Exploring processing adjuvants’ influence on traditional Chinese medicine compatibility of Aconiti Radix Cocta and Pinelliae rhizome using rapid resolution liquid chromatography-quadrupole time-of-flight mass spectrometry

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ABSTRACT

Background: It is known that when crude Pinelliae rhizome and Pinelliae rhizoma preparatum are combined with Aconiti Radix Cocta respectively, the toxicity of the combination varies. However, the component’s transformation between different compatibility have remained unclear.

Objective: In this paper, a novel approach using rapid resolution liquid chromatography-quadrupole time-of-flight mass spectrometry (RRLC-Q-TOF-MS) coupled with multivariate statistical analysis was established for exploring the influence of processing adjuvants (PAs) on the compatibility of Aconiti Radix Cocta and Pinelliae rhizome. Materials and Methods: In order to obtain information about the representative markers between different groups, an exhaustive study of different protocols based on adding or removing different PAs step by step was carried out and the influence of PAs on compatibility was investigated. Results: It was found that lime can facilitate diester diterpenoid alkaloids with high toxicity in Aconiti Radix Cocta to be converted into low-toxic or non-toxic derivatives. Glycyrrhizae Radix et Rhizoma had no remarkable effect on the process. Conclusion: The established method in this study will be of great significance to process research mechanism and study on traditional Chinese Medicine compatibility and clinical application.

Key words: Aconiti Radix Cocta, Pinelliae rhizome, processing adjuvants, RRLC-Q-TOF-MS

INTRODUCTION

Paozhi is a common practice that usually occurs before most traditional Chinese herbs were prescribed. The typical processing methods include steaming, boiling, adding adjuvants and so on. During Processing, PAs are additional aiding materials with specific property, flavor and function. Different adjuvants have different properties and functions. They can co-ordinate medicines, strengthen and change drug efficacy, reduce or eliminate drug toxicity and side-effect, influence the physico-chemical properties of basic remedy and change the medicinal position or drug effect. Therefore, characterizing the variation patterns in composition during processing, as well as the compatibility function, is of great importance in clinical application.

Pinelliae rhizome (Banxia, [BX]) is the dry root tuber of the Araceae plant of Pinellia ternata (Thunb) Breit [Figure 1a]. Pinelliae rhizoma preparatum [Figure 1b], known as Fabanxia (FBX), is a processed product obtained by steeping crude Pinelliae rhizome (Shengbanxia, [SBX]) with Glycyrrhizae Radix et Rhizoma (GR) and lime and then drying them together. Chuanwu (CW) is dry mother root of Aconitum carmichaelii Debx. CW is highly toxic and its processed product, Aconiti Radix Cocta [Zhichuanwu, ZCW, Figure 1c], is commonly used in clinical application. The main ingredients of ZCW are five major groups of alkaloids, namely, diester diterpene alkaloids (DDAs), monoester diterpenoid alkaloids (MDAs), amine diterpenoid alkaloids (AMDAs), etc.
alkaloids (ADAs), non-ester alkaloids (NEAs) and lipo-alkaloids (LOAs). DDAs and MDAs were considered as the major toxic aconite alkaloid in ZCW. BX and ZCW are commonly used as Chinese medicine in clinical application. However, several classic works on traditional Chinese medical science recorded that BX and ZCW should not be used together or should be used cautiously, whereas some scholars also believe that the two are compatible. There are more than 20 commonly used proprietary herbal products for stroke, cancer, bone disease and rheumatism from both historical literature and modern clinical reports, containing BX and ZCW as the main ingredients, such as “Chi wan fang,” “Wutoubanxiaisan,” “Zhenfangbaiwanzi,” etc. However, the components transformation in the compatibility process of BX and ZCW is still unclear.

Some reports showed that compared with the extracting solution of ZCW, the extracting solution from boiling SBX and ZCW contained slightly more, less or the same amount of highly toxic DDAs, whereas the extracting solution from FBX and ZCW contained relatively less highly toxic DDAs. The major difference between FBX and SBX is that PAs (GR and Lime) have been used in FBX. Thus, whether BX and ZCW can be used together may depend on the raw BX, processing method and adjuvant used.

In this paper, we used RRLC-Q-TOF-MS analysis platform to investigate the role of PAs in reducing the toxicity of the compatibility process of BX and ZCW, tried to discover the influence of GR and lime on the chemical transformation in content and/or in amount in the processing procedure and compatibility and revealed features of concerted application of BX and ZCW.

MATERIALS AND METHODS

Chemicals, materials and instruments
Agilent 1200 RRLC-Q-TOF-MS (RRLC-Q-TOF-MS, US Agilent Company), qualitative analysis B.04.00 data analysis software (US Agilent Company), Sartorius BS224S electronic balance (Beijing Sartorius Instrument and System Engineering Co., Ltd.) and TGL-16C centrifuge (Shanghai Anting Scientific Instrument Factory) were used in this study. Acetonitrile (ACN, HPLC grade) was purchased from Merck (Darmstadt, Germany). Other chemical reagents were all of analytical grade.

ZCW was obtained from Sichuan Jiangyou Zhongba Aconiti Science and Technology Development Co. Ltd., the SBX and GR were obtained from the Anguo Medical Herbs Market. They were authenticated as Aconiti Radix Cocta, Pinelliae rhizome and GR by Dr. Zhang Lu. The voucher specimens (No. 20120701-20120703) were deposited in the Laboratory of the Chinese Herbal Medicine School of Tianjin University of TCM, China.

Sample preparation
According to Chinese Pharmacopoeia (2010), SBX is added GR and lime when processing to FBX. Therefore, GR and lime were used as the processing markers in sample preparation. Different BX samples were prepared by adding different PAs. If only lime was missing while other adjuvants and procedures were the same, then the mixture obtained is SBX-GR processed product [BX1, Figure 1d], whereas if only GR was missing and the other adjuvants and procedures were the same, then the mixture obtained is SBX-lime processed product [BX2, Figure 1e].

SBX (1 g), FBX (1 g), BX1 (1 g) and BX2 (1 g) were respectively mixed with 1 g of ZCW. 10 times of distilled water were added and the mixtures were soaked for 30 min, then boiled for 0.5 h. Afterward, the mixtures were centrifuged at 10,000 g for 10 min, then 8 times of distilled water were added to the residue, boiled for another 0.5 h and centrifuged at 10,000 g for 10 min. The filtered liquors were combined, which were mixed solutions and diluted mixed solutions in 100 mL measuring flasks and marked as ZCW-SBX, ZCW-BX1, ZCW-BX2 and ZCW-FBX. Meanwhile, 1 g of ZCW without adding any kind of BX,
was extracted according to the procedure above, the solution was marked as the control solution of ZCW (ZCW-C), to show the integrity of chemical component information of ZCW in the same RRLC-Q-TOF-MS spectrum.

**Conditions of data acquisition**

The sample separation was performed with an Agilent 1200 rapid resolution liquid phase system equipped with a Q-TOF mass spectrometer (RRLC-Q-TOF-MS, US Agilent Company) and with data processing Qualitative Analysis B.04.00 software. Using high-purity nitrogen as assist spray ionization and desolventizing gas, the flow rate of drying gas was 10 mL/min and the temperature was set at 350°C; the atmospheric pressure of the atomizer was 310 kPa; flow rate of \( \text{N}_2 \) was 600 L/h; cone gas flow (\( \text{N}_2 \)), 50 L/h. Electrospray ionization capillary voltage was set at +4.0 kV for positive ion mode. The mass scan was over the range of 50-1000 m/z. A ZORBAX SB-Aq C\(_{18}\) column (2.1 × 100 mm, 1.8 μm) (Agilent, USA) was used. The mobile phase was a gradient prepared from 0.1% formic acid aqueous solution (component (A) and 0.1% formic acid ACN (component (B)). Gradient profile consisted of 0-2 min: B 1%; 2-5 min: B 1-6%; 5-15 min: B 6-10%; 15-20 min: B 10-20%; 20-31 min: B 20-30%; 31-36 min: B 30-50%; 36-38 min: B 50-70%; 38-39 min: B 70-99%; 39-40 min: B 99%; 40-45 min B 99-1%; 45-50 min: B 1%. Total run time, including conditioning of the column prior to the initial conditions, was 50 min. The injection volume was 5 μL. The flow rate was kept constant at 0.3 mL/min and analyzes were conducted at 40 ± 1°C.

**Pattern recognition analysis**

After the extraction, peak alignment, peak match and peak intensity correction of ion pair from collected spectrogram by Mass Profiler Professional (USA, Agilent), the data was imported to SIMCA-P 12.0 (Sweden, Umetrics AB) for principal component analysis (PCA), partial least squares-discriminant analysis (PLS-DA) and orthogonal to PLS-DA. Mass data were introduced into SPSS 13.0 (Chicago, IL, USA) for statistical analysis. Statistical differences were considered to be significant if the test \( P < 0.05 \).

**RESULTS AND DISCUSSION**

**Multivariate statistical analysis and chemical markers exploring**

Given that ZCW compatibility of BX’s main components were alkaloids, positive ion mode was used to monitor and optimize RRLC-Q-TOF-MS conditions.[14] Figure 2 showed the BPC picture of ZCW before and after concerted application with different BX in positive ion mode. About 1500 mass signals were collected for further multivariate data analysis and study on ZCW and BX’s compatibility mechanism. In order to compare the differences between different groups, the compounds were classified with PCA and PLS-DA method to find out the differentiated markers. After Pareto scaling with mean centering, the data were displayed as score plot [Figure 3a]. PCA score plot showed sample aggregation and discreteness.[15] The score plot shows that the experimental samples clearly clustered into different groups, indicating that processing procedures caused changes in the composition and/or content of components between various BX samples and ZCW. ZCW-FBX and ZCW-BX2 gathered on one side of the score plot region, but they were far from ZCW; ZCW-SBX and ZCW-BX1 gathered on the other side which was closer to ZCW. In order to find the potential chemical markers for the discrimination between ZCW’s combination with SBX and FBX, extended statistical analysis was performed to generate a loading plot [Figure 3b]. In the loading plot, every spot represented a variable including its intensity and its corresponding \( t_r-m/z \) pair. Chemical markers were selected according to the distance from the origin on the loading plot. The S-plot was used to identify the chemical markers according to the orders of their contributions to the separation of clustering [Figure 3c]. Variable importance parameters (VIP) value was a coefficient that reflected variable importance to the model, wherein, the larger the value, the more important the variable was; a variable with VIP >1 was considered as an important chemical marker to the model.[16-19] The potential chemical markers were selected according to the former VIP value mentioned and loading plot of PLS-DA model. To ensure the accuracy of final result, the established PLS-DA model must be firstly evaluated. \( R^2 \)-Y (cum) and \( Q^2 \) (cum) parameters usually indicated the fitness and prediction of the model. The model with low ratio of the two parameters was not accepted, as the model has over fitted.[20-23] We checked our model established in this study and all the models were proper and all the analysis were right. According to the criterion of data processing established above, 18 components were identified and selected as chemical markers to distinguish significant differences among different groups.

**Identification of chemical markers**

Under the present chromatographic and spectrometric conditions, chemical markers were identified by comparing accurately measured mass value with the theoretical exact mass. The identities of 18 most changed components were identified or tentatively assigned by comparison with the reference compounds or matching the molecular weight with those of the published compounds of ZCW. Taking mesaconitine as an example VIP value, 1.5; mass-to-charge ratio of ion [M + H]+, 632.3068; MS fragments, 572[M + H–C\(_{2}\)H\(_{4}\)O\(_{4}\)]+, 540[M + H–C\(_{4}\)H\(_{2}\)O\(_{4}\)]+, 512[M + H–C\(_{6}\)H\(_{2}\)O\(_{4}\)]+, 105[M + H–C\(_{2}\)H\(_{4}\)NO\(_{1}\)]+. This data showed the characteristics...
of mesaconitine and its elemental composition may be $C_{33}H_{45}NO_{11}$. After treatment in line with the procedures above, 5 chemical markers were found in synergy of ZCW-C and ZCW-SBX, which were hypaconitine, mesaconitine, 10-OH-benzoylmesaconine, 8-lino-14-benzoyl-hypaconine and 8-pal-14-benzoylehycalconine [Table 1]; 6 chemical markers in synergy of ZCW-C and ZCW-BX1: 10-OH-aconitine, aconitine, mesaconitine, 10-OH-benzoyl-mesaconine, 8-lino-14-benzoylehypaconine and 8-pal-14-benzoylehycalconine [Table 1]; and 18 chemical markers in synergy of ZCW-C and ZCW-BX2: Mesaconine, aconine, hypaconine, deoxyaconine, talatisamine, 10-OH-benzoylmesaconine, 10-OH-benzoyleaconine, 14-benzoylemesaconine, 14-benzoyleaconine, 14-benzoylehypaconine, 10-OH-aconitine, 10-OH-mesaconitine, aconitine, mesaconitine, hypaconitine, deoxyaconitine, 8-lino-14-benzoylehypaconine and 8-pal-14-benzoylehycalconine. [Table 1], 17 chemical markers in synergy of ZCW-C and ZCW-FBX: Mesaconine, aconine, hypaconine, deoxyaconine, talatisamine, 10-OH-benzoyl-mesaconine, 10-OH-benzoyleaconine, 14-benzoylemesaconine, 14-benzoylehypaconine, 10-OH-aconitine, 10-OH-mesaconitine, aconitine, mesaconitine, hypaconitine, 8-lino-14-benzoylehypaconine and 8-pal-14-benzoylehycalconine [Table 1]. These chemical markers belonged to DDAs, MDAs, ADAs, NEAs and LOAs, all of them were from ZCW. Under the same RRLC-Q-TOF/MS condition, there were no markers detected from BX sample, thus, combination may have a greater impact on alkaloid compounds in ZCW.

**Intensity change of chemical markers**

Comparing chemical markers of ZCW before and after its synergy with various BX samples, the DDAs in ZCW, such as hypaconitine and mesaconitine, increased in ZCW-SBX group and decreased in ZCW-BX1 group [Figure 4]. The decrease was especially significantly in ZCW-BX2 group and ZCW-FBX group. Whilst other DDAs in ZCW, such as 10-OH-mesaconitine, aconitine and deoxyaconitine, dropped in ZCW-BX2 group and ZCW-FBX group, no obvious decrease was observed in ZCW-SBX group and ZCW-BX1 group; and the ZCW-BX2 group was similar to ZCW-FBX group in terms of DDAs decrease. As shown in Figure 4, the MDAs, such as 10-OH-benzoylemesaconine, also exhibited some decrease in four combination groups. The ZCW-FBX and ZCW-BX2 groups had similar content of MDAs, such as 10-OH-benzoyleaconine, 14-benzoylemesaconine,
14-benzoylaconine and 14-benzylohycaonine, but much lower than that of ZCW-C group. In respect of ADAs, ZCW-FBX and ZCW-BX2 groups had much higher content of mesaconine, aconine, hycaonine and deoxyaconine than the other three groups, whereas those of ZCW-SBX and ZCW-BX1 were similar to ZCW-C group. In terms of NEAs, such as talatisamine, their content in ZCW-C was similar to ZCW-SBX and ZCW-BX1, but lower than ZCW-FBX and ZCW-BX2. LOAs, such as 8-lino-14-benzoylohycaonine and 8-pal-14-benzylohycaonine, also decreased significantly in four combination groups.

![Figure 3: (a) Principal component analysis score plots of Zhichuanwu (ZCW) and its compatibility of different products of BX. (b) Loading plots corresponding partial least squares-discriminant analysis for pattern recognition. (c) S-plot of ZCW-C and ZCW-FBX.](image)

**Table 1: Potential chemical markers were found between ZCW-C and ZCW-SBX, ZCW-BX1, ZCW-BX2, ZCW-FBX**

| Rt (min) | m/z [M+H]+ | Mass accuracy (ppm) | Identified compounds          | Molecular formula | Structure type | Content variance |
|---------|------------|----------------------|------------------------------|-------------------|---------------|-----------------|
| 10.81   | 486.2698   | 486.2652             | -9.46                        | Mesaconine        | C_{24}H_{39}NO_{9} | ADAs            | ↑ ↑             |
| 13.63   | 500.2854   | 500.2823             | -6.20                        | Aconine           | C_{25}H_{41}NO_{9} |               |                 |
| 13.82   | 470.2748   | 470.2703             | -9.57                        | Hycaonine         | C_{24}H_{39}NO_{9} |               | ↑ ↑             |
| 17.51   | 484.2905   | 484.2910             | 1.03                         | Deoxyaconine      | C_{25}H_{41}NO_{9} |               |                 |
| 19.59   | 422.2901   | 422.2882             | -4.50                        | Talatisamine      | C_{24}H_{39}NO_{9} | NEAs            | ↓ ↑             |
| 23.92   | 606.2699   | 606.2615             | 0.99                         | 10-OH-benzylmesaconine | C_{31}H_{43}NO_{11} | MDAs            | ↓ ↓             |
| 24.83   | 620.3065   | 620.3072             | 1.13                         | 10-OH-benzylaconine | C_{32}H_{45}NO_{11} |               |                 |
| 25.58   | 590.2960   | 590.2909             | -8.64                        | 14-benzylmesaconine | C_{31}H_{43}NO_{10} |               |                 |
| 27.23   | 604.3116   | 604.3063             | -8.77                        | 14-benzylaconine  | C_{32}H_{45}NO_{10} |               |                 |
| 27.89   | 594.3011   | 594.2909             | -9.57                        | 14-benzylhycaonine | C_{31}H_{43}NO_{10} |               |                 |
| 27.65   | 662.3171   | 662.3166             | -0.75                        | 10-OH-aconitine   | C_{32}H_{45}NO_{10} | DDAs            | ↓ ↓             |
| 29.15   | 648.3015   | 648.3022             | 1.08                         | 10-OH-mesaconitine | C_{32}H_{45}NO_{10} |               |                 |
| 29.54   | 646.3222   | 646.3221             | -0.15                        | Acouitine         | C_{32}H_{45}NO_{10} |               |                 |
| 31.18   | 632.3065   | 632.3068             | 0.47                         | Mesaconitine      | C_{24}H_{39}NO_{9} |               | ↑ ↑             |
| 33.62   | 616.3116   | 616.3104             | -1.95                        | Hycaonitine       | C_{24}H_{39}NO_{9} | DDAs            | ↑ ↓             |
| 36.08   | 630.3273   | 630.3278             | 0.79                         | Deoxyaconitine    | C_{25}H_{41}NO_{9} |               |                 |
| 42.17   | 835.6307   | 836.5305             | -0.24                        | 8-lino-14-benzylhycaonine | C_{40}H_{73}NO_{10} | LOAs            | ↓ ↓             |
| 42.34   | 813.5307   | 813.5304             | -0.37                        | 8-pal-14-benzylhycaonine | C_{41}H_{75}NO_{10} |               |                 |

*: Content increased; ↓: Content decreased; "": No significant change happened; DDAs: Diester diterpene alkaloids; MDAs: Monoester diterpene alkaloids; ADAs: Amine diterpene alkaloids; NEAs: Non-ester alkaloids; LOAs: Lipo-alkaloids; FBX: Fabanxia; SBX: Shengbanxia; ZCW: Zhichuanwu; BX: Banxia
Based on former mentioned results, it can be concluded that the compatibility of ZCW and FBX may have an important function on reducing the content of aconite alkaloid in ZCW, which may be mainly due to the different PAs used step by step when processing SBX to FBX. The results demonstrated that FBX can facilitate DDAs reduction or the transformation to non-toxic ADAs. As both SBX and FBX were decocted together with ZCW, the difference on content change of aconite alkaloid after ZCW combination with SBX and FBX was not due to the combination extraction. The different effect of two combinations may be related to the distinction between SBX and FBX. As former discussed, FBX was a processed product obtained by steeping SBX. Moreover, the content
change of aconite alkaloid in ZCW-FBX and ZCW-BX2 showed the same trend and ZCW-SBX and ZCW-BX1 had similar behavior. This finding indicated that lime could effectively lower aconite alkaloid content of ZCW-FBX group, whereas the function of GR was not notable.

At present, there was no consensus on the reason why combination of ZCW and FBX can facilitate DDAs reduction or the transformation to non-toxic ADAs. A number of researchers thought it was due to GR and literature about organic acid in GR can reduce toxicity of aconitum in medical materials. However, the amount of GR must be at least half of the amount of ZCW. Only polyoses and other macromolecular components were very clear because lime caused denaturation of starch, whereas sample solutions of ZCW-FBX and ZCW-BX2 contained macromolecular components such as starch from BX, whereas sample solutions of ZCW-FBX and ZCW-BX2 were very clear because lime caused denaturation of starch, polyoses and other macromolecular components.

CONCLUSION

This study adopted RRLC-Q-TOF-MS technology to analyze the Pas’ influence on compatibility of ZCW's combination with SBX and FBX. Multivariate pattern recognition technology was used to study the overall changes in chemical composition before and after combination. The results indicated that PAs affected the plant chemical composition and combination mechanism. For example, lime can facilitate the change of high toxic DDAs into low-toxic or non-toxic components, whereas GR did not have an important role in this process. This study can help to fully understand the differences of chemical composition of ZCW in synergy with BX, as well as provide useful information for potential chemical markers variation. In this study, the established method will be of great significance to processing mechanism research and study of TCM compatibility and clinical application.

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