



UPLC-Q-TOF-MS Study of the Mechanism of THSWD for Breast Cancer Treatment

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Taohong Siwu decoction (THSWD) is a classic traditional Chinese medicine (TCM) prescription that is widely used in the clinical treatment of gynecological and cerebrovascular diseases. Here we used a method that coupled ultra-performance liquid chromatography to quadrupole time-of-flight mass spectrometry (UPLC-Q-TOF-MS) in which both positive and negative ion modes were established to investigate the major constituents in THSWD. A Waters ACQUITY UPLC BEH C18 column (2.1 mm×100 mm, 1.7 μm) was used to separate the aqueous extract of THSWD. The mobile phase consisted of 0.1% aqueous formic acid (A) and acetonitrile (B). Ninety-five components were identified in two different ion modes, including aromatic acids, flavones, polysaccharides, volatile oils monoterpene glycosides, aromatic cyanogenic glycosides, and others. Pathological changes in tumors and serum expression of interleukin-4 in a mouse model of breast cancer were detected after THSWD treatment. The results showed that THSWD had obvious therapeutic effects. This study establishes a material basis for the use of THSWD in the treatment of breast cancer.

Keywords: Taohong Siwu decoction, breast cancer, UPLC-Q-TOF-MS, components, rats

INTRODUCTION

Traditional Chinese medicine (TCM) is an important part of Chinese culture (Zhen, 2013). As the most representative and integral part of traditional Chinese culture, TCM has a long, profound history. At present, the Party and State attach great importance to TCM, which also has a far-reaching influence all over the world (Weiwei, 2017). Formulae are the main form of TCM clinical medicine (Chen and Luo, 2018). They are characterized by multiple components, which affect multiple pathways and targets, so their mechanisms of action are largely unknown (Li et al., 2012). The “Traditional Chinese Medicine Law of the People’s Republic of China” is its official implementation, marking the first time that China identified the important status, development guidelines, and supporting measures of TCM from a legal perspective and provided a legal guarantee for TCM development.

Taohong Siwu decoction (THSWD) consists of six species of medicinal herbs: *Prunus persica* (L.) Batsch (Taoren, TR), *Carthamus tinctorius* L. (Honghua, HH), *Angelica sinensis* (Oliv.) Diels (Danggui, DG), *Ligusticum chuanxiong hort* (Chuanxiong, CX), *Paeoniae Radix Alba* (Baishao, BS), and *Rehmanniaglutinosa* (Gaertn.) DC. (Shudi, SD) (Ding and Zhang, 2010). THSWD was

documented in “Yi Zong Jin Jian” compiled by Wu Qian in the Qing Dynasty, on the basis that the Siwu decoction enriches the blood, adds TR and HH into the blood, eliminates blood stasis, and promotes blood circulation and nourishment (Liu et al., 2015). It has been widely used in the treatment of gynecological and cerebrovascular diseases (Kim et al., 2016). Ischemic stroke, also known as cerebral infarction due to thrombosis, belongs to the category of “stroke” in TCM (Yu et al., 2007). It is an acute cerebrovascular disease characterized by sudden stupor, unconsciousness, hemiplegia, slurred speech, and speech disorder (Cheng et al., 2018).

Considering the characteristics of a TCM compound, the composition is complex. Moreover, factors like origin, cultivation, harvesting, processing, preparation, and more all influence the clinical treatment effect (Zhang, 2010). Therefore, verifying the chemical composition of TCM compounds through analysis and quality testing is particularly important. It is quite time-consuming to chemically profile TCM preparations with conventional high-performance liquid chromatography coupled with ultraviolet detection (HPLC-UV) or HPLC coupled with mass spectrometry (HPLC-MS) (Han et al., 2007; Li et al., 2014). Compared with traditional HPLC technology, which remains constant over a wider linear range, ultra-performance liquid chromatography (UPLC) has higher efficiency. Its advantages include increasing the mobile phase velocity, shortening the analysis time, and increasing throughput. The columns show superior performance, and the ultra-high pressure liquid chromatography pump offers precise gradient control, low diffusion, a low cross-contamination auto-sampler system, a high-speed detector, and comprehensive hardware and software. The peak capacity, analysis efficiency, and sensitivity are greatly improved compared with conventional HPLC, providing a good platform for the separation analysis of complex systems (Gumustas et al., 2014; Hedaya et al., 2017; Wu et al., 2017). Time-of-flight MS (TOF-MS) can provide accurate molecular mass (Zhang et al., 2018). MassHunter software can calculate the molecular formula of the test substance, so the chemical composition represented by each chromatographic peak can be directly identified with very small errors, generally at 5 ppm. This approach is widely used in the identification of compounds. In this study, UPLC-Q-TOF-MS technology was used to separate and identify the main components of THSWD.

THSWD is used to treat cerebrovascular and gynecological diseases. DG and CX have good therapeutic effects on gynecological conditions, especially breast diseases. However, there have been no reports of the study of THSWD in the treatment of breast cancer. We analyzed the components of THSWD and assessed its therapeutic effect in a mouse model of breast cancer to provide a basis for its clinical use.

MATERIALS AND METHODS

Materials

Prunus persica (L.) Batsch (Taoren, TR, batch number: 17033101), *Carthamus tinctorius* L. (Honghua, HH, batch

number: 17041401), *Angelica sinensis* (Oliv.) Diels (Danggui, DG, batch number: 16070501), *Conioselinum anthriscoides* ‘Chuanxiong’ (syn. *Ligusticum chuanxiong* Hort) (Chuanxiong, CX, batch number: 17061601), *Paeoniae lactiflora* Pall. (Baishao, BS, batch number: 17050301), and *Rehmannia glutinosa* (Gaertn.) DC (Shudi, SD, batch number: 17042501) were purchased from Anqing Huashi Chinese Herbal Medicine Co. Ltd. (Anqing, China). All TCM materials were qualified by Professor Huasheng Peng (hspeng@126.com). All voucher specimens were deposited at the Herbarium of Anhui University of Chinese Medicine, Hefei, China (Herbarium code: ACM, voucher numbers: 17021, 17025, 17034, 17051, 17068, 17080). A Waters ACQUITY™ UPLC I-Class and a SYNAPT-G2-Si MS were obtained from Waters Corporation (Milford, MA, USA). The 4T1 breast cancer cell line and fetal bovine serum (batch number: 20160720) were from Zhejiang Tianhang Biotechnology Co., Ltd. RPMI 1640 medium (batch number: ZI110516) was purchased from Hangzhou Northrend Biotechnology Co., Ltd.

Preparation and Extraction

In this system, the herbs TR, HH, DG, SD, CX, and BS (3:2:3:4:2:3) were soaked then decocted twice with 10 vol. of boiling water for 2 h and 8 vol. of boiling water for 1.5 h. The extraction solutions were then filtered and combined. Next, 95% ethanol was added for alcohol precipitation to obtain Taohong-Siwu decoction alcohol precipitation. A small amount of alcohol solution was filtered through a 0.22- μ m microporous membrane filter, and 2 μ L of filtrate was used for injection analysis.

UPLC-MS and UPLC-MS² Analysis

A Waters ACQUITY UPLC BEH C18 column (2.1 mm \times 100 mm, 1.7 μ m) was used to separate the aqueous extract of THSWD. The mobile phase was 0.1% aqueous formic acid (A) and acetonitrile (B). Chromatographic separation was performed at 35°C in this system. Gradient elution with a flow rate of 0.3 mL/min was performed as follows: 3% B at 0-2 min; 3%-8% B at 2-8 min; 8%-25%B at 8-12 min; 25%-25% B at 12-15 min; 25%-45% B at 15-16 min; 45%-90% B at 16-22 min; 90%-100% B at 22-26 min; 100% B at 26-28 min. The MS analysis was carried out with the electrospray ionization (ESI) source in both positive and negative ion modes, and leucine enkephalin was used as the accurate mass calibration solution. The desolvation gas temperature was 350°C. The flow rates of cone and desolvation gases were set at 50 L/h and 600 L/h, respectively. The capillary, cone, and extraction cone voltages were set to 3.0 and 2.5 V in positive and negative ion modes, respectively. MS^E was applied for MS/MS analysis with a low collision energy of 6 V and a high collision energy of 20-80V. The scan area was set at *m/z* 50-1200.

Animal Treatment and Sample Collection

BALB/c mice (specific pathogen-free grade, 22 \pm 2g), were purchased from the Laboratory Animal Center, Medical University of Anhui Province (permit number: 2016AH-031-11). They were randomly divided into six groups (10 mice each): normal; model; THSWD low-dose, middle-dose, and high-dose

(L, M, and H); and cisplatin. After 4T1 cells were resuscitated, they were routinely cultured in RPMI 1640 medium containing 10% fetal bovine serum at 5% CO₂ and 37°C. The culture medium was changed every 2 to 3 days and passaged. Cells cultured to logarithmic growth phase were collected and diluted 10 times with sterile saline, and 0.2 mL was injected into each mouse. After 48 hours, there was a small pink protrusion of about 2 mm × 2 mm, indicating successful modeling. Mice in the THSWD H, THSWD M, and THSWD L groups were treated with THSWD (25.2, 12.6, and 6.3 g/kg, respectively, three times a day) by gavage. The cisplatin group received cisplatin (0.1 mg/mL) by gavage, the normal and model groups were administered with an equivalent amount of regular saline. The dosing period was 18 days. After the last administration, blood and tumor samples were collected and animals were sacrificed by cervical vertebrae dislocation. All experiments were subject to approval by the Committee on the Ethics of Animal Experiments of Anhui University of Chinese medicine (Permit Number: LLSC20160336).

Hematoxylin and Eosin (H&E) Staining

Tumor tissues were fixed, paraffin-embedded, and sectioned. Changes in tumor cell arrangement and pathological alterations of the tumor were observed with conventional H&E staining and analyzed using ImageJ software (National Institutes of Health, Bethesda, MD, USA).

Interleukin (IL)-4, IL-10, IL-13, and Transforming Growth Factor (TGF)-β1 Expression

Serum was collected from mice and centrifuged at 350 g for 5 min at 4°C. Supernatants were stored at -80°C for cytokine analysis. Enzyme-linked immunosorbent assays (ELISAs) were performed to quantify the concentrations of IL-4, IL-10, IL-13, and TGF-β1, according to the manufacturer's instructions. Optical density (OD) values were read at 450 nm, and the levels of IL-4, IL-10, IL-13, and TGF-β1 were determined based on standard curves.

CD44, CD117, and CD133 Expression

At 24 hours after transfection, the cells were removed and washed with phosphate-buffered saline (PBS; 5 min, 3 times), fixed in methanol (-20°C, 5 min), fixed in 70% ethanol for 5 min, and washed with PBS (5 min, 3 times). They were then blocked in 1% skimmed milk powder for 30 min, and the mixture was incubated for 2 h at room temperature with primary antibody. The cells were washed with PBS, incubated with secondary antibody for 1 h, stained with 4',6-diamidino-2-phenylindole for 2 min, washed with PBS (5 min, 3 times), mounted in DAKO medium (DAKO, Glostrup, Denmark), observed under a fluorescent microscope, and photographed.

Data Processing

Establishment of a Component Analysis Database

We used the UNIFI software library management system to collect data from offline and online MS databases such as PubMed (<http://www.ncbi.nlm.nih.gov/pubmed>), MassBank

(<http://www.massbank.jp/>), Chemspider (<http://www.chemspider.com/>), and METLIN (<https://en.wikipedia.org/>) and identify relevant literature on the chemical composition of THSWD. We collected the names and molecular and structural formulas to build a database of the chemical constituents of THSWD.

Data Acquisition and Analysis

The MS^E Continuum model was used to collect data and determine the chemical composition of THSWD, which was automatically identified by the UNIFI data processing system. MassLynx 4.1 software (Waters Corporation) was used to deal with the base-peak ion current patterns of positive and negative ion modes. The main chemical components were identified and confirmed by the accurate mass of fragment ions theory, relative retention time, offline and online mass spectral database, and related literature.

RESULTS AND DISCUSSION

Optimization of Chromatographic Conditions and Q-TOF-MS/MS Method Development

Due to the diversity of TCM prescriptions, the complexity of each single herb, and the large variety of structures, it is necessary to optimize the chromatographic conditions to obtain and separate the compounds quickly and improve the resolution. The items to be improved are the mobile phase, flow rate, column temperature, and MS parameters. For the mobile phase, we chose aqueous formic acid (A) and acetonitrile (B). We set the column temperature at 35°C. The detection wavelength was based on the literature, and full-band scans were performed at 320 nm with a flow rate of 0.3 mL/min to reduce the column pressure and shorten the analytical time. The elution procedures and MS conditions were set according to section 2.3. The standard sample solution and medicinal sample solution of each ingredient in THSWD were precisely absorbed. The total ion chromatogram of the THSWD solution was obtained according to the liquid-phase and MS conditions and the injection analysis. The base peak chromatograms of THSWD in positive and negative ion modes are shown in **Figures 1** and **2**, respectively.

Analysis of THSWD by UPLC-Q-TOF-MS and Identification of Its Main Constituents

The molecular weights of the 87 peaks extracted from the total ion chromatogram of the extract of THSWD were consistent with the data of the previously built pool, including aromatic acids, flavones, polysaccharides, volatile oils, monoterpene glycosides, aromatic cyanogenic glycosides, and others (**Table 1** and **Table 2**). Their sources were confirmed by comparing the base peak chromatograms of THSWD with its six separate herbal extracts. Standard compounds available were identified by comparing retention time with the exact mass. For

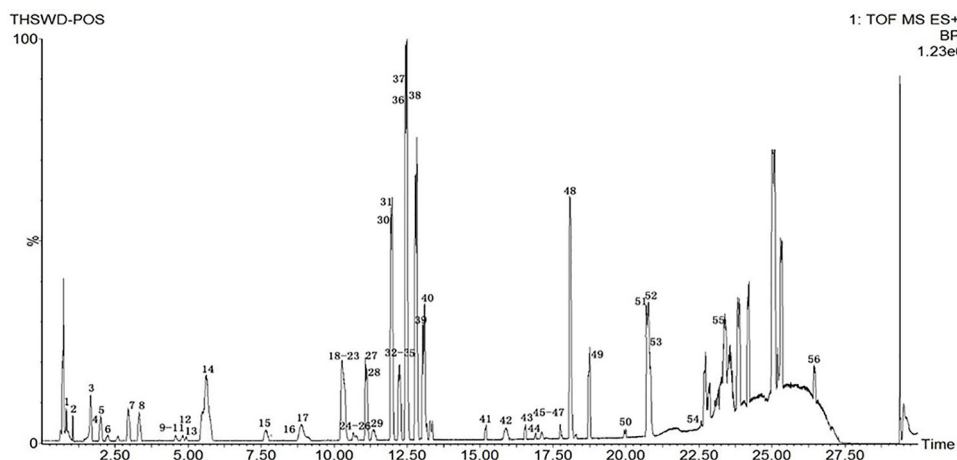


FIGURE 1 | Base peak chromatograms of THSWD in positive ion mode.

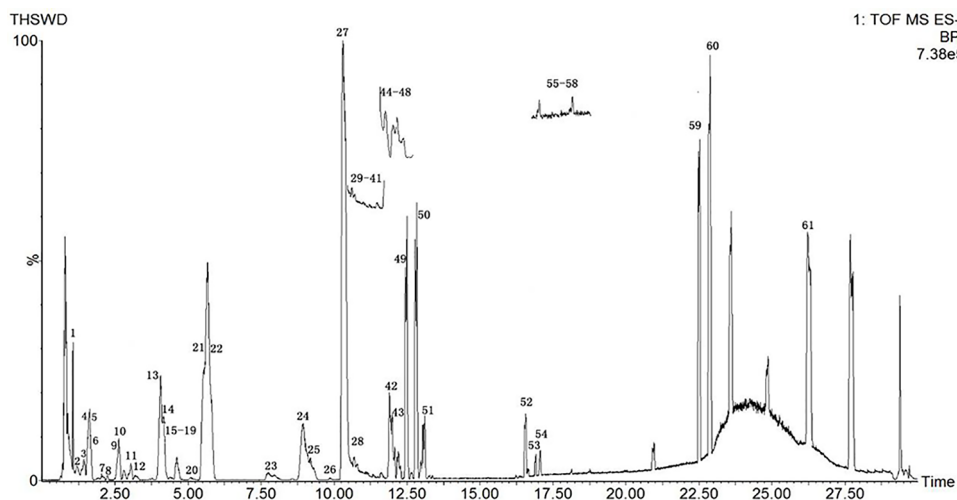


FIGURE 2 | Base peak chromatograms of THSWD in negative ion mode.

compounds that were not standard, the structure was primarily based on accurate mass and tandem MS. In this study, the formulas were based on the high-precision excimer $[M+H]^+$, $[M+Na]^+$, $[M-H]^-$, and $[M+HCOO]^-$ within a mass error of 10 ppm and a partial isotopic abundance. We then searched chemical databases such as Chemspider (www.chemspider.com) and Massbank (<http://www.massbank.jp>) for the most rational formulas. When all isomers were matched, the structures previously reported from the six herbs of THSWD had a higher probability than the other isomers. Finally, fragment ions were used to confirm the chemical structure further. The structures underlying the main activity in THSWD are shown in **Figure 3**.

Phenolic Acids

Seven phenolic acids of THSWD were identified in positive ion mode. Most from *Conioselinum anthriscoides* 'Chuanxiong' and

Angelica sinensis (Oliv.) Diels were ferulic acid, chlorogenic acid, caffeic acid, and vanillic acid, and gallic acid was from *Paeonia lactiflora* Pall. The retention time and MS information of each chemical component in THSWD were determined by UPLC-MS, and the chemical composition was confirmed by combining the extracted ion current map with data in the relevant literature. Compound 41 showed the t_R at the 15.2-min peak obtained in ESI mode at the m/z 194.059 ion peak. The literature (Lin et al., 1998; Huang and Song, 2001; Lu et al., 2004) reports that the relative molecular mass of ferulic acid in *Conioselinum anthriscoides* 'Chuanxiong' (syn. *Ligusticum chuanxiong* Hort) was 194. It was speculated that m/z 194.059 was its quasi-molecular ion peak, and MS² analysis was performed on m/z 193, which were m/z 279.0198, 163.0395, 92.0262, and other fragment peaks. According to the elemental composition analysis, the molecular formula is $C_{10}H_{10}O_4$. It showed a

TABLE 1 | Compounds identified in THSWD by UPLC-Q-TOF-MS in positive ion mode.

Peak No.	Identify	Molecular formula	Mass error (ppm)	RT (min)	Measured mass [M+H] ⁺ /[M+Na] ⁺	Source ^a	MS2
1	8-Epiloganic acid	C ₁₆ H ₂₄ O ₁₀	0.7	1.08	-/399.1264	S	541.1764,201.0082,167.9970
2	Guanosine	C ₁₀ H ₁₃ N ₅ O ₅	1.8	1.52	284.0994/-	H	152.0565,135.0300
3	Gallic acid	C ₇ H ₆ O ₅	0.2	1.6	171.0288/-	H,B	365.1068,153.0187,98.9755
4	1'-O-galloylsucrose	C ₁₉ H ₂₆ O ₁₅	-6.2	1.98	-/517.1132	B	365.1098,261.0313,153.0174,120.0813
5	Rehmannioside D	C ₂₇ H ₄₂ O ₂₀	3.5	2.01	-/709.2186	S	365.1099,153.0174,120.0813
6	5-Hydroxymethyl-2-furfural	C ₆ H ₆ O ₃	-1.7	2.24	127.0388/-	S,D	181.9670,164.9865,109.0290
7	indole-3-carboxaldehyde	C ₉ H ₇ NO	-3.3	3.38	146.0596/-	H	188.0703,146.0596,143.0722,118.0646
8	P-Hydroxybenzoic acid	C ₇ H ₆ O ₃	0.1	3.76	139.039/-	H	201.0096,121.0296
9	Chlorogenic acid	C ₁₆ H ₁₈ O ₉	5	4	355.1041/ 377.0968	D	211.0240,181.0132,147.0446
10	Neo-carthamin	C ₄₃ H ₄₄ O ₂₄	1.1	4.02	451.124/-	H	313.0686,247.0266,211.0240,181.0132,147.0446
11	Safflor yellow A	C ₂₇ H ₃₀ O ₁₆	5.8	4.03	611.1642/-	H	353.1192,201.0065,105.0334
12	Syringin	C ₁₇ H ₂₄ O ₉	0.1	4.73	-/395.1313	H	411.1068,395.1348,161.0590
13	Rehmapicroside	C ₁₆ H ₂₆ O ₈	-8.1	4.77	-/369.149	H	411.1068,395.1348,161.0590
14	Amygdalin	C ₂₀ H ₂₇ NO ₁₁	-0.6	5.63	458.1581/ 480.1474	T	496.1171,480.1473
15	Prunasin	C ₁₄ H ₁₇ NO ₆	0.9	7.68	-/318.0951	T	334.0687,318.0956,201.0082
16	3-O-Feruloylquinic acid	C ₁₇ H ₂₀ O ₉	4	8.62	369.1195/-	T	
17	paeoni-florigenone	C ₁₇ H ₁₈ O ₆	1.1	8.88	319.118/-	B	503.1525,105.0335
18	6-Hydroxyapigenin-6-O-β-D-glucoside-7-O-β-D-glucuronide	C ₂₇ H ₂₈ O ₁₇	2	10.28	625.1412/-	H	519.1216,323.0439,133.0644
19	Ferulaldehyde	C ₁₀ H ₁₀ O ₃	-2.1	10.28	179.0699/-	C,D	519.1216,503.1522,179.0697,151.0750
20	Albiflorin	C ₂₃ H ₂₈ O ₁₁	-0.1	10.28	481.1631/ 503.1523	B	519.1216,503.1522,500.1425,179.0697,151.0750,133.0644
21	Paeonilactone B	C ₁₀ H ₁₂ O ₄	-1.7	10.28	197.0805/-	B	519.1216,503.1522,500.1425,179.0697,151.0750,133.0644
22	Anhydrosafflor yellow B	C ₄₈ H ₅₂ O ₂₆	6.2	10.29	1045.2885/-	H	519.1216,503.1522,500.1425,179.0697,151.0750
23	Lactiflorin	C ₂₃ H ₂₆ O ₁₀	-0.1	10.29	463.1598/-	B	519.1216,503.1522,500.1425,179.0697,151.0750,133.0644
24	Quercetin- 3,7 - O - β - D -glucoside	C ₂₇ H ₃₀ O ₁₇	0.7	10.54	627.156/ 659.1488	H	303.05
25	6-Methoxycoumarin	C ₁₀ H ₈ O ₃	-2.3	10.67	177.0542/-	D	519.1263,287.0570,177.0547,171.9834,89.0389
26	paeoniflorin	C ₂₃ H ₂₈ O ₁₁	0.7	10.75	-/503.1527	B	503.1530,201.0068,163.0387
27	6-hydroxykaempferol	C ₁₅ H ₁₀ O ₇	2.9	11.29	303.0503/-	T,H	491.2935,476.3285,453.3431
28	Cistanoside A	C ₃₆ H ₄₈ O ₂₀	-1.5	11.34	-/823.2619	D	693.1618,635.1593,351.1442,177.0546,145.0277
29	Benzoic acid	C ₇ H ₆ O ₂	7.2	11.36	-/145.027	B	351.1442,289.0703,177.0546
30	Galloylpaeoniflorin	C ₃₀ H ₃₂ O ₁₅	1.6	11.91	633.1752/ 655.1644	B	655.1640,153.0181,147.0443
31	Saffloquinoside A	C ₂₇ H ₃₀ O ₁₅	1	12.04	595.1663/ 617.159	H	503.1527,287.0553,177.0544
32	Isosafrole	C ₁₀ H ₁₀ O ₂	-1	12.13	163.0752/-	C,D	91.0546
33	Isorhamnetin-3-O-nehesperidin	C ₂₈ H ₃₂ O ₁₆	1.2	12.18	625.177/ 657.1698	C	814.5436,317.0659,302.0421,301.0351
34	6-hydroxyapigenin	C ₈ H ₄ O ₃	-4.5	12.19	287.0537/-	D	673.4566,579.2918,363.1112,317.1154,301.1413,160.9908,149.0233
35	Acetoside	C ₂₉ H ₃₆ O ₁₅	4.3	12.26	-/647.1974	S	287.0555,163.0388
36	Kaempferol 3-O-rutinoside	C ₂₇ H ₃₀ O ₁₅	6.5	12.4	595.1696/-	T,H	519.1306,503.1549,317.0680,302.0432,205.0721,181.0127,140.0510
37	cosmosiin	C ₂₁ H ₂₀ O ₁₀	2.6	12.4	433.114/-	C	301.1412,149.0232
38	Carthamidin	C ₁₅ H ₁₂ O ₆	0.6	12.59	289.0708/-	H	153.0181,147.0443
39	Senkyunolide H	C ₁₂ H ₁₆ O ₄	-1.8	13.13	-/247.0936	C	1165.1875,1045.4538,926.4874,789.3951,712.7139,601.8298
40	4-hydroxy-3-butylphthalide	C ₁₂ H ₁₄ O ₃	-0.7	13.15	207.1014/-	C,D	247.0952,147.0446,91.0545
41	ferulic acid	C ₁₀ H ₁₀ O ₄	4.8	15.2	-/217.0482	C,D	279.0198,163.0395,92.0262
42	safflospermidine A	C ₃₄ H ₃₇ N ₃ O ₆	1.8	15.81	584.2766/ 606.2693	H	204.1032,147.0438
43	oxybenzoyl-paeoniflorin	C ₃₀ H ₃₂ O ₁₂	1.3	16.55	-/607.1794	B	607.1794,249.0760,105.0335
44	Benzoylpaeoniflorin	C ₃₀ H ₃₂ O ₁₂	1.1	16.65	585.1973/-	B	133.0663
45	Jionoside B1	C ₃₇ H ₅₀ O ₂₀	-5.9	17.09	-/837.2738	S	228.2324,107.0853,105.0695
46	Z-Butylidene-phthalide	C ₁₂ H ₁₂ O ₂	-0.8	17.15	189.0908/-	C,D	608.1815,219.0692,128.0629
47	Ligustilide	C ₁₂ H ₁₄ O ₂	-2.9	17.22	191.1061/-	C,D	670.3756,447.2399,355.2629,181.1199,165.0908,92.0252

(Continued)

TABLE 1 | Continued

Peak No.	Identify	Molecular formula	Mass error (ppm)	RT (min)	Measured mass [M+H] ⁺ / [M+Na] ⁺	Source ^a	MS ²
48	4,5-dihydro-3-butylphthalide	C ₁₂ H ₁₆ O ₂	4	18.27	193.1231/-	C	242.2845
49	3-Butylphthalide	C ₁₂ H ₁₄ O ₂	1.1	19.12	191.1069/-	C,D	191.1081,128.0624,115.0542
50	Levistolide A	C ₂₄ H ₂₈ O ₄	0.6	20.62	381.2063/ 403.199	D	419.1643,311.2948,256.2643
51	Dibutyl phthalate	C ₁₆ H ₂₂ O ₄	2.7	20.75	-/301.1418	C	787.52097,743.4928,337.3110
52	3-Butylidene-7-hydroxyphthalide	C ₁₂ H ₁₂ O ₃	0.3	20.77	205.086/ 227.0787	C	301.1412,149.0232
53	Z-6,8',7,3'- diligustilide	C ₂₄ H ₂₈ O ₄	-0.6	20.82	381.2058/-	C	510.3556,492.3458,473.2644,327.2904,191.1072
54	Eugenin	C ₄₁ H ₃₀ O ₂₆	6.5	22.3	-/961.098	H	283.2633,281.2476
55	Daucosterol	C ₃₅ H ₆₀ O ₆	-9.4	23.27	-/599.4226	H,C,D	283.2632,243.1214,95.0858
56	Ligustrazin	C ₈ H ₁₂ N ₂	-1.9	26.77	137.1071/-	C	302.0432,205.0720,181.0127

^aH, T, D, C, B, S indicate *Prunus persica* (L.) Batsch, *Carthamus tinctorius* L., *Angelica sinensis* (Oliv.) Diels, *Conioselinum anthriscoides* 'Chuanxiong' (syn. *Ligusticum chuanxiong* Hort), *Paeoniae lactiflora* Pall., and *Rehmannia glutinosa* (Gaertn.) DC, respectively.

TABLE 2 | Compounds identified in THSWD by UPLC-Q-TOF-MS in negative ion mode.

Peak No.	Identify	Molecular formula	Mass error (ppm)	Observed RT (min)	Measured mass[M-H] ⁻ / [M+HCOO] ⁻	Source ^a	MS ²
1	8-epiloganic acid	C ₁₆ H ₂₄ O ₁₀	-2.1	1.09	-/421.1343	S	499.1666,290.0882,128.0352
2	Guanosine	C ₁₀ H ₁₃ N ₅ O ₅	0.2	1.53	282.0844/-	H	377.0849,341.1088,282.0841,161.0454,150.0421
57	Coumalic acid	C ₆ H ₄ O ₄	1	1.61	-/185.0093	H	169.0142
3	gallic acid	C ₇ H ₆ O ₅	-0.3	1.62	169.0142/-	H,B	169.0142
6	5-Hydroxymethyl-2-furfural	C ₆ H ₆ O ₃	-0.5	1.62	125.0244/-	S,D	169.0142
58	6-hydroxykaempferol-7-O-β-D- glucoside	C ₂₁ H ₂₀ O ₁₂	0.2	1.73	463.0883/-	T,H	329.0867,302.0675,287.0878,134.0475
5	Rehmannioside D	C ₂₇ H ₄₂ O ₂₀	3.2	2.06	685.2293/ 731.2275	S	493.1197,331.0622,313.0567,169.0145,164.0721
59	Melittoside	C ₂₁ H ₃₂ O ₁₅	0.4	2.17	523.1743/ 569.1725	H	491.1190,448.1490
7	indole-3-carboxaldehyde	C ₉ H ₇ NO	1.8	2.45	-/190.0513	H	589.1873,379.1206
60	protocatechuic acid	C ₇ H ₆ O ₄	0.9	2.65	153.0195/-	H,C	512.1358,510.1380,474.1612,150.0560
61	Mudanpioside F	C ₁₆ H ₂₄ O ₈	-1	3.37	343.1467/ 389.1449	B	379.1148
62	salicylic acid	C ₇ H ₆ O ₃	-0.5	3.78	137.0244/-	S,D	193.0515,134.0373
9	Chlorogenic acid	C ₁₆ H ₁₈ O ₉	-0.9	4.04	353.0875/-	H	325.0712,191.0561,119.0503
63	hydroxysafflor yellow A	C ₂₇ H ₃₂ O ₁₆	1.3	4.06	611.1626/-	H	503.1185,491.1189,403.1027,191.0561
64	Guaiacol	C ₇ H ₈ O ₂	-0.4	4.37	123.0451/-	C	495.1511,167.0360
65	Vanillic acid	C ₈ H ₈ O ₄	-0.5	4.38	167.0349/-	D	493.1197,331.0622,313.0567,169.0145,164.0721
66	oxypaeoniflorin	C ₂₃ H ₂₈ O ₁₂	-0.2	4.41	495.1507/ 541.158	H	465.1368,177.0568,167.0359,137.0244
12	syringin	C ₁₇ H ₂₄ O ₉	-1.9	4.76	-/417.1394	H	163.0393,119.0501
13	RehMapiroside	C ₁₆ H ₂₆ O ₈	-0.9	4.8	345.1624/ 391.1606	H	463.0886,301.0346,271.0257,151.0042,116.9285
67	caffeic acid	C ₉ H ₈ O ₄	-0.2	5.09	179.035/-	D,C	135.04438
24	Quercetin- 3,7 - O - β - D -glucoside	C ₂₇ H ₃₀ O ₁₇	-1	5.45	625.1404/-	H	463.0898,407.0280,301.0361,299.0217
68	p-Hydroxybenzaldehyde	C ₇ H ₆ O ₂	-1.1	5.46	121.0294/-	H	625.1423,605.0494,480.1163,463.0898,301.0361,299.0218, 146.9627
15	prunasin	C ₁₄ H ₁₇ NO ₆	-1.4	7.73	294.1051/ 340.1033	T	357.0950,332.0729,330.0740
69	p-Hydroxy-cinnamic acid	C ₉ H ₁₀ O ₂	-1	8.56	163.0399/-	H	609.1463,435.0922,285.0407,193.0511,134.0371,133.0294
16	3-O-Feruloylquinic acid	C ₁₇ H ₂₀ O ₉	-0.9	8.71	367.1031/-	T	
20	albiflorin	C ₂₃ H ₂₈ O ₁₁	-0.7	9.2	479.1628/ 525.161	B	542.1508,517.1288,515.1318
70	Mudanpioside E	C ₂₄ H ₃₀ O ₁₃	0.2	10.31	525.1615/-	B	517.1305,515.1323,449.1447,327.1084,165.0555

(Continued)

TABLE 2 | Continued

Peak No.	Identify	Molecular formula	Mass error (ppm)	Observed RT (min)	Measured mass[M-H] ⁻ / [M+HCOO] ⁻	Source ^a	MS ²
18	6-hydroxyapigenin-6-O-β-D-glucoside-7-O-β-D-glucuronide	C ₂₇ H ₂₈ O ₁₇	-0.6	10.43	623.1268/-	H	525.1612,517.1305,515.1323,449.1447,327.1084,121.0294
71	6-hydroxyapigenin-3,6-di-O-β-D-glucoside	C ₂₇ H ₃₀ O ₁₇	1.2	10.56	625.1417/-	H	625.1414,609.1461,463.0871,462.0796,301.0347,271.0248,243.0307,139.0042
41	Ferulic acid	C ₁₀ H ₁₀ O ₄	0.1	10.69	193.0507/-	H,C,D	193.0511,134.0370,133.0294,132.0210
72	Purpureaside C	C ₃₅ H ₄₆ O ₂₀	2.9	10.78	785.2533/-	S	623.2193,495.1525,193.0507,161.0244,137.0246
73	safflor yellow	C ₂₇ H ₃₀ O ₁₅	0.9	11.05	609.1466/-	H	301.0350,300.0277
74	ISOQUERCITRIN	C ₂₁ H ₂₀ O ₁₂	-0.8	11.14	463.0878/-	T,H	271.0248,243.0299,116.9286
75	kaempferol 3-O-β-sophoroside	C ₂₇ H ₃₀ O ₁₆	0.1	11.21	609.1462/-	T,H	609.1462,284.0325,227.0352,151.0040
76	Dihydrophaseic acid-4'-O-β-D-glucoside methyl ester	C ₂₂ H ₃₄ O ₁₀	-0.8	11.25	-/503.213	H	610.1502,493.1855,430.0922,361.1669,283.0260,145.0286
28	Cistanoside A	C ₃₆ H ₄₈ O ₂₀	3.3	11.34	799.2692/ 845.2765	D	799.2694,624.2226,623.2197,193.0509,175.0399,161.0245
11	safflor yellow A	C ₂₇ H ₃₀ O ₁₆	-1.5	11.38	611.1608/ 657.1681	H	609.1464,301.0351,300.0277
77	Rehmaionoside B	C ₁₉ H ₃₄ O ₈	-1.4	11.83	-/435.223	S	695.5073,595.1719,425.1937,389.0877,347.0778,149.0247
39	Senkyunolide H	C ₁₂ H ₁₆ O ₄	0.7	11.91	223.0977/-	C,D	631.1673,169.0140,161.0242
30	galloylpaeoniflorin	C ₃₀ H ₃₂ O ₁₅	0.9	11.92	631.1674/-	B	631.1673,613.1566,399.0927,313.0564,271.0464,169.0140,161.0242
10	neo-carthamin	C ₄₃ H ₄₄ O ₂₄	-1.1	11.93	449.1084/-	H	631.1673,313.0564,271.0464,169.0140,161.0242
36	Kaempferol 3-O-rutinoside	C ₂₇ H ₃₀ O ₁₅	0.7	12.05	593.1516/-	T,H	593.1513,285.0399,284.0327,255.0230,227.0348,175.0399
45	Jionoside B1	C ₃₇ H ₅₀ O ₂₀	2.6	12.08	813.2844/ 859.2916	S	593.1513,285.0399,284.0327,255.0230,227.0348,175.0399
78	p-Hydroxyacetophenone	C ₈ H ₈ O ₂	-1.2	12.13	135.045/-	H	801.1863,637.1470,467.2142,146.9655
37	cosmosiin	C ₂₁ H ₂₀ O ₁₀	-1.8	12.14	431.0976/-	H	801.1863,626.1655,146.9655
33	Isorhamnetin-3-O-nehesperridin	C ₂₈ H ₃₂ O ₁₆	1.6	12.19	623.1627/-	C	623.1624,315.0504,243.0300
79	kaempferol 3-O-D-glucopyranoside	C ₂₁ H ₂₀ O ₁₁	-1	12.27	447.0928/-	T,H	284.0330,227.0351,161.0242,133.0295
80	Isoacteoside	C ₂₉ H ₃₆ O ₁₅	-0.2	12.27	623.1985/-	T	574.1599,424.1033,364.0819,311.0931,161.0238,133.0197
81	Sophoricoside	C ₂₁ H ₂₀ O ₁₀	-0.5	12.41	431.1054/ 477.1036	H	611.1398,403.0806,287.0566,162.8394,119.0501
82	Jionoside A	C ₃₀ H ₃₈ O ₁₅	0	13.12	637.2138/-	D	451.3276,347.0761,174.9555,161.0247
23	lactiflorin	C ₂₃ H ₂₆ O ₁₀	-0.2	13.26	461.1525/ 507.1507	B	539.1899,497.1225,237.0770,121.0299
44	benzoylpaeoniflorin	C ₃₀ H ₃₂ O ₁₂	1.8	16.55	583.1905/ 629.1887	B	619.1593,121.0294
43	oxybenzoyl-paeoniflorin	C ₃₀ H ₃₂ O ₁₂	0.7	16.82	583.19898/ 629.188	B	134.0371,121.0294
83	4,7-dihydroxy-3-butylphthalide	C ₁₂ H ₁₄ O ₄	0.6	17.17	221.0821/-	C	290.0673,205.0884,161.0975
84	Senkyunolide G	C ₁₂ H ₁₆ O ₃	-0.6	17.24	207.1025/-	C	265.0631,221.8433,163.1134
40	4-hydroxy-3-butylphthalide	C ₁₂ H ₁₄ O ₃	-0.8	17.38	205.0869/-	C	237.0929,134.0374
85	senkyunolide E	C ₁₂ H ₁₂ O ₃	-0.5	17.73	203.0713/-	C	203.07211,145.0288,132.0230
48	4,5-dihydro-3-butylphthalide	C ₁₂ H ₁₆ O ₂	-4.7	17.91	191.1069/-	C	925.4829,701.4286,447.1355
86	Myristic acid	C ₁₄ H ₂₈ O ₂	-0.2	22.43	227.2016/-	H	285.1609,227.2017
87	monopalmitin	C ₁₉ H ₃₈ O ₄	-1.6	22.97	-/375.2746	H,C	429.1942,99.9258
56	Daucosterol	C ₃₅ H ₆₀ O ₆	0.2	26.31	-/621.4373	S	255.2330,96.9601

^aH, T, D, C, B, S indicate *Prunus persica* (L.) Batsch, *Carthamus tinctorius* L., *Angelica sinensis* (Oliv.) Diels, *Conioselinum anthriscoides* 'Chuanxiong' (syn. *Ligusticum chuanxiong* Hort), *Paeoniae lactiflora* Pall., and *Rehmannia glutinosa* (Gaertn.) DC, respectively.

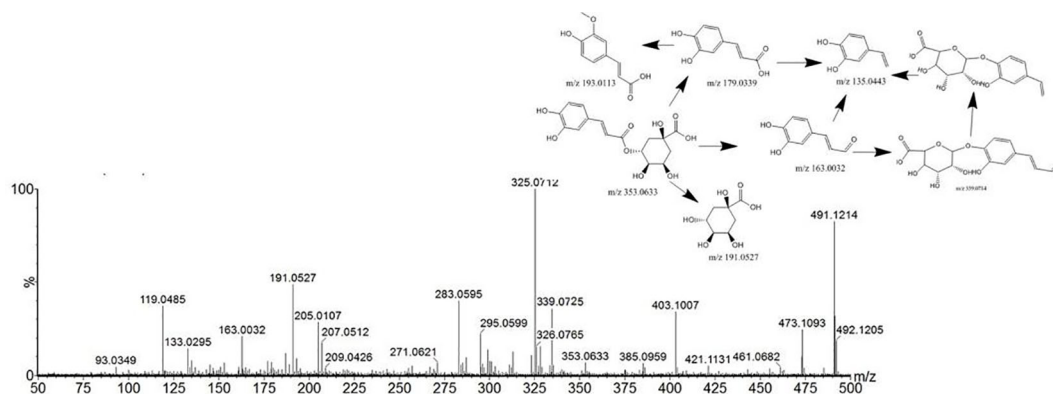


FIGURE 4 | Tandem mass spectra and possible fragment pathways of chlorogenic acid in positive ion mode.

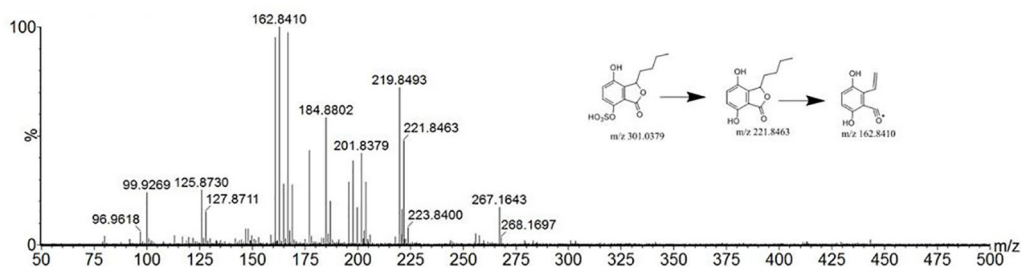


FIGURE 5 | Tandem mass spectra and possible fragment pathways of 4-hydroxy-3-butyl-benzoquinone in positive ion mode.

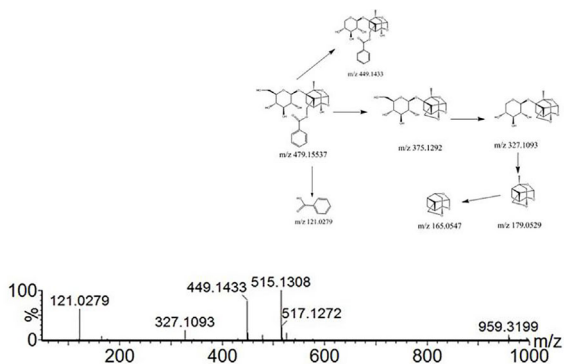


FIGURE 6 | Tandem mass spectra and possible fragment pathways of paeoniflorin in positive ion mode.

Polysaccharides

Polysaccharides are one of the main active ingredients of THSWD and are mainly from *Angelica sinensis* (Oliv.) Diels and *Rehmannia glutinosa* (Gaertn.) DC. Compound 4 had a retention time of 1.98 min, the excimer ion peak was m/z 493

$[M-H]^-$, the fragment m/z 331 was formed as an excimer ion peak, and a glucose residue (162 u) was formed. The characteristic fragment was m/z 169. The ions were derived from galloyl anions, and fragment ions of m/z 125 were generated after the gallic acyl anion underwent hydroxyl alpha cleavage to lose one molecule of CO_2 (44 u). According to the literature, the three isomers 1'-O-galloyl sucrose, 6'-O-galloyl sucrose, and 6-O-galloyl sucrose exist in the white peony, but from the total ion chromatogram A m/z 493 was extracted, and only one peak appeared at 4.549 min. Therefore, it is presumed that compound 4 is galloyl sucrose, but the linking position of its sugar needs to be confirmed.

Volatile Compounds

The major volatile oils were from *Angelica sinensis* (Oliv.) Diels and *Conioselinum anthriscoides* 'Chuanxiong' (syn. *Ligusticum chuanxiong* Hort). Compound 47 showed an $[M+H]^+$ ion at m/z 191.1061. The literature (Wang and Han, 2011) has reported that the relative molecular mass of ligustilide in *C. anthriscoides* 'Chuanxiong' was 190. It was speculated that m/z 191 was its quasi-molecular ion peak. MS² analysis was performed on m/z 191, which yielded fragment peaks such as m/z 670, 447, and 355. It showed a retention time of 17.22 in positive ion mode. According to the elemental composition analysis, the

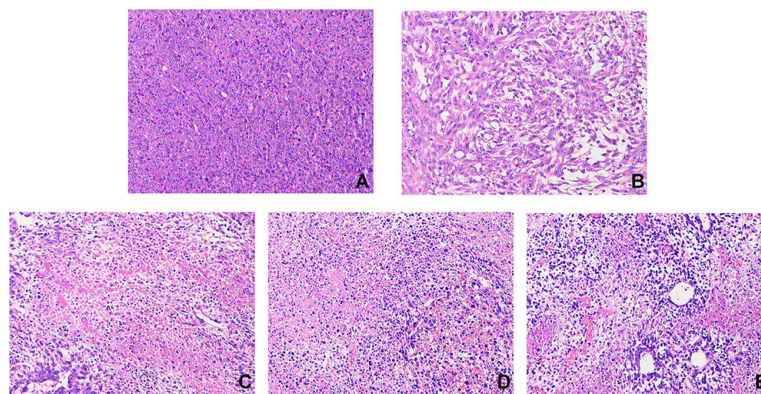


FIGURE 7 | Effect of THSWD on tumor cells in mice. Sections were stained with H&E. Magnification, $\times 200$; scale bars, 50 μm . Model group (A), cisplatin group (B), THSWD H group (C), THSWD M group (D), THSWD L group (E).

TABLE 3 | IL-4, IL-10, IL-13, and TGF- $\beta 1$ expression.

Group	IL-4(pg/ml)	IL-10(pg/ml)	TNF-a	TGF- $\beta 1$ (ng/ml)
N	366.21 \pm 38.07	1097.42 \pm 106.03	1154.64 \pm 174.62	402.80 \pm 31.10
M	336.18 \pm 51.74 ^{##}	1028.10 \pm 43.31 [#]	1657.05 \pm 268.54 ^{##}	445.98 \pm 26.76 [#]
C	506.78 \pm 64.39 ^{**}	1121.88 \pm 105.69 [*]	1338.12 \pm 56.04 [*]	413.13 \pm 11.27 [*]
H	354.10 \pm 44.88	1112.76 \pm 102.59 ^{**}	1336.89 \pm 344.15 [*]	436.98 \pm 34.06
M	386.41 \pm 45.83 ^{**}	1175.52 \pm 115.75 [*]	1607.49 \pm 170.73	400.86 \pm 40.15 [*]
L	343.69 \pm 40.53	1048.27 \pm 143.15	1335.96 \pm 296.74 [*]	424.23 \pm 13.06

Data are presented as mean \pm SD. $n = 8$ per group. ^{**} $P < 0.01$ vs. control group, ^{##} $P < 0.01$ vs. model. H, high-dose; M, middle-dose; L, low-dose; N, normal; M, model.

molecular formula was $\text{C}_{12}\text{H}_{14}\text{O}_2$, the theoretical value of the relative molecular mass was 191.1061, and the measured value was 191.1061. Based on this, it was assumed that the compound was ligustilide.

Compound 40 showed a retention time of 13.15 min in positive ion mode, with an $[\text{M}+\text{H}]^+$ ion at m/z 207.1014. The peak with t_R of 17.38 min yielded an m/z 205.0869 ion peak in the ESI⁺. The literature (Yang et al., 2007) reports that the relative molecular mass of 4-hydroxy-3-butyl- benzoquinone in *C. anthriscoides* 'Chuanxiong' is 206. According to the MS² results, it was assumed that the compound was 4-hydroxy-3-butyl-benzoquinone. The MS² spectrum and possible fragmentation pathways are depicted in Figure 5.

Monoterpene Glycosides

Compound 21 showed a retention time of 10.28 min. The fragment ions in the mass spectrum were m/z 519.1216, 503.1522, 500.1425, 179.0697, 151.0750, and 133.0644, and the excimer ion peak was m/z 479 $[\text{M}-\text{H}]^-$. The MS² spectrum and possible fragmentation pathways are depicted in Figure 6. Fragment ion m/z 525 formed for the molecular ion peak plus

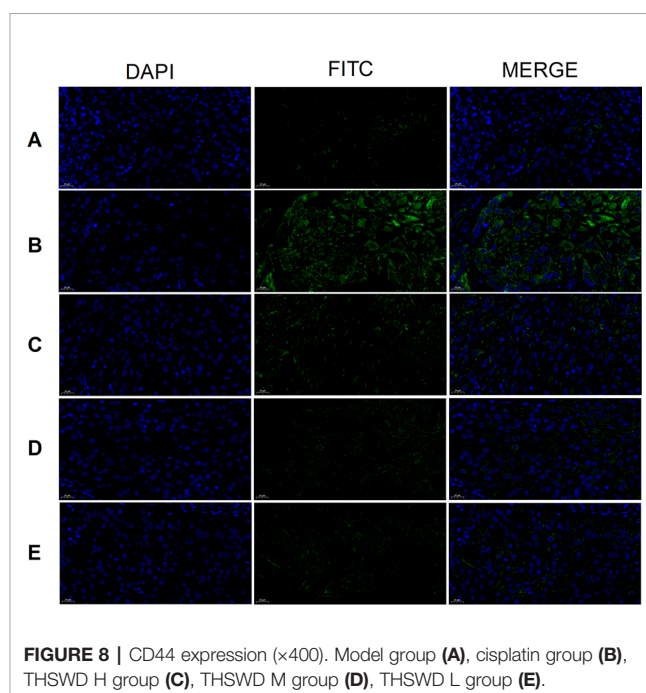
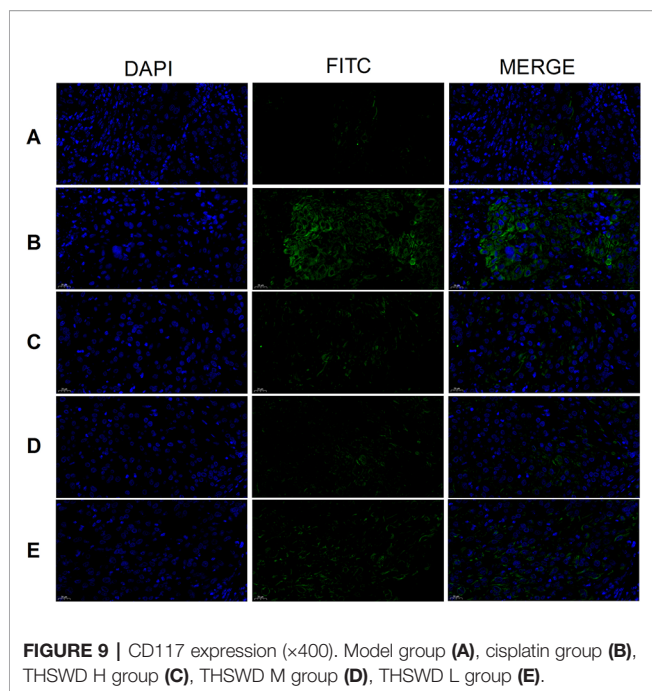


FIGURE 8 | CD44 expression ($\times 400$). Model group (A), cisplatin group (B), THSWD H group (C), THSWD M group (D), THSWD L group (E).

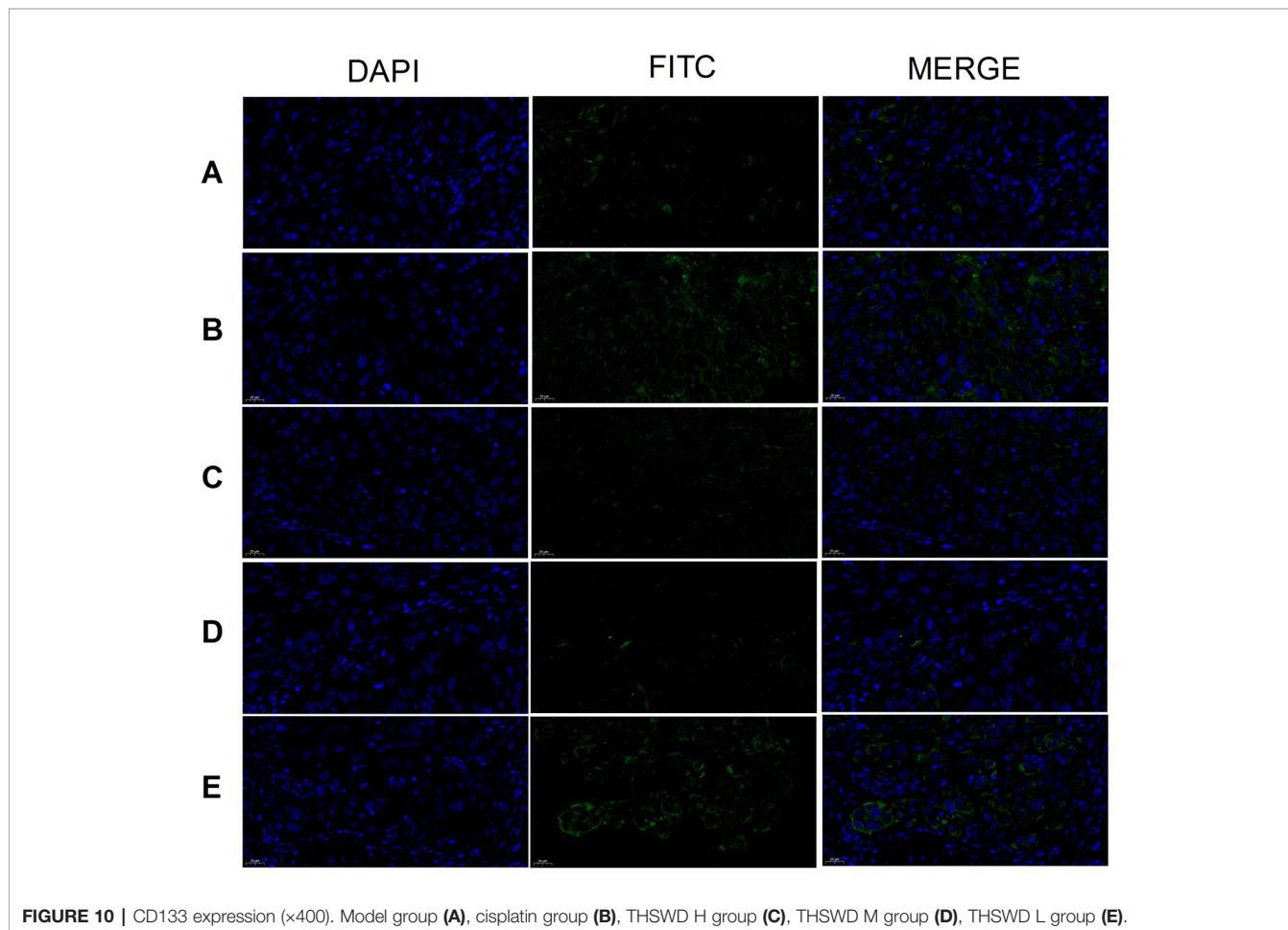
one molecule of formic acid (46 u). Due to the presence of a lactone ring in the structure of paeoniflorin, its acyloxy groups can be broken on both sides, losing one molecule of CO_2 (44 u) at m/z 435. The fragment m/z 357 was formed as the excimer peak and lost one molecule of benzoic acid (122 u), and m/z 121 was the missing one of the benzoic acid fragment ions. A combination of fragment m/z 481 $[\text{M}+\text{H}]^+$ and fragment m/z 503.1522 $[\text{M}+\text{Na}]^+$ appeared in positive ion mode, indicating that compound 21 is paeonolide B.

For compound 26, the retention time was 12.393 min. The fragment ions in the mass spectrum had m/z 525, 449, 327, and 165. The molecular ion peak was m/z 479 $[\text{M}+\text{H}]^+$, and the



fragment ion m/z 525 was the molecular ion peak plus a molecule of formic acid (46 u). m/z 449 is the molecular ion peak (m/z 479 [MH]⁻) that loses a molecule of formaldehyde (30 u) on the 6-carbon residue of the glucosyl group, while the m/z 449 fragment ion further loses a molecule of benzoic acid (122 u) to form the m/z 327 fragment ion. m/z 165 is a fragment of the decyl alkyl skeleton structure, and since the decane skeleton is connected with a benzoyl substituent, m/z 121 fragments were generated here. Combining the fragments m/z 503 [M + Na]⁺ appearing in positive ion mode, the literature suggests that compound 26 is peoniflorin.

Compounds 43 and 44 showed the same molecular weight at 584 Da and almost consistent tandem MS behavior with paeonolide. The fragment ions in the mass spectrum were m/z 629, 553, and 431. The fragment ion m/z 629 was the molecular ion peak (m/z 583 [M-H]⁻) plus one molecule of formic acid (46 u), and m/z 553 was the molecule. The ion peak (m/z 583 [M-H]⁻) loses the formation of a molecule of formaldehyde (30 u) on the 6-carbon residue of the glucosyl group, and the fragment ion m/z 553 further loses a molecule of benzoic acid (122 u), resulting in m/z 431 fragment ions. Positive ion mode yielded m/z 585 [M + H]⁺ and m/z 607 [M + Na]⁺, which identified compound 44 as benzoyl glycoside according to the literature.



Aromatic Cyanogenic Glycosides

Most of the common cyanogenic glycosides have the same skeleton structure. According to the substituents on the cyanohydrin derivatives, they can be divided into two categories: aliphatic and aromatic. A given species of plant generally contains only one or two cyanogenic glycosides, such as Rosaceae. Compounds 14 and 15 were all derived from dry mature seeds of the Rosaceae plant *Prunus persica* (L.) Batsch or *Prunus davidiana* (Carr.) Franch.

Others

THSWD also contains human essential amino acids, vitamins, and other ingredients.

H&E

H&E stained samples for each group are shown in **Figure 7**. Tumor cells in the model group were arranged closely, with less necrosis and a large nuclear-to-plasma ratio. The cell morphology was abnormal, and there was more vascular filling. Compared with the model group, tumor cells in the THSWD groups and cisplatin group showed large areas of necrosis with obvious cell debris. The necrotic cells were loosely arranged with large gaps. The results indicate that THSWD exerts a destructive effect on mouse breast cancer cells. This is the first evidence that THSWD has a significant therapeutic effect on breast cancer tumor cells.

IL-4, IL-10, IL-13, and TGF- β 1 Expression

To explore the mechanism of THSWD on breast cancer, levels of IL-6, IL-8, and IL-1 β were measured by ELISA (**Table 3**). IL-5 was significantly increased in the model group. After treatment with THSWD or cisplatin, IL-5 was significantly reduced. The results indicate that THSWD may regulate immune function in breast cancer mice.

CD44, CD117, and CD133 Expression

Tumor cell CD44 expression was highest in the model group and lowest in the cisplatin group (**Figure 8**). There was no obvious difference between the three THSWD dose groups. The results showed that cisplatin and THSWD significantly inhibited CD44 expression in tumor cells of breast cancer mice.

CD117 expression was highest in the tumor cells of the model group and lowest in the cisplatin and THSWD middle-dose groups (**Figure 9**). The results showed that cisplatin and THSWD significantly inhibited CD117 expression in tumor cells of breast cancer mice.

The expression of CD133 was highest in tumor cells of the model group and lowest in the cisplatin group (**Figure 10**). Results were similar for the THSWD high-, middle-, and low-dose groups. Both cisplatin and THSWD significantly inhibited CD133 expression in tumor cells of breast cancer mice.

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CONCLUSION

In this study, UPLC-Q-TOF-MS was used for the full chemical characterization of THSWD in both positive and negative ion modes. A total of 87 compounds were identified in 32 minutes, including aromatic acids, flavones, polysaccharides, volatile oils, monoterpene glycosides, aromatic cyanogenic glycosides, and others. This method and the chemical material findings may provide useful information for quality control when producing THSWD and related individual herbs, despite the ingredient complexity and specificity for TCM. The results can be used to develop a quality standard of the TCM compound THSWD and provide a theoretical basis for further study of THSWD metabolism *in vivo* to clarify the material basis and mechanism of its clinical application.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation, to any qualified researcher.

ETHICS STATEMENT

All experiments were subject to approval by the Committee on the Ethics of Animal Experiments of Anhui University of Chinese Medicine.

AUTHOR CONTRIBUTIONS

XD is responsible for designing the experiment private. LP is responsible for article writing. QB is responsible for the experimental operation. DP is responsible for data analysis.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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