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Journal of Pharmaceutical and Biomedical Analysis 31 (2003) 175–183



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Short communication

Analysis of 'SHENMAI' injection by HPLC/MS/MS

Zhang Haijiang, Wu Yongjiang, Cheng Yiyu*

Institute of Pharmaceutical Engineering, College of Materials Science and Chemical Engineering, Zhejiang University, Hangzhou 310027, PR China

Received 15 July 2002; received in revised form 27 September 2002; accepted 28 September 2002

Abstract

An HPLC/MS/MS method was developed for the analysis of 'SHENMAI' injection, composed of red ginseng and ophiopogon. The constituents of 'SHENMAI' were found to be similar with those of ginseng and 39 ginsenosides were detected. By the studies of MS and MS/MS spectra and the comparison with literature data, most of these ginsenosides were identified. Based on this study, suggestions were put forward to improve the quality control system of 'SHENMAI' injection.

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Keywords: SHENMAI injection; Red ginseng; Ginsenosides; Quality control; HPLC/MS

1. Introduction

'SHENMAI', made from red ginseng and ophiopogon, is one of the most widely used herbal medicines in Traditional Chinese Medicine. It's used to treat cardiomyopathy, miocardial infarction and hypertension. Although the main constituents in ginseng and ophiopogon were found to be ginsenosides and ophioponins, respectively [1], so far, the active compounds were not exactly clear yet. Thus, it led to the difficulty in the quality control of 'SHENMAI' injection. The present

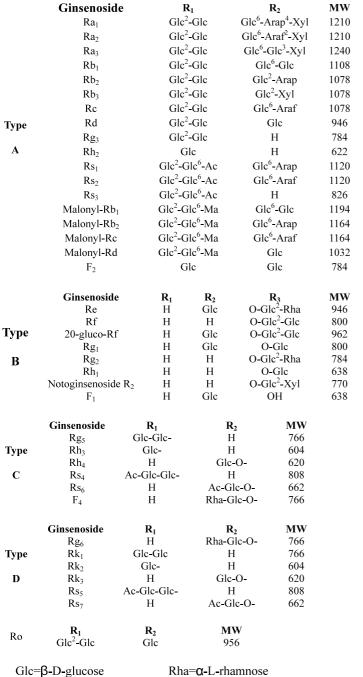
E-mail address: chengyy@zju.edu.cn (C. Yiyu).

quality control system is established on the HPLC fingerprint technology, by which a quality assessment model was built on the data analysis of HPLC chromatograms of different batches of 'SHENMAI' injections. However, due to the unknown of constituents of 'SHENMAI' and the mechanism of the activity, this model could not accurately reflect the pharmaceutical activity of 'SHENMAI'.

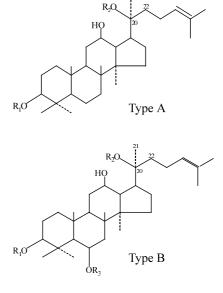
The present study aims to develop a HPLC/MS/ MS method to analyze 'SHENMAI' injection and its composition plants, i.e. ginseng and ophiopogon, to characterize the constituents of the 'SHENMAI' and its difference with ginseng and ophiopogon. This study would be potentially helpful to optimize the fingerprint model.

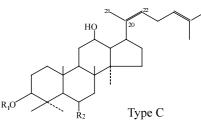
^{*} Corresponding author. Tel.: +86-571-89752509; fax: +86-571-87980668.

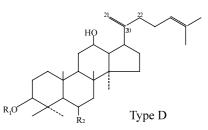
^{0731-7085/02/\$ -} see front matter © 2002 Elsevier Science B.V. All rights reserved. PII: S 0 7 3 1 - 7 0 8 5 (0 2) 0 0 5 6 5 - 4



Arap= α -L-arabinose(pyranose) Araf= α -L-arabinose(furanose) Xyl= β -D-xylose Ac=6-O-acetyl







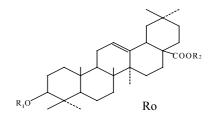


Fig. 1. Structures of ginsenosides.

Table 1Solvent gradient program of HPLC analysis

Time (min)	A (%)	B (%)
0	70	30
10	60	40
15	60	40
25	54	46
25 65 75	35	65
75	25	75

2. Experimental

2.1. Samples preparation

'SHENMAI' injection, together with the extracts of red ginseng and ophiopogon, were supplied by a Chinese pharmaceutical company. All the samples were filtered through a 0.45 μ m film before HPLC analysis.

The plant materials, Panax ginseng C.A. Mey and Ophiopogon japonicus (Thunb)Ker-Gawl, were identified by pharmacognostic specialist, Professor Luan Lianjun. The voucher specimens of red ginseng and ophiopogon were deposited in our laboratory.

Ginsenosides Rb₁, Rb₂, Rb₃, Rc, Rd, Re, Rg₁, Rg₂, 20(s)-Rg₃ and Rh₁ were kindly presented by Professor Xu Jingda, college of pharmacy, Jilin University, PR China.

The structures of ginsenosides were listed in Fig. 1.

2.2. Solvents and reagents

Acetonitrile was HPLC grade from Tedia (Fairfield, OH); acetic acid for analysis from Hangzhou Reagent Company (Hangzhou, PR China); Water was purified by a Milli-Q academic water purification system (Milford, MA).

2.3. Instrumentation and conditions

2.3.1. HPLC-UV analysis

Agilent 1100 series HPLC system (Waldbronn, Germany) equipped with quaternary pump, vacuum degasser, autosampler, diode-array detector, column heater-cooler and ChemStation system was used in HPLC-UV analysis. The chromatographic separation was performed on a Hypersil C_{18} column (5 µm, 4.6 × 250 mm²) from Hanbang Science & Technology (Jiangsu, PR China). A linear gradient elution of A (CH₃COOH:H₂O = 0.01:100) and B (CH₃COOH:H₂O:CH₃CN = 0.01:20:80) was used. The gradient is presented in Table 1. The solvent flow rate was 0.5 ml/min and column temperature was set at 40 °C. The UV detection wavelength was set at 202 nm.

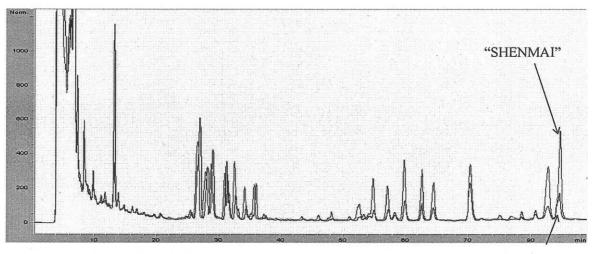
2.3.2. HPLC-ESI-MS analyses

The HPLC conditions for HPLC-MS analyses were the same as those used for HPLC-UV analysis. Esquire LC-00075 series (Bruker, Switzerland) ion-trap mass spectrometer with electrospray ionization was used in HPLC-MS method. ESI-MS conditions of HPLC-MS analysis were as follows: negative ion mode, drying gas N₂, 8 l/min, temperature 320 °C, pressure of Nebulizer 12 psi, octapole voltage 2.35 V, ion-trap voltage 32.2 V, scan range 400–1400 u. ESI-MS/MS conditions were as follows: negative ion mode, separation width 0.9, fragment amplification 1.5, scan range 200–1300 u.

3. Results and discussions

3.1. HPLC analysis of 'SHENMAI' injection and extracts of ginseng and ophiopogon

The typical HPLC-UV chromatograms of SHENMAI injection, ginseng and ophiopogon extract were presented in Fig. 2. The constituents in 'SHENMAI' injection were well separated on reversed-phase column with the linear gradient elution. No obvious peak was detected in ophiopogon extract under different elution gradients with the wavelength scanned from 190 to 390 nm. Furthermore, no ion peak attributed to ophioponins was found in the MS analysis of ophiopogon extract via direct injection. Comparing the chromatograms of 'SHENMAI' and ginseng extract, it's found that their composition were very similar with each other. However, the quantities of each constituent between 'SHENMAI' and ginseng





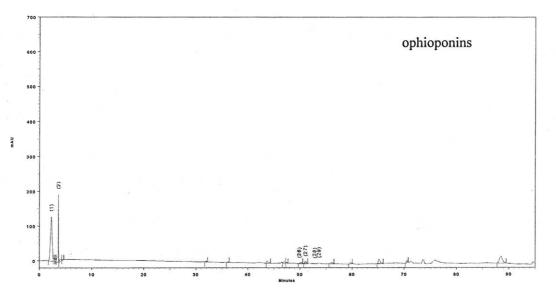


Fig. 2. Chromatograms of 'SHENMAI' injection, ginseng and ophiopogon extract.

showed a big difference, especially the less polarity constituents.

It indicated that the quality of 'SHENMAI' injection relied on ginseng much more than ophiopogon. Moreover, some changes could have taken place in the course of procedure, which resulted in the difference of quantities between the constituents in 'SHENMAI' and ginseng.

3.2. HPLC-UV-ESI-MSⁿ analysis of 'SHENMAI' injection

In HPLC-UV chromatogram, 29 peaks, which were all attributed to ginsenosides, were detected. Each peak had its corresponding peak in HPLC-MS chromatogram except peak 27 (Fig. 3). According to the MS spectra of corresponding

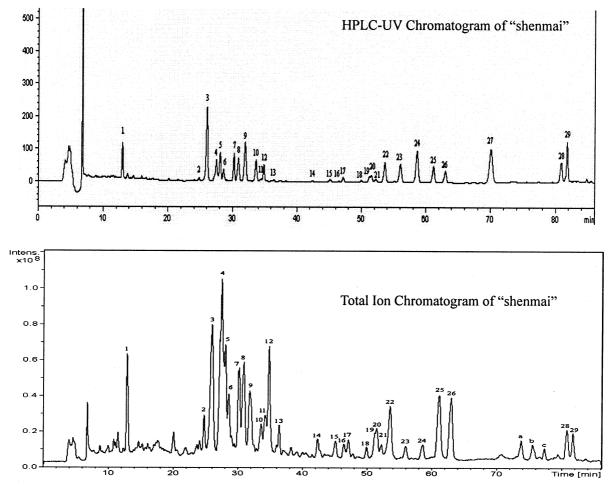


Fig. 3. HPLC-UV Chromatograph and MS-TIC Chromatograph of 'SHENMAI'.

peak in HPLC-UV chromatogram, it could be considered that most constituents in SHENMAI injection were well separated and detected.

In MS spectra, most ginsenosides, excluding malonyl-ginsenosides, exhibited their quasi-molecular ions $[M-H]^-$, adduct ions $[M+Cl]^-$ and $[M+AcO]^-$, some of them showed $[M+M-H]^-$. Since malonyl-ginsenosides are thermally unstable compounds, they would decompose in capillary at 320 °C and lose one molecular CO₂ to produce quasi-molecular ion $[M-CO_2-H]^-$ and adduct ion $[M-CO_2+Cl]^-$, instead of $[M-H]^-$ (Table 2). Chlorine-contained solvents were not employed in HPLC-MS analyses, presence of adduct ions $[M+Cl]^-$ and $[M-CO_2+Cl]^-$ were ascribed to contamination of ion resource.

Compared the retention time and MS spectra of SHENMAI injection sample with those of reference compounds, peak 1, 3, 5, 7–9, 12, 26 were attributed to ginsenoside Re, Rg₁, Rb₁, Rc, Rb₂, Rb₃, Rg₂, Rh₁, Rd and 20(s)-Rg₃, respectively (Table 2). Re and Rg₁, Rb₃ and Rg₂, have same HPLC retention time. They were recognized according to their MS spectral characteristics. Other peaks were identified by careful studies of their MS and MS–MS spectra and by comparison with the literature data ([2–4]). For example, peak 22 and 28 had same m/z value of quasi-molecular

Table 2	
HPLC-UV-ESI-MS	identification

Peak	Retention time	Identification	$[M-H]^-$	$[M+Cl]^-$	[M+AcO] ⁻	[M-H-CO ₂] ⁻	$[M - CO_2 + Cl]^-$	$[M + M - H]^{-}$
1	12.911	Re	945	981	_	_	_	_
1	12.911	Rg ₁	799	835	859	—	—	—
2	24.75	Ra ₁ /Ra ₂	1209	1245	_	_	-	_
3	25.994	Rf	799	835	-	-	-	-
3	25.994	Rb ₁	1107	1143	_	_	-	_
4	27.440	Ro	955	-	-	-	-	-
5	28.044	Rc	1077	1113	-	—	-	—
6	28.563	Notoginsenoside R ₂	769	805	_	_	-	_
7	30.187	Rb_2	1077	1113	-	—	-	—
8	30.875	Rb ₃	1077	1113	_	_	-	_
8	30.875	Rg ₂	783	819	-	-	-	-
9	31.917	Rh ₁	637	673	697	_	-	1275
9	31.917	Rg ₂ iso.	783	819	-	-	-	-
9	31.917	Malonyl-Rb ₁ /iso.	-	-	_	1149	-	_
10	33.596	Rh ₁ iso.	637	673	697	-	-	1275
11	34.34	Malonyl-Rb ₂ /Rb ₃ /Rc	-	-	_	1119	1155	_
12	34.815	Rd	945	981	-	-	-	-
13	36.385	Rb ₂ /Rb ₃ /Rc iso.	1077	_	-	-	-	-
13	36.385	Malonyl-Rb ₂ /Rb ₃ /Rc	-	-	_	1119	1155	_
14	42.37	Malonyl-Rd/iso.	-	-	_	987	1023	_
15	45.11	Not reported	781	817	_	_	-	_
16	46.45	Rg ₂ iso.	783	819	_	_	-	_
16	46.45	Malonyl-Rd/iso.	—	_	-	987	1023	—
17	47.105	Not reported	781	817	_	_	-	_
18	49.96	Not reported	751	787	-	—	-	—
19	51.2	Rg ₆ iso.	765	801	-	—	—	—
20	51.486	Rg ₆ iso.	765	801	_	_	-	_
21	52.180	Not reported	751	787	-	—	—	—
22	53.580	F4/Rg ₆	765	_	-	—	—	—
23	55.993	Rk ₃ /Rh ₄	619	655	679	—	—	1239
24	58.582	Rk ₃ /Rh ₄	619	655	679	—	—	1239
25	61.125	$20(R)Rg_{3}$	783	819	_	_	-	_
26	62.961	20(S)Rg ₃	783	819	-	—	-	—
27	69.997	Not detected	-	-	_	-	-	-
28	80.833	Rk ₁ /Rg ₅	765	801	_	-	_	-
29	81.741	Rk ₁ /Rg ₅	765	801	_	-	-	-
А	73.8	Rs ₃ iso.	825	861	—	-	-	-
В	75.52	Rs ₃ iso.	825	861	_	-	_	-
С	77.35	Ro iso.	955					

ion in the MS spectra. But they could be distinguished using MS-MS spectra. The MS-MS fragment ion (Table 3, Fig. 4) of peak 22 indicated the successive loss of rhamnose and glucose, proved itself to be F4 or Rg_6 , while the MS-MS fragment ion (Table 3, Fig. 5) of peak 28 showed the loss of two glucoses, proved to be Rk_1 or Rg_5 . Several isomers were also detected. Some of them were epimers, including $20(R/S) Rg_3$ as peak 25/26, $20(R/S) Rg_2$ as peak 8/9 and $20(R/S) Rh_1$ as peak 9/10. Some others were geometric isomers, especially those less polarity ginsenosides, for example Rk_1 and Rg_5 (peak 28, 29), which were different just in the position of the double bond.

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Table 3 HPLC-UV-ESI-MSⁿ identification

Peak	Identification	Fragment ion m/z
3	Rf	637[M-H-Glc] ⁻ ; 475[M-H-2Glc] ⁻
4	Ro	793[M-H-Glc] ⁻ ; 631[M-H-2Glc] ⁻ ; 613[M-H-2Glc-H ₂ O] ⁻ ; 587[M-H- 2Glc-CO ₂] ⁻ ; 569[M-H-2Glc-CO ₂ - H ₂ O] ⁻
6	Notoginsenoside	637[M-H-xyl] ⁻ ; 475[M-H-xyl-
	R ₂	Glc] ⁻
19	Rg ₆ iso.	619[M-H-Rham] ⁻ ; 601[M-H- Rham-H ₂ O] ⁻ ;
20	Rg ₆ iso.	619[M-H-Rham] ⁻ ; 601[M-H- Rham-H ₂ O] ⁻ ;
22	F4/ Rg ₆	619[M-H-Rham] ⁻ ; 601[M-H- Rham-H ₂ O] ⁻ ; 457[M-H -Rham- Glc] ⁻
25	20(R)Rg ₃	621[M-H-Glc] ⁻ ; 459[M-H-2Glc] ⁻
26	20(S)Rg ₃	621[M-H-Glc] ⁻ ; 459[M-H-2Glc] ⁻
28	Rk_1/Rg_5	603[M-H-Glc] ⁻ ; 441[M-H-2Glc] ⁻
29	Rk_1/Rg_5	603[M-H-Glc]
А	Rs_3 iso.	783[M-Ac] ⁻ ; 621[M-Ac-Glc] ⁻ ; 459[M-Ac-2Glc] ⁻

MS/MS spectra of peaks, such as 2, 11, 13, 16, 23, 24, and so on, were not detected, probably because of their quantities were so low that the signals were covered by the noise. These peaks were identified by comparing their molecular weight with the literature data. The corresponding 181

literature, they could be supposed to be new ginsenosides which probably were the dehydrated derivatives of notoginsenoside R2 and ginsenoside Rf. The study of these compounds was in process.

4. Conclusions

The quality control has always been the necklace of the development of plant drugs. Although a few strategies were put forward and some progresses were achieved, for example, the fingerprint strategy [5], in which the similarity comparison was used to control the quality, but it's far from solving this problem.

In this study, 'SHENMAI' injection was analyzed by HPLC/MS/MS method and the main constituents were identified. It showed that ginseng would play a much more important role than ophiopogon to affect the quality of 'SHENMAI' injection. Moreover, the big difference between the quantities of the less polarity ginsenosides of ginseng and 'SHENMAI' indicated that these compounds would probably take effects on the bioactivity of 'SHENMAI'. So, some suggestions could be put forward to improve the quality control. Firstly, the quality of ginseng is the key to guarantee the quality of 'SHENMAI'. Sec-

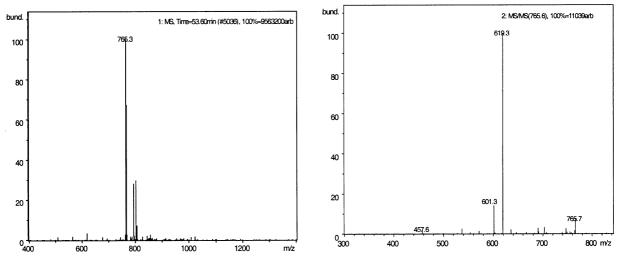
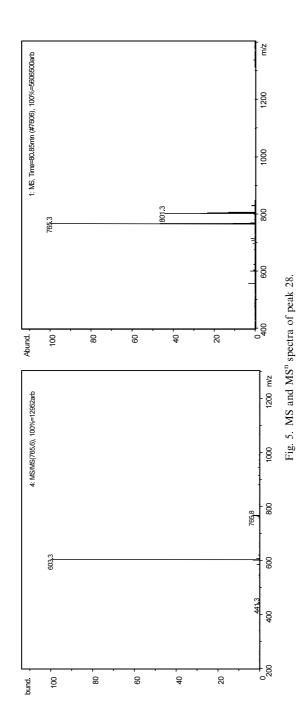


Fig. 4. MS and MSⁿ spectra of peak 22.



ondly, the less polarity ginsenosides could be some of the main active compounds in 'SHENMAI' injection. This suggestion was already confirmed by report [6]. Based on this study and correlative reports of the ginsenosides' bio-activities, further pharmacology tests of the ginsenosides in 'SHEN-MAI' injection could be designed and the indexes for fingerprint could be optimized to improve the quality control efficiency.

Acknowledgements

Thanks to Prof. Xu Jingda for his kind present of standards. The authors are grateful to Dr Zhu Jinke for measuring of HPLC-MS chromatogram. The work was supported by the National Science Foundation of China (No. 20245002).

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