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Development and validation of a GC–MS method for rapid determination of galanthamine in Leucojum aestivum and Narcissus ssp.: A metabolomic approach

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1. Introduction

Galanthamine is an acetylcholinesterase (AChE) inhibitor marketed as a hydrobromide salt (Razadyne®, Reminyl®) for the treatment of Alzheimer's disease [\[1\]. A](#page-10-0)lthough its chemical synthesis has been achieved, plants remain an important source for the pharmaceutical industry. Currently, galanthamine is extracted from Leucojum aestivum and Narcissus pseudonarcissus cv. Carlton in Europe, and from Lycoris radiata and Ungernia victoris in Asia [\[2\].](#page-10-0) The search for new plant sources of galanthamine is of great importance for the competitiveness of companies producing this valuable natural product.

The most reported method for the quantification of galanthamine has been HPLC [\[3–7\], b](#page-10-0)ut its capacity to separate complex alkaloid mixtures is generally limited to about 5–7 alkaloids [\[8\],](#page-10-0) and the separation conditions need to be optimized according to the alkaloid composition (more than 100 alkaloids have been found in the genus Narcissus) [\[9\].](#page-10-0) GC–MS studies have shown that the alkaloid mixtures of amaryllidaceous plants include more than 10–15 compounds [\[10,11\]. G](#page-10-0)C–MS, CE-UV [\[12,13\]](#page-10-0) and HPTLC [\[14\]](#page-10-0)

ABSTRACT

Galanthamine, an acetylcholinesterase inhibitor marketed as a hydrobromide salt for the treatment of Alzheimer's disease, is obtained from some Amaryllidaceae plants. A new method was developed and validated for its quantification by GC–MS in different plant sources: bulbs and leaves from Narcissus confusus; bulbs from N. pseudonarcissus cv. Carlton; and leaves and in vitro cultures from L. aestivum. Samples (50 mg) were extracted with methanol (1 mL) for 2 h, then aliquots of the extracts were silylated and analyzed by GC–MS. The calibration line was linear over a range of 15–800 μ g galanthamine/sample, ensuring an analysis of samples with a content of 0.03–1.54% analyte referred to dry weight. The recovery was generally more than 95%. Good inter- and intra assay precision was observed (RSD < 3%). Principal component analysis of GC–MS chromatograms allowed discrimination of the plant raw material with respect to species, organs and geographical regions. The analytical method developed in this study proved to be simple, sensitive and far more informative than the routine analytical methods (GC, HPLC, CE and NMR), so it may be useful for quality control of plant raw materials in the pharmaceutical industry.

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methods have been also validated for analysis of galanthamine in Narcissus bulbs.

Due to the complex composition of plant extracts, before a chromatographic determination the alkaloids are usually fractionated by means of basic–acidic liquid–liquid extraction [\[13,15,16\]](#page-10-0) or solid-phase extraction [\[7,17\]. T](#page-10-0)he methods described in the literature are time-consuming and include laborious sample preparation procedures. They require a relatively high volume of solvents and the use of prepacked columns to handle a high number of samples, which increases the cost of analysis. Direct quantitative determination of galanthamine in unpurified plant extracts by enzyme immunoassays, radioimmunoassays and NMR analysis has been reported [\[18–20\].](#page-10-0) The first two methods are very sensitive, but they involve raising antibodies or the use of radioactive substances, making the studies laborious and expensive. In addition, they do not provide any information about the other metabolites in the samples. The NMR method quantifies galanthamine directly in methanol/water extracts and can determine the origin of the samples by multivariate data analysis of their metabolite fingerprints. Its main disadvantages, however, are the high cost of the equipment (600 MHz NMR), relatively low sensitivity compared with other methods, occasional problems with overlapping signals and a lower number of compounds identified in the extracts.

Metabolomics, including both targeted and global metabolite profiling strategies, is rapidly becoming the approach of choice across a broad range of sciences including systems biology, drug

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discovery, molecular and cell biology, and other medical and agricultural sciences [\[21\]. E](#page-10-0)arlier metabolite profiling of potato extracts demonstrated that the GC–MS platform is a powerful tool for simultaneous detection and identification of a number of metabolites (amino acids, organic acids, mono-, di-, and trisaccharides, sugar alcohols, and aromatic amines) and also for quantification of selected targets in complex plant matrixes [\[22\].](#page-10-0) Multivariate data analysis of metabolite profiles allows discrimination between plant species and cultivars and can therefore be applied for quality control of plant raw materials [\[23,24\].](#page-10-0)

The aim of the present work was to develop and validate a simple and rapid method combining the advantages of GC–MS metabolic profiling (resolution power, sensitivity, selectivity, analysis of a wide spectrum of compounds after derivatization, MS libraries) and multivariate data analysis (discrimination between samples) for direct quantification of galanthamine in extracts and determination of the origin of plant raw material.

2. Experimental

2.1. Chemicals

Galanthamine hydrobromide was supplied by Galen-N Ltd. (Bulgaria) and its purity and identity were checked by GC–MS and ¹H-NMR. Methanol (HPLC grade), chloroform, sulfuric acid and ammonia (analytical grade) were purchased from SDS (France). The hydrocarbon mixture (C9-C36, Restek, Cat no. 31614) was supplied by Teknokroma (Spain). N,O-bis-(trimethylsilyl)trifluoroacetamide (BSTFA) and pyridine were purchased from Sigma–Aldrich (St. Louis, MO, USA).

2.2. Plant material

Dry leaves from Leucojum aestivum L. grown in the Netherlands and Bulgaria were provided by Ludwig & Co (Lisse, Netherlands) and Galen-N (Sofia, Bulgaria), respectively. In vitro plantlets from L. aestivum were provided by VitroFlora (Trzesacz, Poland) and propagated and maintained until the analysis as previously described [\[25\]. B](#page-10-0)ulbs from Narcissus pseudonarcissus cv. Carlton were supplied by Ludwig & Co. Bulbs and leaves from Narcissus confusus (Pursley) were obtained from plants grown in the greenhouse of the Faculty of Pharmacy at the University of Barcelona, Spain. Voucher specimens were deposited at the herbarium of the University of Barcelona (No. 32936; BCN 71625 and BCN 71626).

The fresh plant material was cut into small pieces and dried at 60 ◦C until constant weight. The dried samples were powdered and stored in a chamber maintaining a constant humidity level of 20% until the analysis.

2.3. Methods of analysis

2.3.1. Sample preparation

50 mg of dried plant material was macerated in screw-top Ependorf tubes (1.5 mL of volume) with 1 mL of methanol adjusted to pH 8 with 25% of ammonia and containing 50 μ g of codeine as an internal standard (IS). After 2 h of extraction at room temperature assisted by an ultrasonic bath for 15 min every 30 min, the samples were centrifuged at 10,000 rpm for 1 min. Then, 300 $\rm \mu L$ aliquots were transferred to glass vials and dried by heating at 45 °C. 100 $\rm \mu L$ pyridine and 100 μ L of BSTFA were added to the dried samples and heated at 70 °C for 2 h. After cooling, 300 μ L of chloroform were added and the samples were analyzed by GC–MS.

For GC–MS analysis of alkaloid profiles, 500 μ L aliquots were transferred to other Eppendorf tubes and 500 μ L of 2% sulfuric acid in distilled water was added. The neutral compounds were eliminated by duplicate extraction (vortexing) with 500 μ L chloroform.

The mixtures were basified with 200 μ L 25% ammonia and the alkaloids extracted in triplicate with 500 μ L chloroform. The organic solvent was evaporated and the dry extract dissolved in 300 μ L chloroform for further GC–MS analysis without derivatization.

2.3.2. Chromatographic conditions

The GC–MS spectra were recorded on a Hewlett Packard 6890+ MSD 5975 (Hewlett Packard, Palo Alto, CA, USA) operating in EI mode at 70 eV. A DB-5 MS column (30 m \times 0.25 mm \times 0.25 μ m) was used. The temperature program was: 100–180 °C at 15 °C min⁻¹, 1 min hold at 180 ◦C and 180–300 ◦C at 5 ◦C min−¹ and 1 min hold at 300 ◦C. Injector temperature was 280 ◦C. The flow rate of carrier gas (Helium) was 0.8 mL min⁻¹. The split ratio was 1:15. Ions at m/z 287, 286 and 174 were used to collect SIM chromatograms in SIM mode. For analysis of underivatized alkaloid fractions, a splitless injection was used. 1 μ L of solutions was injected.

2.3.3. Standard solution

12.8 mg galanthamine HBr (equivalent to 10 mg of galanthamine base) was accurately weighted into a volumetric measuring flask of 10 mL. 2–3 drops of 25% ammonia were added and then dissolved in methanol.

2.4. Method validation

The method was validated according to the International Conference on Harmonisation (ICH) guidelines [\[26\]](#page-10-0) on the validation of analytical methods. All results were expressed as μ g/g of dry weight (DW). For statistical analysis Excel 2000 (Microsoft Office) and GraphPad Prizm v. 3.00 were used. A 5% level of significance was selected.

2.4.1. Response function-calibration model

Eight concentration levels of galanthamine trimethylsilyl (TMS) were prepared ranging from 15 to 800μ g/reference solution (0.5 mL). Each reference solution contained 50 μ g of codeine (IS). Each concentration was analysed twice. The ratios of the peak areas of selected ions in total ion current (TIC) mode of galanthamine TMS (m/z at 358) versus those of codeine (m/z at 371) were plotted against the corresponding concentration of galanthamine to obtain the calibration graph.

242 Precision

For intermediate precision four separate samples (100% or 50 mg) were analysed on day 1 and this was repeated on 3 consecutive days. Every sample was injected once. For repeatability at different concentration levels (linearity of the method) four samples with half the amount (50% or 25 mg) and four samples with twice the amount (200% or 100 mg) were analysed using the same method.

2.4.3. Accuracy

The accuracy of the method was investigated by means of a recovery experiment. To 50% of samples (25 mg), a standard solution of galanthamine base was added at three different concentration levels (50%, 100% and 120%) at the start of the analysis of N. pseudonarcissus cv. Carlton bulbs, N. confusus bulbs and leaves and L. aestivum leaves. Five different concentration levels (50%, 100%, 120%, 200% and 230%) were tested for samples (50% or 25 mg) of in vitro obtained cultures of L. aestivum. For each of the concentrations, four samples were analysed according to the developed method.

2.4.4. Specificity

To test the specificity of the method, the peak purity of galanthamine TMS was investigated by AMDIS 2.64 software (NIST, National Institute of Standardization and Technology, Gaithersburg, MD).

2.5. Data analysis

2.5.1. Identification of the metabolites

The compounds of the methanolic extracts were identified as TMS-derivatives with the help of the NIST 05 database (NIST Mass Spectral Database, PC-Version 5.0, 2005) and other plant-specific databases: the Golm Metabolome Database [\[27\], l](#page-10-0)ipid library [\[28\]](#page-10-0) as well as literature data [\[29\]](#page-10-0) on the basis of matching mass spectra and Kovats retention indexes (RI). The measured mass spectra were deconvoluted by AMDIS 2.64 before comparison with the databases. The spectra of individual components were transferred to the NIST Mass Spectral Search Program MS Search 2.0, where they were matched against reference compounds of the NIST Mass Spectral Library 2005 and the Golm Metabolome Database. The groups of unidentified compounds were determined on the basis of their specific mass spectral fragmentation and in comparison with the mass spectra of known metabolites.

RI values of the compounds were measured with a standard n-hydrocarbon calibration mixture (C9–C36) using AMDIS 2.64 software.

2.5.2. Principal component analysis

A target compound library was constructed from the analysed samples by AMDIS software including MS spectra, retention indexes and retention times. The samples were processed using retention index calibration data and the results were exported to an Excel format. The integrated values of the target compounds were normalized to the value of the internal standard (codeine). Principal component analysis (PCA) was performed with Unscrambler® (version 9.8, COMO software Inc.).

3. Results and discussion

In the present work samples from bulbs and leaves of N. confusus, bulbs of N. pseudonarcissus cv. Carlton, leaves from L. aestivum and in vitro cultures from L. aestivum, representing different matrixes, were used for the validation of galanthamine quantification. These plant raw materials have a broad range of galanthamine content. N. confusus is a galanthamine-rich plant species considered as a promising new source of this valuable alkaloid [\[17\]. P](#page-10-0)lant biotechnology may also play an important role in galanthamine supply [\[30\]. I](#page-10-0)n the last 3–5 years studies of galanthamine biosynthesis have been mainly restricted to in vitro cultures of L. aestivum [\[30,31\].](#page-10-0) Therefore, a single validated method covering a broad range of concentration will be of use for the quality control of available and potentially new sources of galanthamine.

3.1. Method development

3.1.1. Sample preparation

Separation (clean-up) of a target compound or a group of structurally related compounds is a crucial procedure in method development and validation by HPLC, GC, CE, HPTLC, etc. In contrast, sample preparation procedures in GC–MS metabolic profiling aiming at the identification and (relative) quantification of the maximum number of metabolites are considerably more simple, including the separation of the total extract into polar and apolar fractions or when dealing with total extracts. In a former study on the extraction efficiency of various solvents, methanol followed by 1% tartaric acid methanolic solution was found to be the most effective for extracting galanthamine and other alkaloids [\[7\]. W](#page-10-0)e decided to work with the total methanol extracts, thus avoiding the separation step for polar and apolar metabolites.

Galanthamine is a tertiary amine with a pK_a of 8.2 [\[32\]](#page-10-0) and therefore its base form has better solubility in organic solvents (including methanol) than its protonated (salt) form. We optimized the methanol extraction (using leaves of L. aestivum) with respect to pH and time, testing pH 7 and 8 at 1, 2, 4 and 6 h. The results, evaluated by means of an ANOVA single factor, showed that the galanthamine content extracted over 2 h with methanol at pH 7 was less (159.0 \pm 1.7 μ g/DW) than at pH 8 (164.3 \pm 1.3 μ g/DW), with longer extraction times producing no significant improvements. Therefore, the extraction of the final method was performed for 2 h with methanol adjusted to pH 8.

The preliminary observations indicated that sugars are the main group of compounds in the methanolic extracts. Aiming at reproducible derivatization and chromatograms, respectively, which can be further used for PCA analysis, we applied a derivatization procedure adapted for sugars [\[29\]. T](#page-10-0)he derivatization test (at 70 ◦C for 1, 2, 3, 4 and 6 h) indicated that after 2 h of reaction time there were no significant differences in the chromatograms. Derivatization of the standards from galanthamine and codeine for 2 h in the conditions described in Section 2.3.1 was complete. Representative chromatograms are shown in [Fig. 1.](#page-3-0)

3.1.2. 3.1.2.Chromatographic conditions

Working with 300 μ L of extract aliquots, a split ratio of 15:1 was found to be optimal with respect to sensitivity to galanthamine TMS and overloading of the column. The temperature ramp used was previously found to be suitable for separation of various groups of metabolites in other amaryllidaceous species [\[33\]. G](#page-10-0)alanthamine TMS was well separated from the other metabolites in all tested samples. Codeine, an alkaloid structurally similar to galanthamine but not synthesized in the Amaryllidaceae family, was found to be suitable as the IS for quantification of galanthamine by HPLC and GC [\[16,34\].](#page-10-0)

3.2. Validation

3.2.1. Response function-calibration model

The calibration was performed plotting the ratio of peak areas of selected ions of galanthamine TMS (m/z at 358) reference standard (15–800 μ g) versus that of the internal standard codeine TMS (50 μ g, m/z at 371). The regression line was constructed and tested on slope and intersept. In order to evaluate the lack-of-fit (LOF) of the linear model a LOF test was performed and the residuals were calculated and graphically examined ([Fig. 2\).](#page-4-0) The values of standard deviation of the residuals (0.078) and F_{LOF} (0.47, F_{crit} = 4.89) indicated a good linearity response on the selected range.

A wide range of the calibration model was chosen due to the broad variation of galanthamine content in plants. In L. aestivum leaves, for example, the amount of galanthamine ranged from 0.03 to 0.57% of DW (unpublished results). The proposed calibration model ensures quantification of galanthamine in plant samples (50 mg) ranging from 0.03 to 1.59% of DW, which covers the biological variability of the studied species. The in vitro cultures, especially undifferentiated callus cultures, may have a very low galanthamine content [\[31\]. F](#page-10-0)or quantification of lower concentrations, another calibration line and validation are required. In this case, the method may be modified with respect to the split, but the flexibility is limited due to a possibility of column overloading. A better option would be to work in SIM (selected ion monitoring) mode, where the sensitivity of the MS detector is significantly higher [\[16\]. S](#page-10-0)uch a validation was not performed because plant raw materials with a very low galanthamine content are not of practical interest.

3.2.2. Limits of detection and limits of quantification

The limits of detection and quantification were determined by analysis of samples with a known amount of analyte. The limit of

Fig. 1. GC–MS of silylated standards of galanthamine and codeine (A) and methanol extracts from N. confusus bulbs (B), N. confusus leaves (C), N. pseudonarcissus cv. Carlton bulbs (D), L. aestivum leaves (E) and L. aestivum in vitro shoot-clumps (F).

Fig. 2. Calibration curve and residual plot of galanthamine TMS.

Table 1

Validation data: precision.

	N. pseudonarcissus cv. Carlton bulbs	N. confusus bulbs	N. confusus leaves	L. <i>gestivum</i> leaves	L. aestivum in vitro shoot-clumps
Precision on different days $(n=3)$ Repeatability Mean $(\mu g/gDW) \pm SD$ (RSD%) Day 1 Day 2 Day 3	$151 \pm 6(2.09)$ $149 \pm 6(3.69)$ $154 \pm 3(2.15)$	1211 ± 21 (1.71) $1183 \pm 23(1.93)$ $1215 \pm 14(1.27)$	$708 \pm 18(2.52)$ $716 \pm 11(1.51)$ $710 \pm 5(0.65)$	$166 \pm 2(1.30)$ $164 \pm 2(0.98)$ $167 \pm 5(3.20)$	$68 \pm 1 (1.59)$ $67 \pm 2(2.28)$ $68 \pm 1 (0.77)$
Intermediate precision Number of groups Number of replicates RSD (%) between groups/Horwitz ^a	3 2.72/3.53	3 1.93/2.61	1.78/2.82	3 1.89/3.52	3 4 1.95/4.08
Precision on concentration levels Repeatability Number of replicates Mean $(\mu g/gDW) \pm SD$ (RSD%) 50% Mean $(\mu$ g/g DW) \pm SD (RSD%) 200%	4 $164 \pm 4(2.08)$ $154 \pm 1(0.47)$	$\overline{4}$ $1233 \pm 35 (2.86)$ $1254 \pm 56(4.43)$	4 $688 \pm 14(2.10)$ $672 \pm 6(0.78)$	4 $166 \pm 3(1.64)$ $164 \pm 2(1.33)$	$\overline{4}$ $69 \pm 2 (3.15)$ $58 \pm 1(1.41)$

^a 2/3 RSD% Horwitz.

detection was found to be 1 μ g/sample (2 μ g/mL) with a signal to noise ratio (S/N) of ca. 3:1. The limit of quantification was accepted as 5 μ g/sample (10 μ g/mL) showing S/N of 42:1 (ICH requires a value > 10:1) and good precision (RSD 2.29%, $n = 4$).

The sensitivity of the detector in SIM mode was also tested, indicating a detection limit of about 1 ng/sample (2 ng/mL, S/N of ca. 3:1) and limit of quantification of 5 ng/sample galanthamine (S/N of ca. 10:1). Thus, an amount of galanthamine as low as 0.000002% of DW can be detected in the samples (50 mg).

3.2.3. Precision

The precision was investigated at two levels: on different days, including repeatability (precision under the same conditions over a short period of time—one day) and intermediate precision (on different days), and precision at different concentrations. The mean, standard deviation and %RSD for each day and concentration level were calculated. The results indicated good intermediate precision (Table 1). With the exception of the values for the second day of the repeatability test for N. pseudonarcissus cv. Carlton, and those for the precision at concentration levels of N. confusus bulbs (RSD < 5%),

3.2.4. Accuracy

The accuracy of the method was investigated by means of recovery experiments, adding a known amount of galanthamine to the samples at the start of the extraction. A mean recovery and %RSD were calculated. The accuracy was checked at 3 levels (50%, 100% and 120%) for the samples from the intact plants and at 5 levels (50%, 100%, 120%, 200% and 230%) for the in vitro samples considering the possible variation of galanthamine content due to further genetic or nutrient medium improvement. The results, presented in Table 2, showed acceptable recovery regarding the concentrations and the purpose of analysis [\[36\].](#page-10-0)

within and between day and level RSDs were less than the limits set by the modified Horwitz equation [\[35\]. A](#page-10-0)NOVA single factor analysis of the results showed that the mean values of the precision test for each type of plant material were not statistically different, with the exceptions of those for 50% level of bulbs from N. pseudonarcissus cv. Carlton and 200% levels of leaves from N. confusus and in vitro cultures from *L. aestivum*, showing deviations of +8%, −6% and −17%, respectively, which could be attributed to matrix effects.

Table 2

Validation data: repeatability and recovery by spiking of different amounts of galanthamine ($n=3$).

Table 3

Metabolites detected in the studied samples.

Table 3 (Continued)

Table 3 (Continued)

uc-Unknown compound. Results represent the means ± SE of response ratios of measurements in 4 samples (50 mg). Response ratio represents peak area ratio using codeine (50 µg) as a quantitative internal standard.

^a Origin: BG—Bulgaria; TR—Turkey; NL—Netherlands.

3.2.5. Specificity

The peak purity of galanthamine TMS was checked by mass spectral deconvoluting software -AMDIS 2.64. In the samples, the extracted spectrum, retention time and retention index of galanthamine TMS were identical with those obtained from the standard compound.

3.3. GC–MS analysis of plant samples

Along with the samples used formethod validation, two samples of L. aestivum leaves from plants of different geographical origin (Turkey and Netherlands) growing in the Netherlands were analyzed for their alkaloid content by means of the proposed method. The galanthamine content of the sample from Turkey was found to be 0.059% DW and that of plants from the Dutch market was 0.079% DW. Both contents of galanthamine were considerably lower than in L. aestivum plants growing in Bulgaria (0.166% DW).

The methanolic extracts showed metabolite profiles characteristic for each species and plant organ. Organic, amino, and fatty acids, sterols, mono-, disaccharides and alkaloids, showing specific time clustering, were detected [\(Fig. 1\).](#page-3-0) About one hundred metabolites were identified [\(Table 3\).](#page-5-0)

The amount of total extract (extractable compounds, estimated on basis of the response ratio), alkaloid fraction, and proportion of

galanthamine in the alkaloid fraction are important characteristics of plant rawmaterial, determining the technology for galanthamine isolation and purification. The amount of the total extract from N. pseudonarcissus cv. Carlton was about 2-times less than from the bulbs of N. confusus. The highest amount of total extract was obtained from the leaves of Bulgarian L. aestivum, which was significantly higher than those of samples grown in the Netherlands [\(Table 3\).](#page-5-0) The alkaloid fraction varied between 0.35% and 15.86% of total extract in the in vitro shoot-clumps from L. aestivum and N. confusus bulbs, respectively [\(Table 4\).](#page-8-0) Several alkaloids were detected as TMS derivatives in the methanolic extracts. To our knowledge, there is no MS data on TMS-derivatives for amaryllidceous alkaloids in the related literature and databases. In [Fig. 3,](#page-8-0) we present MS spectra of the alkaloid TMS derivatives found in the samples. Homolycorine was detected as a non-derivatized compound due to the lack of a hydroxyl group.

Information on the galanthamine proportion in the alkaloid fractions was difficult to obtain from the total methanol extracts due to the low abundance of the minor alkaloids as well as co-elution with other more abundant metabolites. In addition, identification of the Amaryllidaceae alkaloids as TMS derivatives was hampered by lack of reference MS spectra. For that reason, we fractionated the alkaloids and detected 30 compounds without derivatization ([Table 5\).](#page-9-0) The results indicated that the galanthamine proportion was higher

Results represent the means ± SE of response ratios and% of TIC (total ion current) of measurements in 4 samples (50 mg). Response ratio represents peak area ratio using $codeine(50 µg)$ as a quantitative internal standard.

^a Origin: BG—Bulgaria; TR—Turkey; NL—Netherlands.

Fig. 3. GC–MS spectra of the main alkaloids (TMS-derivatives) in the samples.

Fig. 4. Score plot of principal component analysis using GC–MS chromatograms (PC1 vs PC4). NL–N. confusus leaves; NB–N. confusus bulbs; NC–N. pseudonarcissus cv. Carlton bulbs; IV-In vitro cultures of L. aestivum; LL-L. aestivum plants grown in Bulgaria; L-L. aestivum plants grown in the Netherlands (samples 531, 532, 533 and 534-Turkey, 561, 562, 563 and 564—Dutch market).

in the samples from L. aestivum in vitro cultures (74% of total ion current, TIC) and N. pseudonarcissus cv. Carlton bulbs (63% of TIC). Three to four major (>10% of TIC) accompanying compounds were found in the N. confusus samples, with only two major compounds observed in the others.

3.4. PCA analysis

The GC–MS chromatograms of the samples from different plant species, including those of L. aestivum grown at different geographical regions were analysed by PCA. Good separation was

Table 5

Alkaloids identified in the alkaloid fractions.

The values represent the% of TIC from the total alkaloid mixture.

Identification: 1) NIST database; 2) literature data: a [\[37\], b](#page-10-0) [\[38\],](#page-10-0) [\[39\]; 3](#page-10-0)) standard.

Detected in the silyated samples.

** Tazettine is an extraction artifact of pretazettine [9].

observed between the samples by the principal components PC1 and PC4 [\(Fig. 4\).](#page-9-0) Using this unsupervised multivariate data analysis technique, major principal components clearly distinguished between different species. Furthermore, PCA analysis allowed L. aestivum samples from different geographical regions to be distinguished, although it could not clearly separate those from the Dutch market and Turkey, both growing in the same climatic conditions (in the Netherlands). However, a comparison of the chromatograms revealed that both samples of L. aestivum plants grown in the Netherlands could be distinguished by several minor metabolites, whose presence and abundance is characteristic and indicate genetic differences. The sample of Dutch origin contained O-methylleucotamine (**30**) and significantly higher amounts of both alkaloid fractions and lycorine (**25**) than the sample of Turkish origin.

Thus, the PCA analysis of the metabolite patterns allowed an effective control of the plant raw material (regarding species, plant organ and geographic region) used for galanthamine extraction.

4. Conclusion

The analytical method developed in this study, using GC–MS to determine galanthamine in plant sources, proved to be simple, practical and sensitive. The method is very informative compared with the routine analytical methods (GC, HPLC, CE and NMR) and may be useful for the quality control of plant raw materials used in the pharmaceutical industry, providing information on galanthamine content, alkaloid profiles, plant species, the plant organ and geographic region. Due to the low mass of plant material used for extraction, the method may be applied for metabolic analysis in biotechnological or agrochemical experiments.

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