OPEN ACCESS **MOLECULES** ISSN 1420-3049 www.mdpi.com/journal/molecules

Article

Analysis of the Main Nucleosides in *Cordyceps Sinensis* by LC/ESI-MS

Jian-Wei Xie¹, Lan-Fang Huang^{1,*}, Wei Hu¹, Yun-Biao He² and Kin Ping Wong³

- ¹ Department of Chemistry and Pharmaceutical, Quzhou College, Quzhou 324000, China
- ² Changde Institute for Drug Control, Changde 415000, China
- ³ Biology Department, College of Science and Mathematics, California State University-Fresno, Fresno, CA 93740, USA
- * Author to whom correspondence should be addressed: E-Mail: lf18huang@yahoo.com.cn; Tel.: +86-0570-8026668; Fax: +86-0570-8026667.

Received: 10 November 2009; in revised form: 11 December 2009 / Accepted: 7 January 2010 / Published: 13 January 2010

Abstract: A sensitive, selective and reliable liquid chromatography-mass spectrometry coupled with electrospray ionization interface method for simultaneous separation and determination of thymine, adenine, adenosine and cordycepin in *Cordyceps sinensis* has been established. The optimum separation for these analytes was achieved using a gradient elution system and a 2.0 × 150 mm Shimadzu VP-ODS column. 2-Chloroadenosine was used as internal standard for this assay. $[M+H]^+$ ions at m/z 127, 136, 268, 252 and 302 were chosen and selective ion monitoring (SIM) mode was used for quantitative analysis of the four main nucleosides. The regression equations were linear in the range of 1.0–117.5 µg·mL⁻¹ for thymine, 1.8-127.0 µg·mL⁻¹ for adenine, 0.6-114.0 µg·mL⁻¹ for adenosine and 0.5-107.5 µg·mL⁻¹ for cordycepin. The limits of quantitation (LOQ) and detection (LOD) were 1.0 and 0.2 µg·mL⁻¹ for thymine, 1.8 and 0.6 µg·mL⁻¹ for adenine, 0.6 and 0.1 µg·mL⁻¹ for adenosine and 0.5 and 0.1 µg·mL⁻¹ for cordycepin, respectively. The recoveries of the four nucleosides ranged from 98.47 to 99.32%. The developed method was successfully used to determine nucleosides in *Cordyceps sinensis* from different sources.

Keywords: liquid chromatography; electrospray ionization interface; mass spectrometry; *Cordyceps sinensis*; nucleosides

Introduction

Cordyceps sinensis has been used for centuries in the treatment of hyperglycemia, respiratory and liver diseases, renal dysfunction, renal failure and for its antioxidant properties [1,2]. In addition to its usage, it also has been used extensively in China as a folk tonic food or invigorant. Previous studies have found that the most important bioactive constituents in Cordyceps sinensis are soluble nucleosides. Because of its scarcity in Nature and high price, many fake Cordyceps sinensis have been found on the market, so the determination of nucleosides in Cordyceps sinensis is very important for its quality control [3]. Several techniques are available for the analysis of the main nucleosides in Cordyceps sinensis, such as thin layer chromatography (TLC) [4,5], high performance liquid chromatography (HPLC) [6-9] and capillary electrophoresis [10,11]. However, these methods can typically determine only one or two nucleosides, whose contents in *Cordvceps sinensis* are relatively high, and trace amounts of some nucleosides cannot be detected with the above methods, although these trace amount of nucleosides is also very important to the quality of Cordyceps sinensis from the view that its function as a traditional Chinese medicine (TCM) emphasizes its unity. In an earlier report [12], we developed a simple, fast and sensitive isocratic LC/MS method for simultaneous separation and determination of some active components in Cordyceps sinensis. However, thymine (a main nucleoside in Cordyceps sinensis) could not be determined with this method. In this paper, a simple and reliable liquid chromatography (LC)-mass spectrometric detection (MS) coupled with electrospray ionization (ESI) method for simultaneous separation and determination of the main nucleosides in Cordyceps sinensis is described. The developed method has successfully been used for the determination of the four main nucleosides in Cordyceps sinensis from various sources and can thus be used for identification, differentiation and quality evaluation of Cordyceps sinensis.

Results and Discussion

Optimization of chromatographic conditions

When a LC/ESI-MS method is used, the selection of mobile phase should consider both the analyte separation and the effect on ionization. Due to their similar structure the separation of nucleosides can be very difficult if an isocratic HPLC method is used. For the separation of nucleosides in *Cordyceps sinensis* and its substitutes, some reports [6] have used gradient elution with a mobile phase including (A) KH₂PO₄ buffer and (B) methanol, but this solvent system is not suitable for LC/ESI-MS analysis due to its low volatility and high-concentration of salts. In our study, ammonium acetate was chosen to control the acidity of the mobile phase. Using (A) ammonium acetate and (B) methanol as the mobile phase and a gradient elution program, the four main nucleosides under study were successfully separated. Chromatographic conditions, such as concentration of ammonium acetate, pH and solvent elution program, were optimized on a 2.0 × 150 mm Shimadzu VP-ODS column. The HPLC chromatogram of a mixed working standard solution and internal standard (IS) detected with a photodiode array detector set at 260 nm is shown in Figure 1. Retention time (t_R), UV λ_{max} values are shown in Table 1.

Figure 1. HPLC-DAD chromatogram of nucleoside standards: (1) thymine; (2) adenine; (3) adenosine; (4) cordycepin; (5) 2-chloroadenosine.



Table 1. The values of t_R , $UV\lambda_{max}$, $[M+H]^+$ and other MS ions of nucleoside standard compounds.

| Peak number | Compound | t _R /min | UVλ _{max} /nm | $[M+H]^{+}/m/z$ | [M+Na] ⁺ /m/z |
|-------------|------------|---------------------|------------------------|-----------------|--------------------------|
| 1 | Thymine | 3.49 | 203, 264 | 127 | |
| 2 | Adenine | 5.92 | 207, 261 | 136 | |
| 3 | Adenosine | 10.12 | 206, 260 | 268 | 290 |
| 4 | Cordycepin | 13.75 | 207, 260 | 252 | 274 |

Optimization of the ESI-MS conditions

Both an atmospheric pressure chemical ionization (APCI) and ESI interface were tested in positive and negative ion mode by scanning between m/z 50~350 per second. The ESI interface and positive ion mode were chosen and the LC/ESI-MS mass spectra of thymine, adenine, adenosine, cordycepin and 2-chloroadenosine were obtained by scanning at the tested m/z rate of 50~350 per second. The mass spectra were shown in Figure 2. The spectra of adenine, thymine, adenosine and cordycepin were characterized by a protonated molecular ion $[M+H]^+$ as base peak. In addition to the $[M+H]^+$ base peak, the MS spectra of adenosine and cordycepin had a $[M+Na]^+$. The results are also summarized in Table 1. SIM mode involved the use of the $[M+H]^+$ ions at m/z 127, 136, 268, 252 and 302 chosen for simultaneous determination of the four main active components. A mixture containing four standard compounds and 2-chloroadenosine was injected into the LC/MS system. LC/ESI-MS TIC of thymine, adenine, adenosine, cordycepin standard mixture and 2-chloroadenosine in SIM mode is shown in Figure 3A. It was observed that adenine, thymine, adenosine, cordycepin and IS were baseline resolved. Sensitivity optimization was performed by injection of an adenosine standard (20 μ g·mL⁻¹). **Figure 2.** Mass spectra of nucleosides in scan mode: (A) thymine; (B) adenine; (C) adenosine; (D) cordycepin; (E) 2-chloroadenosine.



Figure 3. LC/ESI-MS total ion chromatograms (TIC) of nucleoside standards: (A) and water extracts of *Cordyceps sinensis* from Qinghai Province (B) (1) thymine; (2) adenine; (3) adenosine; (4) cordycepin; (5) 2-chloroadenosine.The concentration of each standard and 2-chloroadenosine was 10 μ g·mL⁻¹.



Method validation

A set of seven non-zero calibration standards of each component, ranging from 1.0-117.5 μ g·mL⁻¹ for thymine, 1.8-127.0 μ g·mL⁻¹ for adenine, 0.6~114.0 μ g·mL⁻¹ for adenosine and 0.5~107.5 μ g·mL⁻¹ for cordycepin was prepared in order to calculate the standard curves. The calibration curves were calculated by plotting peak area ratio (Y) of each analyte and IS in LC/ESI-MS against concentrations (X, μ g·mL⁻¹). Each standard was analyzed in triplicate and the concentration of IS was 10 μ g·mL⁻¹. Results of regression equation, correlation coefficients and linear range are summarized in Table 2.

| Compound | Degracsion equation | R | Liner | Limit of |
|------------|------------------------|--------|---------------------------|-------------------------------|
| name | Regression equation | | range/µg∙mL ⁻¹ | detection/µg·mL ⁻¹ |
| Thymine | Y = 0.06537X + 0.00614 | 0.9984 | 1.0~117.5 | 0.2 |
| Adenine | Y = 0.1438X + 0.0136 | 0.9981 | 1.8~127.0 | 0.6 |
| Adenosine | Y = 0.1292X + 0.0113 | 0.9991 | 0.6~114.0 | 0.1 |
| Cordycepin | Y = 0.2146X + 0.0194 | 0.9993 | 0.5~107.5 | 0.1 |

Table 2. Regression equation, correlation coefficients (R), linear range and limit of detection.

The method selectivity was studied and no interference at the retention times of each analyte was observed. Because thymine, adenine, adenosine, cordycepin all exist in extracts of *Cordyceps sinensis*, no real blank was available for preparation of standards or controls. A solvent blank was analyzed and no peaks at m/z 127, 136, 268 or 252 were observed. The limits of detection of the nucleosides were in the order of $0.1-0.6 \ \mu g \cdot m L^{-1}$, which were determined from signal to noise ratios of 3. Nevertheless, after applying the method to actual samples, the limits of quantification for nucleosides were considered more important. The lower limits of the range of the calibration curves can be considered as the limits of quantification of the method. The intra-assay precision and accuracy were determined by six replicate analyses of each nucleoside standards at concentrations of 2.0, 20.0 and 100.0 $\mu g \cdot m L^{-1}$ on

the same day. The intra-assay precision and accuracy results are listed in Table 3. The precisions were in the range of 1.4-6.3%. The accuracies were on the order of -3.5-4.0%.

| A 14 | Concentration | Concentration | RE | RSD/ |
|------------|---------------------------|---------------------------|------|------|
| Analyte | added/µg∙mL ⁻¹ | found/µg∙mL ⁻¹ | 1% | % |
| Thymine | 2.0 | 2.06 | 3.0 | 6.3 |
| | 20.0 | 19.7 | -1.5 | 3.2 |
| | 100.0 | 101.6 | 1.6 | 1.4 |
| | 4.0^{a} | 3.94 | -1.5 | 4.2 |
| Adenine | 2.0 | 1.93 | -3.5 | 5.4 |
| | 20.0 | 20.4 | 2.0 | 3.1 |
| | 100.0 | 99.3 | -0.7 | 1.6 |
| | 2.0 ^a | 1.95 | -2.5 | 2.9 |
| Adenosine | 2.0 | 1.96 | -2.0 | 5.4 |
| | 20.0 | 20.3 | 1.5 | 3.4 |
| | 100.0 | 99.1 | -0.9 | 1.7 |
| | 2.5 ^a | 2.47 | -1.2 | 3.8 |
| Cordycepin | 2.0 | 2.08 | 4.0 | 6.1 |
| | 20.0 | 20.4 | 2.0 | 2.9 |
| | 100.0 | 99.3 | -0.7 | 1.9 |
| | 2.0 ^a | 2.07 | 3.5 | 5.7 |

Table 3. Intra-assay precision and accuracy of thymine, adenine, adenosine and cordycepin.

^a Aqueous extract of a *Cordyceps sinensis* sample from Sichuan spiked with thymine, adenine, adenosine and cordycepin standards.

The inter-assay precision and accuracy were determined by analyzing above samples that were tested on six consecutive days. The results of inter-assay precisions and accuracies are listed in Table 4. The inter precisions were in the range of 1.6-6.7%. The inter accuracies were in the order of -4.5-4.0%. For determination of recovery, $8.0 \mu g$ thymine, $4.0 \mu g$ adenine, $5.0 \mu g$ adenosine and $4.0 \mu g$ cordycepin were added to 100 mg *Cordyceps sinensis* from Sichuan and extracted as described under "sample preparation" in the Experimental section. This spiked sample was also used to evaluate precision and accuracy. The recoveries were 98.47%, 99.16%, 98.72% and 99.32% for thymine, adenine, adenosine and cordycepin, respectively.

Table 4. Inter-assay precision and accuracy of thymine, adenine, adenosine and cordycepin.

| Analyte | Concentration added/µg∙mL ⁻¹ | Concentration found/µg∙mL ⁻¹ | RE/ % | RSD/% |
|---------|--|--|----------|-------|
| Thymine | 2.0 | 2.08 | 4.0 | 6.7 |
| | 20.0 | 19.4 | -3.0 | 4.2 |
| | 100.0 | 98.6 | 1.4 | 1.8 |
| | 4.0^{a} | 4.10 | -2.5 | 3.8 |

| | 10 | | | | |
|------------|-------------------|------|------|-----|---|
| Adenine | 2.0 | 1.91 | -4.5 | 6.2 | _ |
| | 20.0 | 20.6 | 3.0 | 3.7 | |
| | 100.0 | 98.7 | -1.3 | 1.9 | |
| | 2.0 ^a | 2.04 | 2.0 | 5.4 | |
| Adenosine | 2.0 | 1.94 | -3.0 | 6.4 | |
| | 20.0 | 20.7 | 3.5 | 3.8 | |
| | 100.0 | 98.1 | -1.9 | 2.1 | |
| | 2.5 ^a | 2.58 | 3.2 | 5.3 | |
| Cordycepin | 2.0 | 1.92 | -4.0 | 6.5 | |
| | 20.0 | 20.6 | 3.0 | 3.9 | |
| | 100.0 | 98.2 | -1.8 | 1.6 | |
| | 2.0^{a} | 1.94 | -3.0 | 6.2 | |

 Table 4. Cont.

^a Aqueous extract of a *Cordyceps sinensis* sample from Sichuan spiked with thymine, adenine, adenosine and cordycepin standards.

Application

The developed method was used to determine nucleosides in *Cordyceps sinensis* from different sources. A representative LC/ESI-MS TIC of water extracts of *Cordyceps sinensis* from Qinghai Province in SIM mode were shown in Figure 3B. The determination results are summarized in Table 5. The results showed that there were differences between the nucleoside contents of *Cordyceps sinensis* from different sources. The contents of the four determined nucleosides obtained from different sources also varied considerably from locality to locality. Since nucleosides are important biologically active components, therefore, quality control of this commonly used drug is in great demand. The developed method could be of significant importance for the identification, differentiation and quality evaluation of *Cordyceps sinensis*.

Table 5. Results for the determination of main nucleosides in *Cordyceps sinensis* from different sources.

| Source | Thymine/µg∙g ⁻¹ | Adenine/µg∙g ⁻¹ | Adenosine/µg·g ⁻¹ | Cordycepin/µg·g ⁻¹ |
|---------|----------------------------|----------------------------|------------------------------|-------------------------------|
| Qinghai | 174.2 | 29.7 | 186.5 | 91.2 |
| Tibet | 138.5 | 38.1 | 79.6 | 48.7 |
| Sichuan | 142.6 | 20.4 | 126.1 | 31.3 |

Experimental

Apparatus

The liquid chromatography system (Shimadzu, Kyoto, Japan) consisted of an LC-10Advp solvent delivery pump, an FCV-10ALvp low pressure gradient unit, a DGU-14A degasser, a CTO-10Avp column oven, a SPD-M10Avp photodiode array detector. The mass spectrometer was a LCMS-2010 (Shimadzu, Kyoto, Japan). The column utilized for separation was a 2.0 × 150 mm Shimadzu VP-ODS

column with a particle size of 5 μ m. The analytical column was protected by a C₁₈ Guard-pak cartridge (Waters, Milford, MA, USA).

Materials and standards

Standards of thymine, adenine, adenosine, cordycepin and 2-chloroadenosine (their structures are shown in Figure 4) were purchased from Sigma (St. Louis, MO, USA), A stock solution of each standard ($0.2 \text{ mg} \cdot \text{mL}^{-1}$) was prepared in methanol and stored in a refrigerator. Dilutions of 2.0 and 20.0 µg·mL⁻¹ of each compound were prepared by diluting the stock solution with methanol. Acetic acid and ammonium acetate used were analytical grade. HPLC-grade methanol was obtained from Hanbang Science and Technology CO. (Jiangsu, China) and Milli-Q quality water was used in the preparation of the mobile phase. Solvents were filtered through a 0.45 µm membrane and degassed. *Cordyceps sinensis* from Qinghai, Sichuan and Tibet Autonomous Regin of the P.R. China was purchased from local herb stores in Changsha.

Figure 4. Structures of thymine, adenine, adenosine, cordycepin and 2-chloroadenosine.



Sample preparation

Cordyceps sinensis was dried at 35 °C for two hours and first ground into powder (approximately 20 mesh). Subsequently, it was weighed (about 100 mg) and added to distilled water (20 mL) for extraction of nucleosides. The sample-water mixture was placed into an ultrasonic bath for two hours. Then the sample-mixture was filtered and the filtrate was vacuum-dried. The residue was dissolved in methanol (2 mL).

Chromatographic conditions

The mobile phase including (A) ammonium acetate (40 mM, pH = 5.2) and (B) methanol was degassed ultrasonically before use. Each component of the mobile phase was filtered through a 0.22 μ m membrane. Chromatographic separation was performed at room temperature and a flow-rate

of 0.2 mL/min. The solvent elution program was shown in Table 6. The wavelength of the photo-diode array detector was $190 \sim 300$ nm. The injection amount was 5 μ L.

| | 14 | bie of Solvent elution program for the Le. | |
|------|----------|--|------------|
| Step | Time/min | Ammonium acetate (40 mM, pH=5.2)/% | Methanol/% |
| 1 | 0 | 85 | 15 |
| 2 | 6 | 85 | 15 |
| 3 | 12 | 80 | 20 |
| 4 | 17 | 80 | 20 |
| 5 | 20 | 95 | 5 |

Table 6. Solvent elution program for HPLC.

Mass spectrometric detection conditions

MS coupled with an electrospray ionization (ESI) interface in both scan and SIM mode was used. [M+H]⁺ at m/z 127, 136, 268, 252 and 302 (IS ion) was selected as the SIM ions for qualitative and quantitative analysis. The ESI temperature was 400 °C. The curved desolvation line (CDL) and block temperature were 250 °C and 200 °C, respectively. Probe voltage was +4.5 KV. Detector voltage was 1.5 KV. CDL Voltage was -20 V. Q-array Bios was 50 V. Nebulizing gas flow was 4.5 L/min.

Conclusions

By use of a gradient elution system and 2-chloroadenosine as internal standard, a suitable LC/ESI-MS method for simultaneous separation and determination of the four main nucleosides in *Cordyceps sinensis* has been established and validated. The developed method can be used for the identification, differentiation and quality evaluation of this material. The obtained results show that the LC/ESI-MS method is a powerful technique for the analysis of bioactive components in TCMs.

Acknowledgements

This research work is financially supported by Province Natural Science Foundation of Zhejiang, China (Grant No. Y4080442). Financial support from the College of Science and Mathematics, California State University, Fresno, CA, USA is also gratefully acknowledged. The authors gratefully acknowledge the experimental assistance of Associate Professor Fang-qiu Guo (College of Chemistry and Chemical Engineering of Central South University, Changsha 410083, China) for LC-MS analysis.

References and Notes

- 1. Zhu, J.S.; Halpern, G.M.; Jones, K. The scientific rediscovery of an ancient Chinese herbal medicine: Cordyceps sinensis: part I. J. Altern. Complement. Med. **1998**, *4*, 289–303.
- 2. Zhu, J.S.; Halpern, G.M.; Jones, K. The scientific rediscovery of a precious ancient Chinese herbal regimen: Cordyceps sinensis: part II. *J. Altern. Complement. Med.* **1998**, *4*, 429–457.

- 3. Zheng, F.Z.; Dong, Z.H.; Yu, Q. *Modern Study of Traditional Chinese Medicine*. Xue Yuan Press: Beijing, China, 1999; Volume 6, pp. 99–100.
- 4. Ma, Y.; Wang, Y. Determination of nucleosides in Cordyceps sinensis preparation by dualwavelength TLC-scanning. *China Pharm.* **2008**, *20*, 2375–2377.
- 5. Hu, K.; Fang, D.S. Comparison of Cordyceps Sinensis and solid fermentation of Cordyceps Militaris by TLC. *China Pharm.* **2008**, *19*, 1180–1182.
- 6. Guo, C.; Zhu, J.; Zhang, C.; Zhang, L.J. Determination of adenosine and 3□deoxyadenosine in Cordyceps militaris (L.) Link by HPLC. *Zhongguo Zhong Yao Za Zhi* **1998**, *23*, 236–237.
- 7. Huang, H.N; Hua, Y.Y.; Lin, H.L. Research on rapid and simultaneous determination of functional factor in Cordyceps by HPLC-PDA method. *Strait J. Prev. Med.* **2008**, *14*, 1–3.
- 8. Song J.F.; Liu C.Q.; Li D.J.; Jin B.Q. Determination of cordycepin from cultured cordyceps sinensis by HPLC-DAD. *Food Sci.* **2008**, *40*, 352–354.
- 9. An, J.; Lu, H.C. Determination of adenosine content in Cordyceps militaris(L.) Link by HPLC. *J. Guangdong Coll. Pharm.***2008**, *24*, 123–125.
- Sun, Y.Q.; Sun, G.X.; Jin, Y. Quality control of traditional Chinese medicines by the capillary electrophoresis fingerprint and capillary electrophoresis-mass spectrometry. *Chin. J. Chromatogr.*2008, 26, 160–165.
- 11. Li, J.; Feng, C.Q.; Ni, X.M.; Zhang, W.S. Determination of nucleosides of natural Cordyceps sinensis in Qinghai Province by capillary electrophoresis. *Chin. Pharm. J.* **2008**, *43*, 1105–1107.
- Huang, L.F.; Liang, Y.Z.; Guo, F.Q.; Zhou, Z.F.; Cheng, B.M. Simultaneous separation and determination of active components in Cordyceps sinensis and Cordyceps militarris by LC/ESI-MS. J. Pharm. Biomed. Anal. 2003, 33, 1155–1162.

Sample Availability: Samples of the extracts from Cordyceps sinensis are available from the authors.

© 2010 by the authors; licensee Molecular Diversity Preservation International, Basel, Switzerland. This article is an open-access article distributed under the terms and conditions of the Creative Commons Attribution license (http://creativecommons.org/licenses/by/3.0/).