Analysis of *E. rutaecarpa* Alkaloids Constituents *In Vitro* and *In Vivo* by UPLC-Q-TOF-MS Combined with Diagnostic Fragment

Shenshen Yang, Meng Tian, Lei Yuan, Haoyue Deng, Lei Wang, Aizhu Li, Zhiguo Hou, Yubo Li, and Yanjun Zhang

1. School of Traditional Chinese Materia Medica, Tianjin University of Traditional Chinese Medicine, 312 Anshan West Road, Tianjin 300193, China
2. Tianjin State Key Laboratory of Modern Chinese Medicine, Tianjin University of Traditional Chinese Medicine, 312 Anshan West Road, Tianjin 300193, China

Correspondence should be addressed to Yubo Li; yubolil163.com and Yanjun Zhang; tianjin_tcm001@sina.com

Received 22 December 2015; Revised 3 May 2016; Accepted 5 May 2016

Academic Editor: Filomena Conforti

Copyright © 2016 Shenshen Yang et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

*Evodia rutaecarpa* (Juss.) Benth. (Rutaceae) dried ripe fruit is used for dispelling colds, soothing liver, and analgesia [1]. *E. rutaecarpa* can ease depression, relieve dyspepsia, and exert analgesic, sedative, antibacterial, and antioxidant activities, as well as other pharmacological effects. Up to now, alkaloids are the main active ingredients of *E. rutaecarpa* [2]. However, previous studies have focused on the indole biological components of *E. rutaecarpa* [3, 4]. TCM contains many components, but only the ingredients absorbed into the bloodstream produce an effect [5]. Therefore, the ingredients of TCM found in the animal blood should be analyzed using a vast majority of the drugs administered to animals to determine the active sites of actual and effective ingredients.

The chemical constituents and absorbed components of *E. rutaecarpa* should be analyzed to completely control the medicinal qualities of this plant and simultaneously detect its main ingredients.

Previous studies have employed TLC, HPLC with UV detector or MS, and CE to quantify or identify the alkaloids in Evodia [6–10]. An effective and reliable analytical method should be established to quickly classify and identify *E. rutaecarpa* chemical compositions *in vitro* and *in vivo*. Ultraprecision liquid chromatography quadrupole time-of-flight mass spectrometry (UPLC-Q-TOF-MS) is advantageous because of its high speed, high resolution, and high accuracy. MS scanning can provide accurate mass information on characteristic molecular ions and fragment ions to offer a reliable basis for the qualitative and quantitative analyses of target compounds.
compounds [11–13]. UPLC-Q-TOF-MS is also increasingly applied in many areas, which analyzes sterol glycosides and uncovers the effects of light intensity and temperature under shading treatments on the metabolites in tea [14, 15]. Diagnostic fragment was the characteristic fragment of a certain type of compound applied for screening and identification. Based on these characteristics, we also employed UPLC-Q-TOF-MS in our experiment to quickly classify and identify the alkaloids compositions of *E. rutaecarpa*. Additionally, we detected the absorbed components after intragastric administration by *E. rutaecarpa* alcohol extract and observed the exogenous active substances.

The above screening method can quickly classify and identify *E. rutaecarpa* chemical compositions. The proposed method provided a fast, accurate, and feasible analytical process to distribute and determine the complex components in traditional Chinese medicine (TCM). First, we used UPLC-Q-TOF-MS to perform a fingerprint analysis of *E. rutaecarpa* and obtain the fragments’ information of *E. rutaecarpa* chemical constituents. Afterward, we rapidly classified and identified the *E. rutaecarpa* alkaloids compositions by using diagnostic fragments as screening tools. Finally, combined with the literature and standard we could confirm the compounds in the plant. According to this method, a total of 17 alkaloids constituents and 6 absorbed components were identified from *E. rutaecarpa*. The established method is expected to be widely accepted and approved, given the popularity of MS and the requirements of fast and efficient assays in routine studies.

### 2. Materials and Methods

#### 2.1. Materials

The UPLC-Q-TOF-MS system (Waters, USA) consisted of an autosampler and a DAD detector and a column compartment was used for the analysis. Acetonitrile, formic acid, and methanol were chromatographically pure. The 95% alcohol used was also analytically pure and *E. rutaecarpa* (Juss.) Benth. adopted in this experiment was purchased from Hubei Herbs Company. A total of 10 male Wistar rats weighing 180 ± 20 g were obtained from the Academy of Military Medical Sciences.

#### 2.2. UPLC-Q-TOF-MS Analysis Conditions

The chromatographic column was used was Waters ACQUITY UPLC BEH C18 Column (2.1 × 100 mm, 1.7 μm). Column temperature was set to 35°C. Phase A was the water phase, which consisted of 0.1% aqueous solution of formic acid. Phase B consisted of 0.1% formic acid in acetonitrile solution at a flow rate of 0.3 mL/min and injection volume of 5 μL. Gradient elution was used for chromatographic separation and the gradient programs were 0–1 min, 11% B; 1–2 min, 11%–21% B; 2–4 min, 21%–33% B; 4–7 min, 33%–70% B; 7–9 min, 70%–82% B; 9–16 min, 82%–100% B; 16–18 min, 100% B; 18–19 min, 100%–11% B; and 19–20 min, 11% B.

UPLC-Q-TOF-MS equipped with ESI in positive ion mode was used to scan and analyze the chemical constituents of *E. rutaecarpa*. MS parameters were set as follows: dry gas (N2) flow rate, 10 mL/min; gas temperature, 325°C; desolvation gas flow rate, 600 L/h; capillary voltage, 2.1 kV; and collision-induced dissociation voltage, 6 kV. For the nebulizer, the boil-off gas was at 350 psi of pressure and the auxiliary gas was high-purity nitrogen. The reference ion was [M+H]+ = 5652771 to ensure the acquisition accuracy in spectra. Data were acquired in the range of 50–1000 Da.

#### 2.3. Solution Preparation

*E. rutaecarpa* powder (40 mesh) was extracted twice by refluxing extraction. The first extraction was performed 10 times with 70% ethanol extract for 1.5 h and the second extraction was 8 times of 70% ethanol extract for 1.5 h. After filtration, the mixed filtrate was concentrated to a relative density of 0.1 g crude drug/mL by recovering ethanol, and the mixture was stored for further use.

#### 2.4. Blood Samples Preparation

Ten Wistar rats (180 ± 20 g) were fasted with free access to water for 12 h prior to the experiment. The rats were then intragastrically given *E. rutaecarpa* ethanol extract (0.6 g crude drug/kg body weight). After 2 h, blood samples were collected from the eye venous plexus. The samples were centrifuged at 3000 r/min for 15 min and 3500 r/min for 8 min to obtain the serum. The serum was immediately placed in a refrigerator at −80°C and stored until analysis. 200 μL of the serum was precisely pipetted and placed in a 5 mL stopped centrifuge tube. An equivalent amount of acetonitrile was added to the tube for protein precipitation. After stirring for 3 min, the tube was centrifuged at 4°C at 13000 r/min for 15 min. Supernatant was filtered through a 0.45 μm membrane for UPLC-Q-TOF-MS analysis in accordance with chromatographic conditions.

### 3. Results and Discussion

#### 3.1. Established Methods of Compound Classification and Identification

TCM involves many plants with complex chemical compositions. However, substances with similar frameworks are placed in the same category, and these substances exhibit similar material fracture behavior during collision-induced mass spectrometry. Therefore, we used this feature to explore for fragmentation patterns of a substance in its mass spectrum [16]. Based on the structural characteristics of the two alkaloids and the information collected from a large number of literatures, and compared with the standard of mass spectrometry, the rule of the diagnosis of these two kinds of alkaloids was found and summarized. Therefore we established the method which combined diagnose fragments with UPLC-Q-TOF-MS to characterize the chemical compositions of *E. rutaecarpa*. As previously mentioned, alkaloids are the main active ingredients of this herb. The alkaloids in *E. rutaecarpa* are mainly indole alkaloids and quinolone alkaloids. Figure 1 shows the typical total ion current (TIC) chromatograms of the results obtained via UPLC-Q-TOF-MS of this herb and blood.

We quickly identified the composition of *E. rutaecarpa* and speculated the possible structures of its compounds through their mass spectral fragmentation patterns and by combining the data reported in the literature (Table 1). A total of 5 indole alkaloids and 12 quinolone alkaloids were identified on the basis of the contrasting cleavage rules, fragment ion characteristics, and mass spectral data. In the
3.2. Identification of the Compounds in *E. rutaecarpa* Herb

3.2.1. Indole Alkaloid. We divided these alkaloids into two categories based on their structural characteristics. Category A contained one indole quinoline alkaloid with five complete rings. Category B contained ring openings instead of five ring structures in their parent nucleus.

According to the literature and reference mass spectrum information of standard available, the significant structural feature of category A is the transformation of the cyclohexene. Under the collision of MS, the characteristic pathway of category B is the transformation of the additional double bonds. In addition, the characteristic peaks of category A are the transformation of the cyclohexene rings. Category B contained ring openings instead of five ring structures in their parent nucleus.

The mass spectrum of compound 3 (Figure 2) showed the retention time at 6.97 min and estimated formula was \( \text{C}_{19}\text{H}_{17}\text{N}_2\text{O} \). The mass charge ratios of the main fragments were 304, 171, 161, 144, 134, 116, and 106; \( m/z \) 304 was the quasimolecular ion peak \([M+H]^+\). We get the fragment ion at \( m/z \) 171 \([M+\text{H-C}_8\text{H}_8\text{NO}]^+\) and \( m/z \) 134 \([M+\text{H-C}_{11}\text{H}_{10}\text{N}_2]^+\) ion molecule of complementary fragment ions which is the typical RDA fragmentation characteristics, so determining that it belongs to category A firstly. Furthermore, fragment ions at \( m/z \) 161 \([M+\text{H-C}_{10}\text{H}_8\text{N}]^+\) and 144 \([M+\text{H-C}_9\text{H}_7\text{N}_2\text{O}]^+\) were found because another RDA fragmentation occurred, which resulted from the rearrangement of intramolecular double bonds. In addition, the \( m/z \) 144 \([M+\text{H-C}_7\text{H}_5\text{N}_2\text{O}]^+\) and 134 \([M+\text{H-C}_7\text{H}_5\text{N}_2]^+\) of debris were missing a -C_2H_4 and -CO formed \( m/z \) 116 \([M+\text{H-C}_9\text{H}_8\text{N}_2\text{O-C}_2\text{H}_4]^+\) and 106 \([M+\text{H-C}_{11}\text{H}_{10}\text{N}_2\text{CO}]^+\) of the fragment ion. Hence, compound 3 was identified as evodiamine, which was consistent with the literature and standard [19]. At the same time, these alkaloids also contain rutecarpine, 14-formylidihydrorutecarpine, and dehydroevodiamine.

Category B showed significant structural differences from category A. RDA fragmentation did not occur in category B. These characteristics were used to distinguish the two categories. Consider that the cleavage active sites were amide and quaternary ammonium bonds. From the structures of category B, the easily broken sites were the C-N bonds in the amide groups. Other bonds linked to quaternary ammonium nitrogen atoms were also easy to break.

The mass spectrum of compound 5 (Figure 3) showed the retention time at 6.81 min; estimated formula was \( \text{C}_{19}\text{H}_{17}\text{N}_2\text{O} \). The mass charge ratios of the main fragments were 308, 175, 165, and 134, which is not the typical RDA fragmentation characteristics, so determining that it belongs to category B firstly. Among them, \( m/z \) 308 was the quasi-molecular ion peak \([M+H]^+\). \( m/z \) 134 \([M+\text{H-C}_{11}\text{H}_{10}\text{N}_2]^+\) and 175 \([M+\text{H-C}_9\text{H}_7\text{N}_2\text{NO}]^+\) were produced by amide structure fragmentation, which was the main fragmentation pathway of category B. Furthermore, fragment ions at \( m/z \) 165 \([M+\text{H-C}_{10}\text{H}_9\text{N}]^+\) and 144 \([M+\text{H-C}_9\text{H}_7\text{N}_2\text{O}]^+\) were found resulting from the transformation of the cyclohexene.
<table>
<thead>
<tr>
<th>Number</th>
<th>RT</th>
<th>Identification</th>
<th>Formula</th>
<th>Theoretical mass (m/z)</th>
<th>Experimental mass (m/z)</th>
<th>Error (ppm)</th>
<th>Blood components</th>
<th>MS data (+) (m/z)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>3.07</td>
<td>14-Formyldihydroxyrutaecarpine</td>
<td>C_{19}H_{15}N_{3}O_{2}</td>
<td>318.1243</td>
<td>318.1245</td>
<td>0.63</td>
<td>Y</td>
<td>273, 261, 217, 171, 144</td>
</tr>
<tr>
<td>2</td>
<td>4.29</td>
<td>Dehydroevodiamine</td>
<td>C_{19}H_{15}N_{3}O</td>
<td>302.1295</td>
<td>302.1309</td>
<td>4.63</td>
<td>Y</td>
<td>302, 286, 261, 217, 177, 133</td>
</tr>
<tr>
<td>3</td>
<td>6.97</td>
<td>Evodiamine</td>
<td>C_{19}H_{17}N_{3}O</td>
<td>304.1450</td>
<td>304.1439</td>
<td>−3.62</td>
<td>Y</td>
<td>304, 171, 161, 144, 116, 106</td>
</tr>
<tr>
<td>4</td>
<td>7.13</td>
<td>Rutaecarpine</td>
<td>C_{18}H_{13}N_{3}O</td>
<td>288.1137</td>
<td>288.1141</td>
<td>1.39</td>
<td>Y</td>
<td>288, 273, 171, 169, 120</td>
</tr>
<tr>
<td>5</td>
<td>6.81</td>
<td>Evodiamide</td>
<td>C_{19}H_{21}N_{3}O</td>
<td>308.1763</td>
<td>308.1753</td>
<td>−3.24</td>
<td>N</td>
<td>308, 175, 165, 144, 134</td>
</tr>
<tr>
<td>6</td>
<td>7.87</td>
<td>1-Methyl-2-nonyl-4(1H)-quinolone</td>
<td>C_{19}H_{19}NO</td>
<td>286.2171</td>
<td>286.2171</td>
<td>0</td>
<td>N</td>
<td>286, 261, 217, 186, 173, 144, 102</td>
</tr>
<tr>
<td>7</td>
<td>8.20</td>
<td>1-Methyl-2-[(Z)-6-undecenyl]-4(1H)-quinolone/1-methyl-2-[(Z)-5-undecenyl]-4(1H)-quinolone</td>
<td>C_{21}H_{29}NO</td>
<td>312.2327</td>
<td>312.2334</td>
<td>2.24</td>
<td>N</td>
<td>312, 217, 186, 173, 144</td>
</tr>
<tr>
<td>8</td>
<td>8.31</td>
<td>1-Methyl-2-[(Z)-6-undecenyl]-4(1H)-quinolone/1-methyl-2-[(Z)-5-undecenyl]-4(1H)-quinolone</td>
<td>C_{21}H_{29}NO</td>
<td>312.2327</td>
<td>312.2321</td>
<td>−1.92</td>
<td>N</td>
<td>312, 217, 186, 173, 144</td>
</tr>
<tr>
<td>9</td>
<td>8.69</td>
<td>1-Methyl-2-[(4Z,7Z)-4,7-tridecadienyl]-4(1H)-quinolone</td>
<td>C_{23}H_{31}NO</td>
<td>338.2484</td>
<td>338.2480</td>
<td>−1.18</td>
<td>N</td>
<td>338, 217, 186, 173, 144</td>
</tr>
<tr>
<td>10</td>
<td>8.95</td>
<td>1-Methyl-2-undecyl-4(1H)-quinolone</td>
<td>C_{21}H_{21}NO</td>
<td>314.2484</td>
<td>314.2484</td>
<td>0</td>
<td>N</td>
<td>314, 217, 186, 173, 144</td>
</tr>
<tr>
<td>11</td>
<td>9.24</td>
<td>Evocarpine</td>
<td>C_{23}H_{33}NO</td>
<td>340.2640</td>
<td>340.2642</td>
<td>0.59</td>
<td>Y</td>
<td>340, 187, 174, 173, 159, 144, 132</td>
</tr>
<tr>
<td>12</td>
<td>9.58</td>
<td>1-Methyl-2-dodecyl-4(1H)-quinolone</td>
<td>C_{22}H_{33}NO</td>
<td>328.2640</td>
<td>328.2633</td>
<td>−2.13</td>
<td>N</td>
<td>328, 186, 173, 144, 132</td>
</tr>
<tr>
<td>13</td>
<td>9.63</td>
<td>1-Methyl-2-[(6Z,9Z)-6,9-pentadecadienyl]-4(1H)-quinolone</td>
<td>C_{25}H_{35}NO</td>
<td>366.2797</td>
<td>366.2814</td>
<td>4.64</td>
<td>N</td>
<td>354, 328, 305, 261, 217, 186, 173, 144, 132</td>
</tr>
<tr>
<td>14</td>
<td>10.25</td>
<td>Dihydroevocarpine</td>
<td>C_{23}H_{33}NO</td>
<td>342.2797</td>
<td>342.2805</td>
<td>2.34</td>
<td>Y</td>
<td>342, 340, 261, 217, 186, 173, 144, 122</td>
</tr>
<tr>
<td>15</td>
<td>10.47</td>
<td>1-Methyl-2-[(Z)-9-pentadecenyl]-4(1H)-quinolone/1-methyl-2-[(Z)-10-pentadecenyl]-4(1H)-quinolone</td>
<td>C_{25}H_{35}NO</td>
<td>368.2953</td>
<td>368.2942</td>
<td>−2.99</td>
<td>N</td>
<td>368, 202, 186, 173, 144, 122</td>
</tr>
<tr>
<td>16</td>
<td>10.54</td>
<td>1-Methyl-2-[(Z)-9-pentadecenyl]-4(1H)-quinolone/1-methyl-2-[(Z)-10-pentadecenyl]-4(1H)-quinolone</td>
<td>C_{25}H_{35}NO</td>
<td>368.2953</td>
<td>368.2949</td>
<td>−1.09</td>
<td>N</td>
<td>368, 342, 217, 186, 173, 144, 122</td>
</tr>
<tr>
<td>17</td>
<td>11.83</td>
<td>1-Methyl-2-pentadecyl-4(1H)-quinolone</td>
<td>C_{25}H_{39}NO</td>
<td>370.3100</td>
<td>370.3105</td>
<td>−1.35</td>
<td>N</td>
<td>370, 261, 217, 186, 173, 144, 122</td>
</tr>
</tbody>
</table>

(a) Y represents detected in the blood and N denotes not detect.
in the quaternary amine bond cleavage. Therefore, compound 5 was identified as evodiamide by comparing the mass spectral data with the related literature [20].

3.2.2. Quinolone Alkaloids. The mother nuclei of quinolone alkaloids were identical, and they only differed in the position of C-2 in the side chains. According to the structure and literature known that quinolone alkaloids can occur replacement rearrangement and McLafferty rearrangement lose part of side chain formation with a stable total conjugate system of fragment ions \( m/z \) 186 and \( m/z \) 173. Therefore, \( m/z \) 186 and 173 ion peaks were used as diagnostic fragments to evaluate quinolone alkaloids. Except for their identical mother nuclei, these compounds only differ in the position of the C-2 of side chains, where their carbon chain lengths and unsaturation degrees are distinct. However, only the fragment ions of carbon chain were difficult to detect, and the characteristic fragment ions of certain compounds were consequently absent in the mass spectral detection of quinolone alkaloids [21]. Thus, we documented the molecular ion peak \([M+H]^+\) for authentication and identified quinolone alkaloid chemical constituents combined with information from the literature.
and mass spectral data. Hence, we preliminarily determined 12 quinolone alkaloids.

Compound 11, for example, with a retention time at 9.24 min and inference formula, may be C_{23}H_{33}NO. In the experiment get m/z 340, 328, 187, 186, 174, 173, 159, 144, and 132 fragment ions. Firstly, the mother nuclear fragment ions with high abundance at m/z 186 [C_{12}H_{18}NO]^{+} and 173 [C_{11}H_{17}NO]^{+} were found to be unrelated to the side chains, which can be diagnostic compounds 11 belonging to quinolone alkaloids. Secondly, we also received by the m/z 144 and 159 of debris. The mass difference between m/z 186 [C_{23}H_{33}NO]^{+} and the fragment at m/z 144 [C_{11}H_{22}NO-C_{2}H_{2}O]^{+} indicated a lost ketene molecule. A fragment ion at m/z 159 [C_{11}H_{17}NO-CH_{2}]^{+} was formed because a molecule of -CH_{2} was removed from the fragment at m/z 173 [C_{11}H_{16}NO]^{+}. In addition, the mass spectrum of compound 11 showed a quasimolecular ion peak at m/z 340 [M+H]^{+}; therefore, compound 11 was identified as evocarpine by comparing the mass spectral data with the standard. The major fragment ions of evocarpine are shown in Figure 4.

From the mass spectra of compounds 6–17, the mother nuclear fragment ions with high abundance at m/z 186 and 173 were found to be unrelated to the side chains. Thus, compounds 6–17 were quinolone alkaloids.

3.3. Identification of the Compounds in Blood. In this study, 6 alkaloids were identified in the blood. The 6 active components were dehydroevodiamine, evodiamine, rutaecarpine, and 14-formylidihydroxyrutaecarpine of indole alkaloids and evocarpine and dihydroevocarpine of the quinolone alkaloids. Through consulting the literature known, evodiamine and rutaecarpine in human serum via LC–MS by analyzing the blood components in E. rutaecarpa [22].

Compound 2 as an Example. Compound 2, a known compound in E. rutaecarpa, is dehydroevodiamine; the retention time is at 4.29 min, and estimated formula was C_{18}H_{15}N_{2}O and the m/z of [M+H]^{+} is 302.1309. Firstly, screening for the data of blood samples using MassLynx software, the results of screening are that the m/z of [M+H]^{+} is 302.1296 and retention time is at 4.38 min; based on the retention time and the quasimolecular ion peak we can initially determine which is dehydroevodiamine. The screening result was shown in Figure 5. Then, the main fragments of screening compound were compared with the fragments of compound 2. The mass charge ratios of the main fragments of compound 2 were 302, 286, 261, 217, 177, and 133, and the mass charge ratios of the main fragments in screen compound were 302, 261, 217, 177, and 133. The debris information is basically consistent with the fragment information of compound 2. Finally, comparing the main fragments of screen compound with the standard, the mass charge ratios of the main fragments of dehydroevodiamine were 302, 287, 261, 217, 210, 177, and 133, which were consistent with the compound in blood samples, so determined the compound that the m/z of [M+H]^{+} is 302.1296 of screening in blood samples was dehydroevodiamine. The main fragments were shown in Figures 6 and 7.

Combining the diagnostic fragments with the technology of UPLC-Q-TOF-MS and comparing the ion fragment information with that of the alkaloids in E. rutaecarpa, a total of 17 chemical components were identified. At the same time, the data of blood samples by using UPLC-Q-TOF-MS were compared with the identified chemical components in E. rutaecarpa and we classified and identified the 6 active components of E. rutaecarpa. Thus, this method can quickly and accurately identify the main components and active components of the traditional TCM.
Figure 5: The screening result of blood sample.

Figure 6: The main fragments of screening from blood.
3.4. Discussion. In this experiment, 17 compounds were identified in *E. rutaecarpa*. After animals were intragastrically given *E. rutaecarpa*, 6 absorbed components were found. The existence of other forms of metabolites remains to be further verified.

We investigated *E. rutaecarpa* constituents and blood components in positive and negative ionization modes. After we detected the alkaloids in *E. rutaecarpa*, we observed a high abundance of [M+H]⁺ ion peaks in the excimer (+) ESI mass spectra, whereas [M−H]⁻ signals nearly did not appear in the excimer (−) ESI mass spectra. Thus, mass spectrum in positive ion mode was finally obtained, which was consistent with the literature [23].

There are different methods established for analysis of the chemical composition of Evodia reported in previous studies. One is use of HPLC-ESI-MS to identify five kinds of alkaloid in Evodia [24]; another is use of ESI-IT-TOF-MS method to analyze rutaecarpine and fragmentation pathway of the two derivatives based on accurate ion mass and multilevel spectroscopy [25]. Compared with the above two documents, this study not only combined the advantages of their method, using UPLC-Q-TOF-MS combined with fragments diagnostic technology, through the diagnostic fragments for rapid compound classification, but also can more accurately identify the chemical composition sensitively, and our research identified 17 chemical composition of Evodia.

In this study, the 6 active components were dehydroevodiamine, evodiamine, rutaecarpine and 14-formylhydroxyrutaecarpine of indole alkaloids and evocarpine and dihydroevocarpine of the quinolone alkaloids. Evodia has analgesic, anti-inflammatory, antiulcer, relaxing blood vessels and lowering blood pressure, cardiac, antitumor, and other pharmacological effects. According to the literature, evodiamine from *Evodia rutaecarpa* induces apoptosis via activation of JNK and PERK in human ovarian cancer cells [26]. Evodiamine selectively targets cancer stem-like cells through the p53-p21-Rb pathway and may be used for the treatment of breast cancer [27]. Dehydroevodiamine could antagonize triggered arrhythmias such as ACs and DADs induced by cardiotonic agents in human atrial and ventricular tissues through a general reduction of the Na⁺ and Ca²⁺ inward currents, with an increase of pH, and NHE activity [28]. Therefore, the role of these active ingredients is consistent with pharmacological effects of Evodia.

The mass spectral fragment ions were produced by the single-molecule cleavage reaction of molecular ions or large fragment ions. The abundance of a specific fragment ion relative to the molecular ions and other fragment ions provided the position structures in the molecules, environment, and other valuable information on fragment ions. UPLC-Q-TOF-MS is characterized by high analysis speed, high examination sensitivity, and strong anti-interference ability, among others. This process can effectively reduce body interference and improve detection accuracy when analyzing complex samples. This process is also applied to the simultaneous determination of various complex components in TCM. Furthermore, UPLC-Q-TOF-MS is extensively used to determine the structures of compounds. This method can quickly classify compounds and plays important roles in the identification of effective materials and components of TCM.
4. Conclusion

A rapid and sensitive method was developed using UPLC-Q-TOF-MS coupled with diagnostic fragment technique to separate and identify *E. rutacearpa* chemical constituents *in vitro* and absorbed components *in vivo*. A total of 17 chemical constituents *in vitro* and 6 absorbed components *in vivo* were successfully identified in *E. rutacearpa* in terms of the precise relative molecular mass, mass spectral fragment structural information, chromatographic retention rules of the chromatographic peaks in the mass spectra, and the previous literature. UPLC-Q-TOF-MS rapidly analyzed the chemical constituents and medicinal components of TCM and provided a new concept for the in-depth study on the medicinal composition of TCM. Hence, the proposed method is a relatively good qualitative determination technique to establish an experimental basis for further analysis of *E. rutacearpa* chemical constituents, control of medicinal qualities, and development of TCM chemical constituents. Furthermore, this research on absorbed components *in vivo* is important for clinical research.

Competing Interests

The authors declare that there is no conflict of interests regarding the publication of this paper.

Authors’ Contributions

Shenshen Yang and Meng Tian contributed equally to this work and should be considered co-first authors.

Acknowledgments

Yubo Li and Yanjun Zhang participated in the conception and design of the experiments and provided critical advice on final paper. Shenshen Yang, Meng Tian, Lei Yuan, Haoyue Deng, Lei Wang, Aizhu Li, and Zhiguo Hou carried out the experiment and prepared the first draft of the paper. All authors contributed to the preparation of the paper and read and approved the final version. This work was supported by the National Science and Technology Support Program of China (2011BA107B00 and 2011BA107B08) and the Program for Changjiang Scholars and Innovative Research Team in University (IRT_14R41).

References


Submit your manuscripts at http://www.hindawi.com