



Investigation of Steroidal glycoalkaloid concentrations in different potato cultivars through Liquid chromatography-Tandem Mass Spectrometry

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ABSTRACT

A robust analytical method employing solid-phase extraction (SPE) and liquid chromatography-tandem mass spectrometry (LC-MS/MS) was optimized and validated for the determination of α -solanine and α -chaconine in different potato cultivars. Calibration and validation were conducted using a blank matrix derived from sweet potato powder. The method demonstrated excellent linearity, low detection limits, and high precision and accuracy.

Keywords: Liquid chromatography-Tandem Mass Spectrometry, Glycoalkaloids, α -Solanine, α -Chaconine, Potato analysis

1. INTRODUCTION

Steroidal glycoalkaloids (SGAs) are secondary metabolites naturally occurring in plants of the Solanaceae family [1]. These compounds serve as chemical defenses against pests, pathogens, and environmental stresses. Among SGAs, α -solanine and α -chaconine are the most abundant toxic compounds in potatoes, constituting up to 90% of the total glycoalkaloid content [2, 3]. As shown in Figure 1, the core structures of α -solanine and α -chaconine are similar, with the only difference being in the structure of the trisaccharide chain attached to the solanidane triterpenoid molecule. Studies have indicated that α -chaconine is more toxic than α -solanine [4]. While SGAs can have antimicrobial and anticancer properties at low concentrations, their excessive accumulation poses significant risks to human health, including gastrointestinal and neurological effects [5]. To ensure consumer safety, regulatory guidelines limit the total glycoalkaloid (TGA) content to 200 $\mu\text{g g}^{-1}$ of potato fresh weight or 1 $\mu\text{g g}^{-1}$ of potato dry weight [6]. Exceeding these thresholds makes potatoes unsuitable for consumption. The concentration of SGAs in potatoes is influenced by various factors, including cultivar, agricultural practices, and storage conditions. These compounds are typically concentrated in the peel, with lower levels found in the tuber's inner tissues. Therefore, accurate quantification of α -solanine and α -chaconine is essential for monitoring compliance with safety standards. Advances in analytical techniques, particularly liquid chromatography-tandem mass spectrometry (LC-MS/MS), offer



sensitive and selective methods for SGA analysis. Integrating LC-MS/MS with solid-phase extraction (SPE) enhances method accuracy by mitigating matrix effects[7], [17].

This study develops and validates an LC-MS/MS method for the quantification of α -solanine and α -chaconine in four popular commercial potato cultivars. The optimized approach ensures compliance with global safety standards while providing a reliable tool for monitoring glycoalkaloid levels in agricultural products.

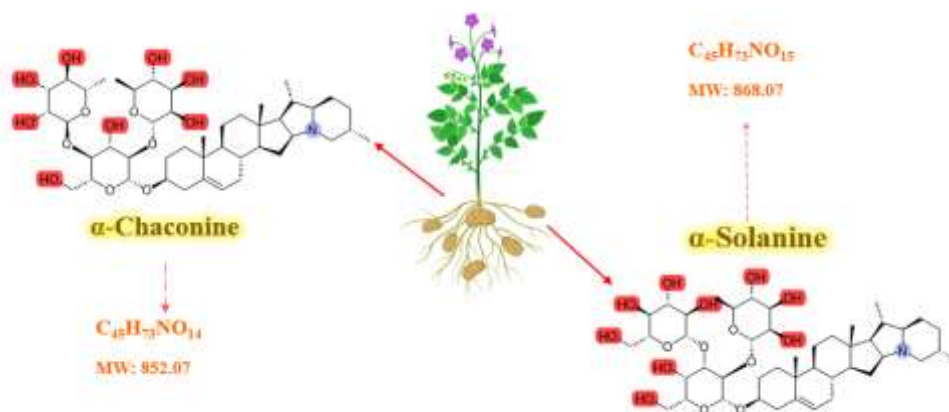


Fig. 1. Chemical structures and molecular weights of α -chaconine and α -solanine

2. Materials and Methods

2.1. Standards and Reagents:

- Glycoalkaloids: α -Solanine, α -Chaconine, and α -Tomatine ($\geq 90\%$ purity, Sigma-Aldrich).
- Solvents: LC-MS grade methanol and acetonitrile (Merck).
- Buffer: Ammonium formate (1 mM , pH adjusted with 0.1% formic acid).

2.2. Optimization of Mass Spectrometry

To optimize the mass spectrometry parameters, direct infusion analysis was performed using a standard solution at a concentration of $0.0 \text{ } \mu\text{g/mL}$. The following settings were determined empirically: capillary voltage -0.10 V , skimmer voltage 30 V , Oct 1 DC voltage 14 V , Oct 2 DC voltage 1.60 V , trap drive voltage $0.8.2 \text{ V}$, Oct RF voltage 183.7 V , capillary exit voltage 121.8 V , Lens 1 voltage -6 V , and Lens 2 voltage -70 V . Each analyte was monitored under single multiple reaction monitoring (MRM) mode in the positive ESI mode using the optimized "SPS mode" of the mass spectrometer's tuning program. Verification was achieved through flow injection analysis of a $0.0 \text{ } \mu\text{g/mL}$ standard solution under the optimized chromatographic conditions. Table 1 presents the specific MRM transitions for α -chaconine, α -solanine, and α -tomatine standards. Protonated ions $[M+H]^+$ were observed at m/z 852.6 , 868.6 , and 1034.7 for α -chaconine, α -solanine, and α -tomatine, respectively.

Table 1. Optimized MRM transitions for α -chaconine, α -solanine, and α -tomatine.

Compounds	Parent ion (m/z)	Production (m/z)
α -Chaconine	852.6	707.0, 060.4, 398.1
α -Solanine	868.6	722.4, 060.4, 398.1



α -Tomatine	۱۰۳۴.۶	۱۰۱۶.۶, ۹۰۲.۵, ۴۱۶.۱
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۲. Instrumentation:

Chromatographic separation was achieved using a HILIC column (4.6×150 mm, $5 \mu\text{m}$) with a mobile phase of 10 mM ammonium formate:acetonitrile ($20:80$, v/v) at a flow rate of 0.3 mL/min. The HPLC system was coupled to an Agilent 7330 ESI-ion trap mass spectrometer equipped with an electrospray ionization interface. Hardware operation and data analysis were performed using Agilent ChemStation software. Direct injection of standard solutions into the mass spectrometer was carried out using a syringe pump (model KDS-CE 100, KDSscientific, U.S.A.).

۴. Sample Preparation

Potato samples were collected from local markets across different provinces of Iran. The potatoes were washed, chopped, and ground using an industrial grinder. The ground material, including both peel and tuber, was freeze-dried and further pulverized into a fine powder. A 0.5 g portion of the dried sample was extracted with 10 mL of 10% acetic acid and sonicated for 30 minutes. The extracts were centrifuged, filtered, and purified using solid-phase extraction (SPE) under a vacuum manifold.

۵. Result and Discussion

Under the optimized chromatographic conditions, the retention times for α -chaconine, α -solanine, and the internal standard (α -tomatine) were determined to be 40.5 , 61.2 and 70.5 minutes, respectively (Figure 2). The method demonstrated excellent quantification capability for both α -solanine and α -chaconine, with calibration curves exhibiting high linearity ($R^2 = 0.9993$ and $R^2 = 0.9955$, respectively). The sensitivity of the method was further highlighted by its low detection (LOD) and quantification (LOQ) limits, as detailed in Table 2.

The accuracy and precision of the method were evaluated at three concentration levels (low, medium, and high) for each analyte. Precision was assessed through intra-day ($n=5$) and inter-day ($n=3$) measurements, expressed as the relative standard deviation (RSD%). The intra-day RSD values ranged from 1.22% to 7.67% , while inter-day RSD values ranged from 2.05% to 7.82% , all of which are below the acceptable threshold of 10% , as recommended by the USFDA for bioanalytical methods. The highest RSD observed was 7.67% , indicating high reproducibility across different analyses (Table 3). Accuracy was determined by calculating the relative recovery percentage (RR%) using Equation 1. The recovery values for α -solanine and α -chaconine were close to the theoretical value of 100% , ranging from 95.31% to 107.53% . These results demonstrate that the method is both accurate and precise for the quantification of α -solanine and α -chaconine in potato samples.

To evaluate the practical application of the method, the total glycoalkaloid (TGA) content was analyzed in four popular commercial potato cultivars in Iran: Agria, Sante, Jelly, and Spirit. The measured TGA levels are presented in Table 4 and Figure 3. Among the cultivars, Agria had the highest TGA concentration (139.9 mg/ 100 g dry weight), followed by Sante, Jelly, and Spirit, which had concentrations of 120.5 , 110.3 , and 95.7 mg/ 100 g, respectively. As illustrated in Figure 3, the bar chart clearly demonstrates these differences, with Agria showing a significantly higher TGA level compared to the other cultivars, while Spirit exhibited the lowest concentration. These findings highlight significant variations in glycoalkaloid content among different cultivars, likely due to genetic differences and environmental factors.

Importantly, the results indicate that the TGA levels in most cultivars exceeded the safety threshold of 100 mg/ 100 g dry weight, which is considered the upper limit for safe consumption. This underscores the need for intervention measures to reduce glycoalkaloid concentrations. One effective strategy involves peeling the potatoes, as glycoalkaloids are predominantly localized in the peel. Studies have shown that removing the peel can significantly lower the overall glycoalkaloid content, making the potatoes suitable for

consumption^[^]. The robustness of the developed method and its applicability for assessing glycoalkaloid levels were demonstrated through its successful application to these cultivars. These findings not only validate the method but also emphasize the importance of routine monitoring and processing interventions in ensuring the safety of potato-based foods, particularly for cultivars with high glycoalkaloid content.

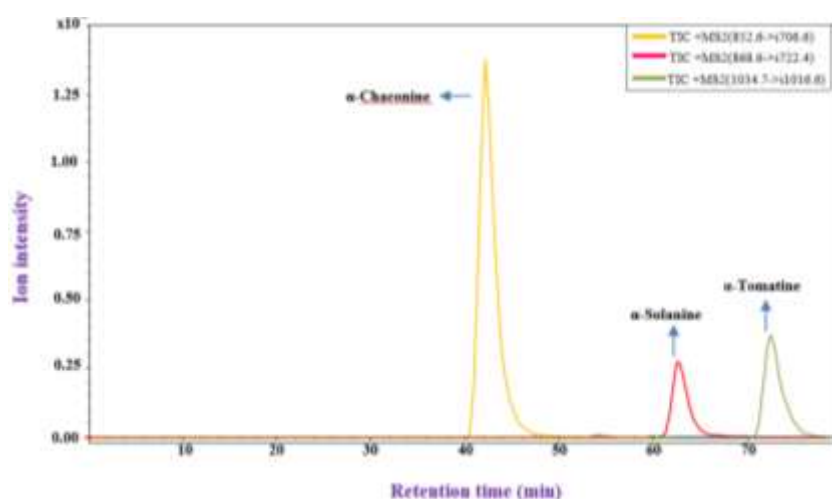


Fig. ۲. Total ion current chromatogram of α -Chaconine ($832.6 \text{ m/z } [M+H]^+$ ion), α -Solanine ($868.6 \text{ m/z } [M+H]^+$ ion) and α -Tomatine ($1034.5 \text{ m/z } [M+H]^+$ ion) in standard solutions.

Table ۲. Correlation coefficients, LODs and LOQs of α -solanine and α -chaconine

Compounds	Correlation coefficient	LOD (mtas)	LOQ (mg/۱۰۰ g DW)
α -Solanine	۰.۹۹۹۳	۲.۰۲×۱۰^{-۳}	۵.۸×۱۰^{-۳}
α -Chaconine	۰.۹۹۵۵	۰.۸۸×۱۰^{-۳}	۳.۴۰×۱۰^{-۳}



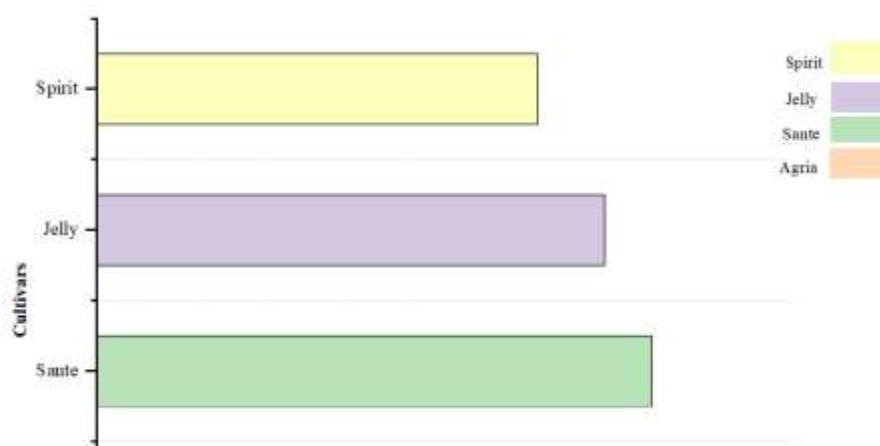
Table 4. TGA content in potato cultivars

Cultivar	TGA (mg/100 g dry weight)
Agria	139.9
Sante	120.5
Jelly	110.3
Spirit	95.7

Table 5. Evaluation of the precision (RSD%) and accuracy (Relative recovery%) of the LC-MS/MS method for quantification of α -Solanine and α -Chaconine at different concentration levels.

Compounds	Spiked concentration (mg/100 g DW)	RSD % Intra-day (n=5)	RSD % Inter-day (n=3)	Relative recovery (RR%) (n=3)
α -Solanine	0.0085	3.06	7.82	107.53 ± 1.03
	0.032	1.50	2.40	95.31 ± 2.15
	0.700	7.67	6.20	103.55 ± 2.37
α -Chaconine	0.0055	2.53	3.16	98.93 ± 1.46
	0.032	1.22	2.05	105.74 ± 1.45
	8.435	5	6.21	100.21 ± 1.22

$$RR\% = \frac{\text{Observed Amount} - \text{Actual concentration in real sample}}{\text{Spiked amount}} \times 100 \quad (1)$$





۶. Conclusion

This study successfully developed and validated a sensitive and reliable method for the quantification of α -solanine and α -chaconine in potato samples using SPE coupled with LC-MS/MS in MRM mode. The method demonstrated excellent linearity, low detection limits, and high precision, making it suitable for glycoalkaloid analysis in diverse potato cultivars. Application of the method to four commercial potato varieties in Iran (Agria, Sante, Jelly, and Spirit) revealed significant variations in total glycoalkaloid (TGA) content among cultivars. Most of the tested varieties exceeded the safety threshold of $100 \text{ mg}/100 \text{ g}$ dry weight, highlighting the necessity of monitoring glycoalkaloid levels to ensure consumer safety. Peeling was shown to be an effective intervention, significantly reducing TGA content and improving the safety of the final product. This validated method provides a robust tool for the accurate assessment of glycoalkaloids in potatoes, supporting food safety efforts and compliance with international health standards. Future studies should focus on understanding the effects of agricultural and environmental factors on glycoalkaloid content and exploring additional strategies to further improve potato safety.

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