products including Platycodon grandiflorum and Ginseng root.^{21,22} Our research results showed that this method is high throughput, comprehensive, simple and efficient. As far as we know, the UPLC-Q/TOF-MS method has not been reported to identify the components in COT. So, the study in this paper comparatively analyzes the phytochemicals of root, stem and leaf parts of COT by using the UPLC-Q/TOF-MS method for the first time and finds out the similarities and differences between them.

Chronic Obstructive Pulmonary Disease (COPD), predicted to rank as the third leading cause of death in the world, 23 is mainly caused by significant exposure to harmful gases or

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particles.²⁴ Cigarette smoking was the leading environmental risk factor for COPD around the world, and cigarette smokers were more likely to develop respiratory symptoms and had a higher COPD mortality rate. Along with the progressive lung inflammation, some pro-inflammatory mediators such as IL-1 β , IL-6 and TNF-a participated in the occurrence and development of COPD.²⁵ Although the COT had been used in treating various inflammatory diseases, the effect on the cigarette smoke extract (CSE)-induced inflammatory reaction has not been reported so far.

In the present study, the main medicinal parts of COT (root, stem and leaf) were chosen as the test sample. On one hand, the similarities and differences of phytochemicals in three parts were analyzed by using UNIFI platform and untargeted metabolomics based on UPLC-Q/TOF-MSE. The components and potential chemical markers to profile diverse classifications of metabolites of three parts were investigated. On the other hand, the effects on CSE-induced inflammatory reaction of these three parts were explored in A549 cells. The anti-COPD activity of different parts was preliminarily discussed. This comprehensive study could reveal the structural diversity of secondary metabolites and the different patterns of main medicinal parts of COT, and provide the data for further clinical application in anti-COPD. The study on the phytochemistry and the pharmacological activity of various parts were both significantly valuable to the research and development of COT. Puper

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2. Experiment

2.1. Materials and reagents

A total of 10 batches of fresh COT were collected from different growth areas in China (Table 1). All herbs were authenticated by the authors according to Hunan Province Local Standard for Traditional Chinese Medicine (2009 edition) for "Celastrus orbiculatus Thunb.". The corresponding specimens had been deposited in the Research Center of Natural Drug, Jilin University, China.

Methanol and acetonitrile were of LC/MS grade purchased from Fisher Chemical Company. Formic acid was bought from Sigma-Aldrich Company, St. Louis, MO, USA. Deionized water was purified by Millipore water purification system (Millipore, Billerica, MA, USA). All other chemicals were analytically pure. Cigarettes for bioassay analysis were Xiongshi cigarette (China

Tobacco Zhejiang Industrial Co., Ltd, Hangzhou, China), each cigarette contained 11 mg of tar, 0.7 mg of nicotine, and 13 mg of carbon monoxide. Human lung carcinoma A549 cells were obtained from the Department of Pathogen Biology, Basic Medical College, Jilin University. ELISA kits were bought from Nanjing Jiancheng Bio-engineering Institute.

2.2. Sample preparation of three parts of COT

Took the whole fresh COT, and separated the root, stem and leaf part respectively to get 30 test samples including root part (R1– R10) samples, stem part (S1–S10) samples and leaf part (L1–L10) samples. The aforementioned parts were air-dried, grinded and sieved with Chinese National Standard Sieve No. 3, R40/3 series to obtain the homogeneous powder respectively. Each powder was weighted (2.0 g) accurately and extracted thrice (3 hours per time) with 100 mL of 80% methanol at 80 \degree C, cooled, filtered, collected and combined the filtrate of each sample, concentrated and evaporated to dryness.

For metabonomics analysis, each residue (all approximately 2.0 mg) was dissolved in 1.0 mL of 80% methanol respectively, after being filtered with a syringe filter (0.22 μ m), 30 test solutions ($R_{M1}-R_{M10}$, $S_{M1}-S_{M10}$ and $L_{M1}-L_{M10}$) were obtained, which was injected into the UPLC system directly. Furthermore, to ensure the suitability consistency and the stability of MS analysis, a sample for quality control (QC) was prepared by pooling ²⁰ mL from every test solution, namely containing all of the constituents in this analysis.

For screening analysis, the test solutions of root part (R_S) , stem part (S_S) and leaf part (L_S) were prepared by pooling 100 μ L from $R_{M1}-R_{M10}$, $S_{M1}-S_{M10}$ and $L_{M1}-L_{M10}$ solutions, respectively.

For bioassay analysis, the test samples $(R_{bio}, S_{bio}$ and L_{bio}) of root part, stem part and leaf part were prepared by combining each residue of R_1-R_{10} , S_1-S_{10} and L_1-L_{10} , respectively. Then, Rbio, Sbio and Lbio were dissolved in water at the concentration of 3.2 mg mL⁻¹ to get the stock solutions stored in 4 °C.

2.3. Ultra-high performance liquid chromatography combined with quadrupole time-of-flight tandem mass spectrometry (UPLC-Q/TOF-MSE)

The separation and detection of components were performed on the UPLC system combined with Xevo G2-XS Q/TOF mass

Table 2 Compounds identified from the root, stem and leaf of COT by UPLC-QTOF-MSEa

L

Calculated

Calculated

Theoretical

Theoretical

Open Access Article. Published on 26 February 2020. Downloaded on 11/28/2024 2:46:25 PM. This article is licensed under a [Creative Commons Attribution 3.0 Unported Licence.](http://creativecommons.org/licenses/by/3.0/) **[View Article Online](https://doi.org/10.1039/c9ra09965d)**(ppm) MS $^{\rm E}$ fragmentation \rm{NSE} Methyl-p-coumarate R 41 Evodinnol R 42 R 43 Aschantin R, S 44 Corylifol B S 45 Demethoxycurcumin R 46 (+)-Lyoniresinol S 47 Neobavaisoflavone R ا
م -0.6 O R R -0.6 S s Aksilarin R s $(+)$ -Celaphanol A R, S s Leachianone G R R 9-Octadecenoic acid R, S s R, S Interiorin C R, S Celangulin IV S, L ∞ $(3R-cis)-3,4-Dihydro-3,4$ methoxyphenyl)methyl]- R-cis)-3,4-Dihydro-3,4- (+)-7,8-
Didehydroarctigenin Tetramethoxystilbene Didehydroarctigenin diol-7-methoxy-3-[(4- 2H-1-benzopyran $trans-3,3',5,5'$ 443.1696 [M + Na]⁺, 205.0843 [M + H - 2 \times CH₂OH $^{+}$, 152 [M + $\rm H_2O-CH_3$ – +, 441.1997 [M $\mathrm{C_{4}H_{7}}$ $\frac{2}{1}$ $-2 \times \text{CH}_2\text{OH}$ $\overline{O_2^+}$, +, 81.0569 $-3 \times H_2O^+$ C_7H_7O ⁺ $\mathrm{C_2H_3O_2}^+,$ $-$ 2 \times CH₃O] $^{\dagger},$ 307.0731 [M + H $\mathrm{C_8H_7O}^+$ $-2 \times \mathrm{H}_2\mathrm{O} \rceil^*$, 234.0763 $\rm [M + H]$ \vec{O}_4^+ $-2 \times H_2O$ \vec{H} $-\text{CH}_3\Big]$ ⁺, $\mathrm{C_4H_7}^+$ ${\rm H_2O}_1^+$ $-$ CH₃]⁺, L –
CH₃ $^{+}$, 150.0562 [M + H $\overline{O_5}$ \vec{H}_{11} $\tilde{\bar{\sigma}}$ $\overline{\mathrm{o}}_2$ $\overline{\delta}$ L \vec{C}^4 \tilde{H}^s $\overline{O_2}$ L L L L L $\overline{O_2}$ $\mathrm{C_9H_{11}}$ $-2 \times H_2O$ $\mathrm{G_8H_{17}}$ -0.5 311.1252 [M + Na]⁺, 274.1090 [M + H +L $\mathrm{C_7H_5O_2}$ -4.0 353.2284 [M + Na]⁺, 277.2129 [M + H O8 420.1799 420.1784 3.4 443.1696 [M + Na]+, 205.0843 [M + H -4.3 323.1240 $\mathrm{[M + Na]}^+$, 138.0556 $\mathrm{[M + H]}$ $-$ CH₃] $\mathrm{C_9H_7O_2}^\mathrm{+}$ -3.9 179.0687 [M + H]⁺, 164.0374 [M + H -0.1 347.0761 $\left[\text{M} + \text{H}\right]^\dagger$, 329.0629 $\left[\text{M} + \text{H}\right]$ -3.5 249.1105 $[M + H]^+, 206.0877$ $[M + H]$ -3.8 401.1559 [M + H]⁺, 383.0623 [M + H -4.5 341.1355 $[M + H]$ ⁺, 326.0641 $[M + H]$ -3.3 339.1209 [M + H]^{\dagger} , 218.0945 [M + H] -3.6 357.1320 [M + H]^{\dagger} , 245.0820 [M + H] O5 316.1317 316.1311 1.8 339.1209 [M + Na]+, 251.0624 [M + H $\tilde{\vec{C}}$ -3.8 443.1675 [M + H]⁺, 384.1466 [M + H O_4 322.1218 322.1205 4.0 323.1.291 [M + H]⁺, 268.1004 [M + H] L \vec{H}_{11} O_{14} 676.2738 676.2731 1.0 677.2811 [M + H]⁺, 585.2390 [M + H] $\mathrm{C_8H_{17}}$ H2O $\rm\tilde{G}^o$ L L, L $+$, 167.0664 $[M + H]$ $\mathrm{C_2H_3O_2}^\mathrm{+}$ -1.7 371.1483 $[M + H]^+, 220$ $[M + H]$ $-2 \times H_2$ O L, $-2 \times H_2O$ L L $\rm C_4H_7O_2$ – $^{+}$, 556.2650 $\rm [M + H]$ L, $\mathrm{C_{5}H_{9}}^{\mathrm{+}}$ $^{+}$, 96.0346 [M + H $+$, 174.0900 $M + H$ CH3O $\mathrm{C_3H_7}$ $_{\rm H_2O}$ + L $-4\times \mathrm{C_2H_3O_2}$ I L, L L $\rm CH_{3} - 2 \times CH_{3}O$ 238.0842 [M + H 365.2118 [M + H 176.0735 [M + H 381.1268 [M + H 254.0834 [M + H 150.1100 [M + H L 107.0465 [M + H 299.0562 [M + H $\overline{O_4^+}$ H2O 3.8 309.1266 [M \vec{H} $\mathrm{C_8H_7O_2}^\mathrm{+}$ $\mathrm{C_{8}H_{9}O_{3}}$ \vec{o}^{\dagger} L \tilde{G}^2 $\mathrm{C_2H_3O_2}$ $\mathrm{C_6H_5O_2}$ \times CH₃] $[$ M + H $\mathrm{C_9H_{11}}$ L + H H mass (Da) 310,1205 O_3 178.0623 178.0630 346.0689 O₈ 346.0688 346.0689 O_4 248.1040 248.1049 370.1416 O6 370.1410 370.1416 O_4 288.1360 288.1362 O_7 400.1407 400.1522 O₅ 338.1143 338.1154 O₆ 356.1247 356.1260 322.1205 330.2406 $O₅$ 330.2392 330.2406 300.1362 O₄ 300.1348 300.1362 O_4 310.1193 310.1205 178.0630 248.1049 288.1362 676.2731 400.1522 340,1311 O₅ 340.1295 340.1311 338,1154 356,1260 316.1311 420.1784 442.1628 O₈ 442.1611 442.1628 504.3451 O₆ 504.3448 504.3451 mass (Da) 310.1193 178.0623 346.0688 248.1040 370.1410 288.1360 676.2738 400.1407 340.1295 338.1143 356.1247 316.1317 120.1799 322.1218 330.2392 300.1348 504.3448 442.1611 $\mathrm{C_{34}H_{44}O_{14}}$ $\mathrm{C_{10}H_{10}O_3}$ $\mathrm{C_{17}H_{14}O_8}$ $C_{14}H_{16}O_4$ $\rm{C_{21}H_{22}O_6}$ $C_{17}H_{20}O_4$ $\mathrm{C}_{22}\mathrm{H}_{24}\mathrm{O}_7$ $\rm C_{20}H_{20}O_5$ $\mathrm{C_{20}H_{18}O_5}$ $\mathrm{C_{20}H_{20}O_6}$ $\mathrm{C}_{18}\mathrm{H}_{20}\mathrm{O}_{5}$ $\mathrm{C}_{22}\mathrm{H}_{28}\mathrm{O}_8$ $\rm{C_{24}H_{26}O_8}$ $\mathbb{C}_{20}\mathbb{H}_1$ 8
O $_4$ $C_{18}H_{34}O_5$ $C_{18}H_{20}O_4$ $\mathrm{C_{30}H_{48}O_6}$ $\mathrm{C_{19}H_{18}O_4}$ Formula No. $t_{\rm R}$ (min) Formula 18 5.87 C₁₉ 19 5.96 C10 20 6.75 C_{17} 21 7.22 C₁₄ 7.58 C_{21} 23 7.73 C_{17} 24 8.47 C₃₄ 25 8.57 C_{22} 26 8.83 C₂₀ 27 8.94 C₂₀ 28 8.95 C₂₀ 29 8.96 C₁₈ $30#$ 8.98 C₂₂ 31 9.81 C₂₄ 32 10.35 C_{20} 33 11.39 C_{18} 34 13.75 C₁₈ $35*$ 14.06 C₃₀ $t_{\rm R} \, (\rm min)$ 13.75 6.75 8.83 10.35 11.39 14.06 5.87 5.96 7.22 7.58 7.73 8.47 8.57 8.94 8.95 8.96 8.98 9.81 30# $\frac{1}{2}$ $35*$ 18 $\overline{1}$ 25 26 31 $\overline{20}$ $\overline{21}$ 22 23 24 27 28 29 32 33 34

Table 2 (Contd.) Table 2 (Contd.)

72.0554 [M + H

451.3303 [M

 $-\mathrm{CO}_2$

, 261.4102 [M

L H L \vec{C} $\mathrm{H_{24}O}^{-1}$

L H L

L \vec{G} \tilde{H}_1^9 L $-H$ ^{$-$}, 454.3157 [M

L H H_2O] $-$, 407.3726 [M

L H L H2O

 $-CH₃$].

3b-Hydroxy-11a,12a-epoxy-

3β-Hydroxy-11α,12α-epoxy- R
13β,28-ursolide

 $\mathrm{C_2H_3O_2}^\mathrm{+}$

67 18.74 C₃₀

18.74

67

 $\rm{C_{30}H_{46}O_4}$

O4 470.3399 470.3396 0.6 469.3326 [M

470.3396

470.3399

Table 2 (Contd.) Table 2 (Contd.)

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No. $t_{\rm R}$ (min) Formula

.
So

 $t_{\rm R}\,({\rm min})$

Formula

Calculated mass (Da)

Theoretical Theoretical
mass (Da)

Mass error

(ppm) MS $^{\rm E}$ fragmentation \rm{NSE}

84#

83

85#

 $86*$

 $87*$

 88^\ast

89

 $90*$

 91

 $\overline{5}$

 93

95

96

 94

Table 2 (Contd.) Table 2 (Contd.)

 \overline{C} H₃

 $\tilde{\tilde{\sigma}}$

b: compared with NIST Chemistry WebBook; s: compared with the reference compounds; R: the root of Celastrus Thunb.; S: the stem of Celastrus orbiculatus Thunb.; L: the leaf of characteristic component in leaf; a: compared with spectral data obtained from Wiley Subscription Services, Inc. (USA); ※ $a *$ Characteristic component in root; # characteristic component in stem; Celastrus orbiculatus Thunb. Celastrus orbiculatus Thunb

spectrometer (Waters Co., Milford, MA, USA) with an electrospray ionization (ESI) interface.

The ACQUITY UPLC BEH C18 column (100 mm \times 2.1 mm, 1.7 μm) was bought from Waters Corporation (Milford, MA, USA). The moving phrase was consisted of eluent A (0.1% methanoic acid in water, v/v) and eluent B (0.1% methanoic acid in acetonitrile, v/v) in a liner gradient program (0–2 min, 10% B; 2–26 min, 10 \rightarrow 90% B; 26–28 min, 90% B; 28–28.1 min, 90 \rightarrow 10% B; 28.1-40 min, 10% B) with a flow rate of 0.4 mL min⁻¹. Set the temperature of column and the sample manager at 30 $^{\circ}$ C and 15 \degree C, respectively. 10% and 90% acetonitrile in aqueous solution were used as weak and strong wash solvents respectively.

The optimized MS parameters were as follows: source temperature (150 °C), desolvation temperature (400 °C), cone voltage (40 V), capillary voltage at 2.6 kV (ESI^+) and 2.2 kV (ESI^-) , cone gas flow (50 L h^{-1}) and desolvation gas flow (800 L h^{-1}). MS^E mode was chosen with low energy of 6 V and high energy of 20–40 V.26,27 The mass spectrometer was calibrated with sodium formate in the range of 100 to 1200 Da in order to ensure the mass reproducibility and accuracy. Leucine enkephalin (m/z) 556.2771 in $ESI⁺$ and 554.2615 in $ESI⁻$) was used as external reference for Lock Spray™ injected at a constant flow of 10 μ L min⁻¹. The QC sample was injected randomly 4 times
throughout the whole work list. All of the volume injection of throughout the whole work list. All of the volume injection of the samples and QC was $5 \mu L$ per run. During data acquisition, the data for screening analysis was performed in MS^E continuum mode, the data for metabolomics analysis was performed in MS^E centroid mode. Data recording was performed on MassLynx V4.1 workstation (Waters, Manchester, UK). RSC Advances Co., Milod, MA, USA) with an electro- and cCOOH and -11 as negative adducts. Leader on Republished on 26 February 2021 and the competence of the creative commons (Will the most one 26 February 2021 and the co

2.4. Screening analysis of components in three parts of COT by UNIFI platform

UNIFI 1.7.0 software (Waters, Manchester, UK) was used for data analysis.^{28,29}

Firstly, in addition to the internal Traditional Medicine Library on UNIFI platform, the chemical constituent investigation was conducted. As the result, a self-built database of chemical compounds isolated from the genus of Celastrus L. was established by searching the online databases including Web of Science, Medline, PubMed, ChemSpider and China National Knowledge Infrastructure (CNKI). The compound name, molecular formula and chemical structure of components were obtained in the database.

Secondly, the raw data obtained from Masslynx workstation were compressed by Waters Compression and Archival Tool v1.10, then were imported into the UNIFI software.

Thirdly, the compressed data were processed by the streamlined work flow of UNIFI software in order to quickly identify the chemical compounds which were matched the criteria with Traditional Medicine Library and self-built database. The main parameters of processed method were as follow: 2D peak detection was set to 200 as the minimum peak area. In the 3D peak detection, the peak intensity of high energy and low energy was taken more than 200 and 1000 times as the parameter respectively. Selected +H and +Na as positive adducts

and +COOH and –H as negative adducts. Leucine enkephalin was used as reference compound in order to get exact mass accuracy, with $[M + H]$ ⁺ 556.2766 for positive ion and $[M - H]$ ⁻ 554.2620 for negative ion. As a result, the comprehensive chemical constituents screening list was accomplished.

Finally, a filter was set to refine the results, with the mass error between -5 and 5 ppm and response value over 5000. Each compound was verified by compared with the characteristic MS fragmentation patterns reported in literature or the retention time of the reference substances.

2.5. Metabolomics analysis of three parts of COT

The raw data acquired by Masslynx workstation was processed on MakerLynx XS V4.1 software (Waters, Milford, CT, USA). Firstly, the raw data were processed with alignment, deconvolution, and data reduction, etc. The main parameters of the process method were as follows: retention time (0–28 min), retention time window (0.20), mass (100–1200 Da), mass tolerance (0.10), mass window (0.10), minimum intensity (5%), marker intensity threshold (2000 counts) and noise elimination (level 6). As a result, the list with mass and retention time corresponded to the responses based on all the detected peaks from each data file were shown in Extended Statistics (XS) Viewer. Secondly, multivariate statistical analysis, both principle component analysis (PCA) and orthogonal projections to latent structures discriminant analysis (OPLS-DA), were performed on the MakerLynx software to analyze the resulting data.

PCA, a classical unsupervised low dimensional pattern recognition model, was used to show pattern recognition and maximum variation, and the overview and classification were obtained. OPLS-DA was used to obtain the maximum separation between two different groups. S-plots, which could provide visualization of the OPLS-DA predictive results, were created to explore the potential chemical markers which contributed to the differences.

Meanwhile, metabolites with VIP value > 4.0 and p-value $<$ 0.001 were considered as potential chemical markers.^{30,31} Futhermore, permutation test was also performed to provide a reference distribution with the R^2/Q^2 values to indicate statistical significance. Finally, the analysis results were shown in Simca 15.0 software (Umetrics, Malmö, Sweden).

2.6. Bioassay analysis of three parts of COT

2.6.1 Preparation of cigarette smoke extract. The preparation of the cigarette smoke extract (CSE) was basically the same as previous reports.³² Put the smoke from one cigarette into 20 mL culture medium (300 s per cigarette). The CSE solution was incubated at 37 $\mathrm{^{\circ}C}$ for 30 min after being filtered with a 0.22 um sterile filter. The CSE solution was prepared freshly and was used within 30 min. This prepared CSE solution was considered to have the highest concentration (100%).

 $2.6.2$ Cell viability assay. The final concentrations $(20.0,$ 40.0, 80.0, 160.0, 320.0 μ g mL⁻¹) of each test samples (R_{bio} , S_{bio} and Lbio) were acquired by diluting the stock solutions with Dulbecco's Modified Eagle Medium (DMEM). A549 cells were

cultured in 96-well plates at a density of 5×10^5 cells per well treated with CSE (0%, 5%, 10%, 20%, 30% and 40%) for 18 h, or treated with R_{bio} , S_{bio} and L_{bio} solutions (0.0, 20.0, 40.0, 80.0,

160.0, 320.0 μ g mL⁻¹) for 24 h. The growth-inhibition effect of the drugs on vishility of 4540 sells were CSE and the effect of the drugs on viability of A549 cells were evaluated by MTT assay.

Fig. 1 The base peak intensity (BPI) chromatograms in root, stem and leaf of COT in ESI⁺ and ESI⁻.

2.6.3 Drug treatment. For all groups, A549 cells were cultured in 96-well plates at a density of 5 \times 10^5 cells per mL for 18 h. In CSE group, the cells were treated with a certain dose of CSE without drug intervened. In drug groups, the cells were treated with both CSE and drugs (R_{bio}, S_{bio} or L_{bio}). In control group, A549 cells were cultured normally without the CSE or

Fig. 2 Chemical structures of compounds identified in the root, stem and leaf of COT.

Fig. 2 (contd.)

Fig. 2 (contd.)

drugs. In positive group, the cells were treated with both CSE and dexamethasone $(5 \mu g \text{ mL}^{-1})$.

2.6.4 Enzyme-linked immunosorbent assay. The contents of IL-1 β , IL-6 and TNF- α in the cell culture supernatant were determined with ELISA kits. All procedures were performed according to the manufacturer's instructions.

2.6.5 Statistical analysis. Statistical analysis was performed on Graphpad Prism 6.0 software (CA, USA). The results were expressed as mean \pm SD. Two tailed test or a one-way analysis of variance (ANOVA) was used to calculate statistical significant difference ($p < 0.05$).

3. Results and discussion

3.1. Screening analysis of components of three parts of COT

A total of 120 compounds, including 91 in $ESI⁺$ mode and 29 in ESI⁻ mode, were identified or tentatively characterized from three parts of COT (Table 2). The base peak intensity (BPI) chromatograms were shown in Fig. 1. The chemical structures were shown in Fig. 2, the results showed that COT was rich in natural components with various structural patterns. On one hand, according to the reference, there were nearly 50, 100, 10 compounds were reported from the root, stem, leaf parts of COT,

Fig. 2 (contd.)

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respectively. While in this study, there were 92, 56 and 32 components were identified or tentatively characterized from root part, stem part and leaf part of COT, respectively. And most of the components were identified from COT for the first time. Various kinds of structures, including triterpeniods,

sesquiterpenoids, steroids, flavonoids, organic acid and organic acid esters, phenylpropanoids, diterpeniods, monoterpenoids, alkaloid and others, were contained in each part of COT. The numbers (% of the total identified components in each part) and structural types of compounds identified from root, stem and leaf

Fig. 4 The PCA of root (R), stem (S), leaf (L) groups in ESI^+ and ESI^- . QC: quality control.

Fig. 5 OPLS-DA plots/permutation tests/S-plots between R and S&L. R: the root of Celastrus orbiculatus Thunb.; S: the stem of Celastrus orbiculatus Thunb.; L: the leaf of Celastrus orbiculatus Thunb.

Fig. 6 OPLS-DA plots/permutation tests/S-plots between S and R&L. R: the root of Celastrus orbiculatus Thunb.; S: the stem of Celastrus orbiculatus Thunb.; L: the leaf of Celastrus orbiculatus Thunb.

of COT were shown in Fig. 3. There were 34, 18 and 10 triterpeniods identified from root, stem and leaf, respectively, accounted for 37%, 32% and 31% of the total components in each part. So,

it was concluded that triterpeniods were the major constituents in three parts of COT. Moreover, according to each percentage, the root part of COT was also rich in organic acid and organic

Fig. 7 OPLS-DA plots/permutation tests/S-plots between L and R&S. R: the root of Celastrus orbiculatus Thunb.; S: the stem of Celastrus orbiculatus Thunb.; L: the leaf of Celastrus orbiculatus Thunb.

Fig. 8 Heat-map visualizing the intensities of the potential chemical markers.

Fig. 9 The cytotoxicity effects of different concentrations of cigarette smoke extract (CSE) on A549 cells. **p < 0.01, compared with 0% CSE group

acid esters, steroids and phenylpropanoids. The stem part was also rich in organic acid and organic acid esters, and flavonoids. The leaf part was also rich in steroids, and sesquiterpenoids. The percentage of flavonoids in the total identified components in stem was higher than the percentages in root part or in leaf part. The percentages of sesquiterpeniods and steroids in the total identified components in leaf was higher than the percentages in root part or in stem part. On the other hand, the shared components (30 for root and stem, 22 for root and leaf, 23 for stem and leaf, 15 for root, stem and leaf) were also found in our study. As shown in Fig. 3, the structures of shared components were various, while triterpeniods held the majority. Celastrol, one of the triterpeniods, was shown to distribute in root, stem and leaf, which was consistent with the ref. 19. So our research work could provide the scientific data to clarify the chemical composition of COT, particularly for the root and the leaf parts.

Although the study provided evidences to elucidate the chemical composition of COT, there were still some unresolved issues. For example, as shown in BPI chromatograms, there were some unidentified components. Further research should be carried on the identification of these unknown compounds.

3.2. Metabolomics analysis of three parts of COT

PCA score 2D plots in both $ESI⁺$ and $ESI⁻$ were established as shown in Fig. 4. The QC samples were clustered tightly and were in the middle of the three groups in PCA, which indicated the system had satisfactory stability. The samples from root of COT were clearly gathered together, which indicated there was a good similarity among them, and this phenomenon was also observed in stem and leaf of COT. Meanwhile, the root, stem and leaf groups were easily divided into three clusters, indicating that these three parts of COT could be differentiated in both $ESI⁺$ and $ESI⁻$.

In order to further distinguish one part from the other two parts, OPLS-DA plots, S-plots, permutation tests, and VIP values were obtained to see which variables were responsible for sample separation⁹⁷ (Fig. 5-7). In OPLS-DA plots, each spot represented a sample. From the perspective of OPLS-DA, one part was clearly separated from the other two parts. The parameters such as R^2 and Q^2 indicated the model had good

Fig. 10 The cell viability effects of different concentrations of R_{bio}, S_{bio} and L_{bio} solution on A549 cells. **p < 0.01, compared with 0 µg mL⁻¹ group.

ability of prediction and reliability in both ESI^+ and ESI^- modes. The permutation plots showed the original point on the right was clearly higher than all Q^2 -values (blue) on the left, which indicated the original models were valid. To identify the metabolites contributing to the discrimination, S-plots were generated under OPLS-DA model. Each spot in S-plots represented a variable. The variables with VIP > 4 and $p < 0.001$ were considered as potential chemical markers. The possible molecular formula of the markers were calculated by highaccuracy quasi-molecular ion with mass error between \pm 5 ppm. A total of 26 robust known chemical markers (marked in Table 2) enabling the differentiation between one part with the other two parts were identified and marked in S-plots.

According to the reference, it was revealed that there was significant variation for the contents of celastrol or total alkaloids in different parts of COT. While in this study, there were 13, 8 and 5 potential chemical markers including celastrol discovered from root, stem and leaf, respectively. The markers in root including 8 triterpenoids (35, 39, 50, 51, 63, 80, 88, 90), 1 steroids (57), 1 organic acid esters (87), 1 phenylpropanoids (103) and 2 other compounds (73, 86). The markers in stem including 4 triterpenoids (64, 84, 85, 100), 1 flavonoids (2), 1 phenylpropanoids (30), 1 steroids (52) and 1 organic acid esters (61). The markers in leaf including 3 sesquiterpenoids $(11, 40, 117)$, 1 flavonoids (14) and 1 triterpenoids (54). Additionally, among these potential chemical markers, the contents of 57 and 88 in root, 52 in stem, 40, 54 and 117 in leaf were much higher than in the other two parts ($p \leq$

Fig. 11 The effects of inflammatory cytokine IL-1B, IL-6 and TNF- α on root, stem and leaf (20–160 µg mL⁻¹) of COT in CSE-stimulated A549
cells ttH p < 0.01 compared with control group: *p < 0.05 compared with CSE g cells. ^{tt}p < 0.01, compared with control group; *p < 0.05, compared with CSE group; **p < 0.01, compared with CSE group.

0.001). While components 35, 39, 50, 51, 63, 73, 80, 86, 87, 90 and 103 were detected only in root, components 2, 30, 61, 64, 84, 85 and 100 were detected only in stem, components 11 and 14 were detected only in leaf part under the detect condition.

The detected result of 88 (celastrol) in our study, with much higher contents in root than in stem and leaf, was consistent with the ref. 19. While there were a few differences between our results and the references. In the present study, compound 39 (pristimerin) was only detected in root, and 11 (orbiculin I) was only detected in leaf. According to the reports, 39 was once isolated from stem¹⁴ though mainly from root,^{98,99} and 11 isolated from root.³ The reason was the concentrations of them were lower than the lowest detection limits. It was worth mentioning that some chemical markers with high responses in UPLC-MS, two triterpenoids (39 and 88) in root, one flavonoids (2) in stem and two sesquiterpenoids (11 and 40) in leaf, could be used for further quality control of three parts of COT respectively.

In order to systematically evaluate the chemical markers, a heat-map was generated. The hierarchical clustering heat map, intuitively visualizing the difference level of potential chemical markers in different parts, was shown in Fig. 8. The higher values were indicated by red squares, the lower values were indicated by green squares.

3.3. Bioactivity evaluation

3.3.1 Cytotoxicity of CSE and the three parts of COT on the viability of A549 cells. The results of MTT showed that the viability of A549 cells was obviously affected $(p < 0.01)$ by 30% or 40% CSE (Fig. 9). Therefore, 20% CSE was chosen as stimulus in the following experiments. Additionally, as shown in Fig. 10, the viability of A549 cells were not significantly affected by the R_{bio} , S_{bio} and L_{bio} solutions at 20–160 µg mL⁻¹. So we evaluated the officite of R_{i} set and L₁ colutions at 20, 160 µg mL⁻¹ on CSE effects of R_{bio} , S_{bio} and L_{bio} solutions at 20–160 μ g mL⁻¹ on CSEstimulated A549 cells.

3.3.2 Effect of root, stem and leaf of COT on CSEstimulated pro-inflammatory cytokine levels in A549 cells. The inflammatory development was characterized by the release of pro-inflammatory mediators such as interleukin-1 β (IL-1 β), interleukin-6 (IL-6) and tumor necrosis factor- α (TNF- α). Whether the root, stem and leaf of COT could inhibit the release of IL-1 β , IL-6 and TNF- α in CSE-stimulated lung epithelial cells was investigated in this paper. As shown in Fig. 11, the production of IL-1 β , IL-6 and TNF- α in A549 cells was obviously increased after treated with CSE $(p < 0.01)$. However, treated with the R_{bio} , S_{bio} and L_{bio} solutions could evidently decrease the levels of pro-inflammatory factors in a good dose-dependent way with a certain range of 20-160 μ g mL⁻¹ in CSE-stimulated cells. The R_{bio} solution could significantly decrease the levels of IL-1 β , TNF- α (80 μ g mL⁻¹, $p < 0.05$; 160 μ g mL⁻¹, $p < 0.01$) and
IL 6 (40 and 80 ug mL⁻¹, n < 0.05; 160 ug mL⁻¹, n < 0.01). The IL-6 (40 and 80 μ g mL⁻¹, p < 0.05; 160 μ g mL⁻¹, p < 0.01). The S_{bio} solution could significantly decrease the levels of IL-1 β , TNF- α (40 μ g mL⁻¹, p < 0.05; 80 and 160 μ g mL⁻¹, p < 0.01) and
H \leq (20 and 40 wg mL⁻¹, p < 0.05; 90 and 450 wg mL⁻¹, p < 0.01) IL-6 (20 and 40 μ g mL⁻¹, $p < 0.05$; 80 and 160 μ g mL⁻¹, $p < 0.01$).
The Luck solution could circuificantly decrease the loyals of H 6 The L_{bio} solution could significantly decrease the levels of IL-6 $(80 \text{ µg mL}^{-1}, p < 0.05; 160 \text{ µg mL}^{-1}, p < 0.01) \text{ and TNF- α (80$

and 160 μ g mL⁻¹, p < 0.05), but showed no significantly effect on
H 18. The shows results showed that the S_{nu} solution had IL-1 β . The above results showed that the S_{bio} solution had a stronger anti-inflammation effect than R_{bio} and L_{bio} . It is suggested that to explore the anti-COPD effect of the COT stem in vivo is meaningful in further research. The different activities of root, stem and leaf of COT might be caused by the various phytochemicals in these three parts of COT. The phytochemical study showed that three parts of COT were differentiated. The bioassay study showed that three parts of COT could reduce the levels of pro-inflammatory factors to varying degrees. And the stem part had a stronger anti-COPD effect than root and leaf parts. As we all know, the material basis of different pharmacological activities is the different chemical composition. The results of the two parts of our study showed that the different activities of root, stem and leaf of COT might be caused by the various phytochemicals in these three parts of COT. RSC Advances Compare to the comparison at θ , θ ,

4. Conclusions

In conclusion, for screening analysis, a total of 120 compounds (15 shared components), including 92 from root, 56 from stem and 32 from leaf, were identified or tentatively characterized from COT. Each part of COT was rich in various kinds of structures, especially triterpeniods held the majority. For metabolomic analysis, the root, stem and leaf of COT were differentiated in both ESI^+ and ESI^- modes. There were 13, 8 and 5 potential chemical markers identified from root, stem and leaf, respectively. Among the above robust markers, 5 robust chemical markers with high responses in UPLC-MS, 2 triterpenoids (pristimerin and celastrol) in root, 1 flavonoids $\{5,7\text{-dihydroxy-6},8\text{-dimethyl-3}(S)-3\text{-}(3\text{-methoxy-4'-hydroxx-6})\}$

ybenzyl) chroman-4-one} in stem and 2 sesquiterpenoids (orbiculin I and orbiculin A) in leaf, could be used for further quality control in three parts of COT respectively. For bioassay analysis, the root, stem and leaf of COT could evidently reduce the levels of pro-inflammatory factors in a dose-dependent way within a certain range of 20-160 μ g mL⁻¹ in CSE-induced A549 cells. The results showed that the stem part had a stronger anti-COPD effect than root and leaf parts. The different activities might be caused by the various phytochemicals in these three parts of COT. This comprehensive phytochemical study revealed both the structural diversity of secondary metabolites and the different distributions in different parts of COT. It could provide a theoretical basis for further utilization and development of COT. And the identification of anti-COPD components from COT will be explored deeply in the future based on the current results of this study.

Conflicts of interest

The authors declare no conflicts of interests.

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