

Glucosinolates Determination in Tissues of Horseradish Plant

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[Abstract] Glucosinolates (GLS) are secondary metabolites mainly found in plants belonging to the Brassicaceae family, including also horseradish (*Armoracia rusticana* G. Gaertn., B. Mey. & Scherb), a popular spice with a characteristic pungent flavor due to the abundance of GLS. Such compounds exhibit antibacterial, antifungal, and insecticidal activities, as well as human health properties. Therefore, it is very important to have a full understanding of their levels and profiles in plants. However, the characterization of GLS from horseradish crude extracts is a tough task, due to the complexity of the vegetal matrix and the occurrence of many GLS in trace amounts. Here we describe two alternative effective and rapid methods for GLS characterization in horseradish plants: Liquid chromatography coupled to high resolution mass spectrometry (LC-MS) for determination of intact GLS and HPLC-UV for determination of desulfo-GLS.

Materials and Reagents

1. Horseradish tissue (hypogeous and epigeous portion) (see Note 1)
2. Methanol (MeOH) (LC/MS grade) (Carlo Erba Reagents, catalog number: 414831)
3. Acetonitrile (ACN) (LC/MS grade) (Carlo Erba Reagents, catalog number: 412342)
4. Ultrapure Milli-Q water
5. Liquid nitrogen
6. Sinigrin hydrate from horseradish (99%) (Sigma-Aldrich, catalog number: S1647)
7. Rapeseed ERM certified Reference Material containing gluconapin, 4-hydroxyglucobrassicin, glucobrassicinapin, glucobrassicin and gluconaturtiin (Sigma-Aldrich, catalog number: ERMBC367)
8. Glucoiberin (C2 Bioengineering, catalog number: 10-JS 12-05-02)
9. Glucobarbarin (C2 Bioengineering, catalog number: 18-DM 19-10-99)
10. Glucotrapeolin (C2 Bioengineering, catalog number: 16-PM 19-10-99)

11. 70% MeOH (see Recipes)

In addition only for intact GLS determination

12. Formic acid (gradient grade) (Sigma-Aldrich, catalog number: F0507)
13. 0.1% Formic acid (HCOOH) (see Recipes)

In addition only for desulfo-method

14. DEAE-Sephadex A-25 (formiate form) obtained by using DEAE-Sephadex A-25 (chloride form) (Sigma-Aldrich, catalog number: A25120) and imidazole (Sigma-Aldrich, catalog number: 56750)
15. Sulfatase type H-1 (Sigma-Aldrich, catalog number: S-9626)
16. Sulfatase type H-1 (1 to 2.5) (see Recipes)
17. DEAE-Sephadex A-25 (formiate form) (see Recipes)

Equipment

1. Freeze Dry Systems (*e.g.* Labconco, model: Freezone 4.5)
2. Laboratory mill
3. Disposable 50 ml and 15 ml polypropylene tubes
4. 2 ml sample vials
5. Water bath: beaker filled with water and placed on a heating device (electric hotplate or similar device).
6. Thermometer
7. Vortex mixer
8. Refrigerated centrifuge (50 ml tubes) (*e.g.* Heraeus, model: Varifuge F)
9. Glass Pasteur pipettes
10. 0.22 μm nylon filter (Whatman)
11. HPLC system with a photodiode array detector (*e.g.* Agilent, model: Agilent 1200 HPLC Liquid Chromatography System)
12. HPLC 2 ml glass vials (Phenomenex, model: AR1-3910-12) with caps (Phenomenex, model: AR0-8959-13-B)
13. Liquid chromatography (LC) coupled with electrospray ionization (ESI) and high resolution mass spectrometry (MS) (*e.g.* Thermo Fisher Scientific, model: LC-ESI-FTICR MS)
14. Intact glucosinolates Discovery C₁₈ column, 250 x 4.6 mm, 5 μm particle size (pore size, 180Å) (Sigma-Aldrich, catalog number: 504971), with a Discovery C₁₈ 20 x 4 mm security guard cartridge (Sigma-Aldrich, catalog number: 505129)

15. Desulfoglucosinolates (Nucleodur C₁₈ column, 125 mm x 3 mm) (MACHEREY-NAGEL, catalog number: MN760051.30)

Procedure

Intact GLS

A. GLS extraction

1. Clean plant material with distilled water and dry with paper towels. Separate roots from epigeous portion, weight and immediately freeze at -80 °C to inhibit myrosinase activity.
2. Lyophilize the frozen tissues and grind to a fine powder using a laboratory mill. Chill roots in liquid nitrogen before the lyophilization to allow the crushing.
3. Weigh 200 mg of frozen dry material in 50 ml polypropylene tubes and place in a water bath heated with an electric hotplate at 70-80 °C for 10 min. During this process, carry out the following steps: After 1 min add 2 ml of 70% methanol solvent and after 5 min mix with vortex for 20 sec.
4. At the end of 10 min remove the polypropylene tube from the water bath and mix again with vortex for 20 sec.
5. Centrifuge at 4 °C for 10 min at 2,400 x g.
6. Collect the supernatant in a disposable 15 ml polypropylene tube by using glass Pasteur pipettes.
7. Extract again the remaining pellet with 2 ml of 10% methanol in water bath heated with an electric hotplate at 70-80 °C for 10 min; after 5 min mix with vortex for 20 sec and the follow the same procedure from step A4-6.
8. Combine the supernatants and vortex the extract for 20 sec.
9. Filter through 0.22 µm nylon filter and transfer into 2 ml sample vials.

B. GLS detection by LC MS

1. Inject 25 µl of each sample in LC-ESI-FTICR MS system equipped with Discovery C₁₈ column, 250 x 4.6 mm, 5 µm particle size (pore size, 180 Å), with a Discovery C₁₈ 20 x 4 mm security guard cartridge.

Settings:

Flow rate: 1 ml/min

Column temperature: 25 °C

Solvent gradient for chromatographic separation:

Time (min)	% Solvent A		% Solvent B	
	0.1% HCOOH water		ACN	
0	90		10	
10	76		24	
12	40		60	
15	90		10	
20	90		10	

Connect the LC system to the mass spectrometer by a laboratory-made splitter with a split ratio of 1:4 after the analytical column to allow 200 $\mu\text{l}/\text{min}$ to enter the ESI source.

2. Perform full-scan experiments in both the linear trap and the ICR cell. Collect data in profile mode in the range of m/z 50–1,000 by setting:

Ionization mode: negative

Resolution: 100.000 (FWHM) at m/z 400

ESI needle voltage: -4.60 kV

Capillary voltage: -22 V

Temperature of the heated capillary: 350 $^{\circ}\text{C}$

Sheath gas (N_2) flow rate: 80 (arbitrary units)

In these conditions GLS elute with a retention time and molecular exact mass to charge ratio as reported in the following table (Agneta *et al.*, 2012; Agneta *et al.*, 2014):

Chemical name	Molecular formulae	Monoisotopic exact value as [M-H] ⁻ (m/z)	t _R
3-(Methylsulfinyl)propyl-GLS	C ₁₁ H ₂₁ NO ₁₀ S ₃	422.02549	4.3
2-Propenyl-GLS	C ₁₀ H ₁₇ NO ₉ S ₂	358.02720	4.4
2-Methylsulfonyl-oxo-ethyl-GLS	C ₁₀ H ₁₇ NO ₁₂ S ₃	437.98402	4.6
3-Butenyl-GLS	C ₁₁ H ₁₉ NO ₉ S ₂	372.04285	5.5
1-Methylpropyl-GLS	C ₁₁ H ₂₁ NO ₉ S ₂	374.05850	6.2
2-Methylpropyl-GLS	C ₁₁ H ₂₁ NO ₉ S ₂	374.05850	6.4
4-Mercaptobuthyl-GLS	C ₁₁ H ₂₁ NO ₉ S ₃	406.03057	6.5
7-Methylsulfinylheptyl-GLS	C ₁₅ H ₂₉ NO ₁₀ S ₃	478.08808	7.3
4-Hydroxyindol-3-ylmethyl-GLS	C ₁₆ H ₂₀ N ₂ O ₁₀ S ₂	463.04866	7.4
Unidentified isomer of 4-hydroxyindol-3-ylmethyl-GLS	C ₁₆ H ₂₀ N ₂ O ₁₀ S ₂	463.04866	7.5
2(S)-Hydroxy-2-phenylethyl-GLS and/or 2(R)-Hydroxy-2-phenylethyl-GLS	C ₁₅ H ₂₁ NO ₇ S	438.05341	7.8
4-Pentenyl-GLS	C ₁₂ H ₂₁ NO ₉ SO ₂	386.05850	7.8
Benzyl-GLS	C ₁₄ H ₁₉ NO ₉ S ₂	408.04285	8.5
Indol-3-ylmethyl-GLS	C ₁₆ H ₂₀ N ₂ O ₉ S ₂	447.05375	10.3
2-Phenylethyl-GLS	C ₁₅ H ₂₁ NO ₉ S ₂	422.05850	12.1
4-Methoxyindol-3-ylmethyl-GLS	C ₁₇ H ₂₂ N ₂ O ₁₀ S ₂	477.06431	12.7
7-Methylthioheptyl-GLS	C ₁₅ H ₂₉ NO ₉ S ₃	462.09317	16.7

Note: It is possible to use the extracts to quantify GLS by preparing a calibration curve for each GLS using a series of dilution and calculating the area under the peak of each compound.

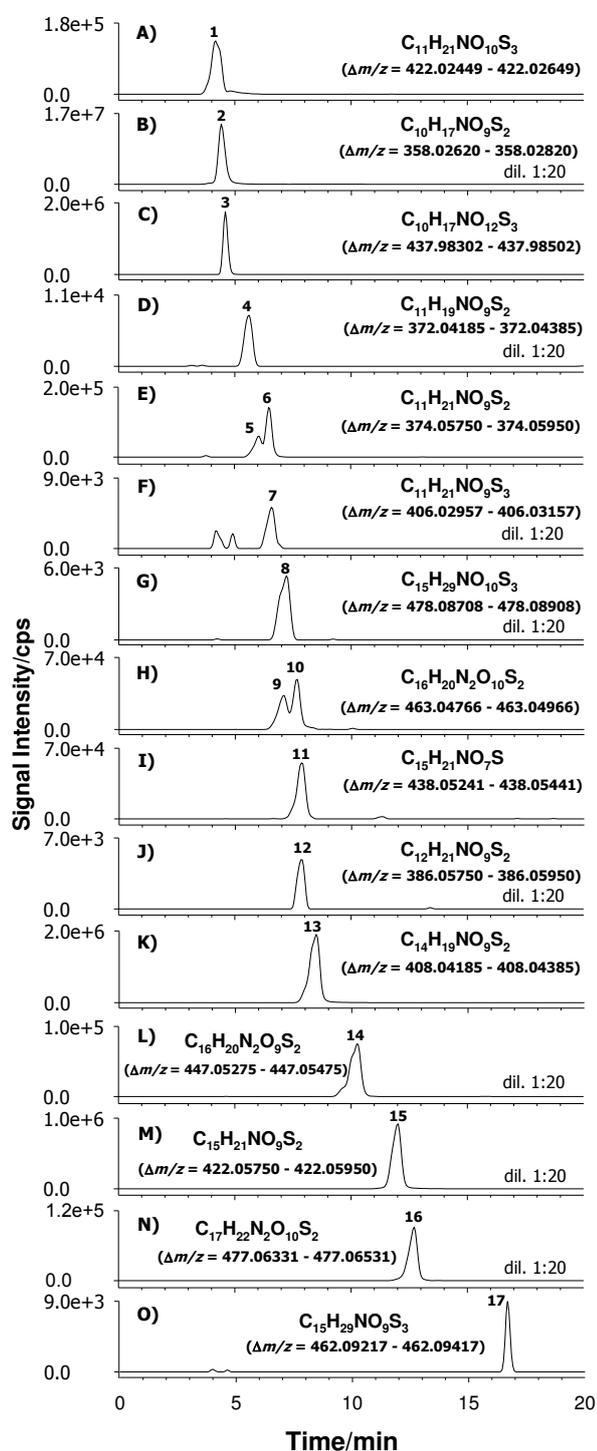


Figure 1. Example of eXtracted Ion Chromatograms (XICs) using high resolution

LC-ESI-FTICR MS acquired in negative ion mode of root crude extract of *A. rusticana*. The ions monitored are displayed in each trace (plots A-O) and correspond to the most abundant deprotonated molecules, $[M-H]^-$, using a restricted window of ± 0.0010 m/z unit centered around each selected ion. Peak numbers in bold correspond to: (1) 3-(methylsulfinyl)propyl-GLS; (2) 2-propenyl-GLS; (3)* 2-methylsulfonyl-oxo-ethyl-GLS; (4) 3-butenyl-GLS; (5)* 1-methylpropyl-GLS; (6)* 2-methylpropyl-GLS; (7)* 4-mercaptobuthyl-GLS; (8)* 7-methylsulfinylheptyl-GLS; (9) 4-hydroxyindol-3-ylmethyl-GLS; (10) unidentified isomer of 4-hydroxyglucobrassicin; (11) 2(S)-hydroxy-2-phenylethyl-GLS and/or 2(R)-hydroxy-2-phenylethyl-GLS; (12) 4-pentenyl-GLS; (13) benzyl-GLS; (14) indol-3-ylmethyl-GLS; (15) 2-phenethyl-GLS; (16) 4-methoxyindol-3-ylmethyl-GLS; (17)* 7-methylthioheptyl-GLS (peaks with asterisk are tentatively assigned). The XIC signals of some peaks were acquired using a sample extract diluted 1:20 with the mobile phase. In bold are indicated also the molecular formula and the range of monoisotopic value as $[M-H]^-$ ion (Δ m/z) (Agneta *et al.*, 2014).

Desulfo GLS

A. GLS extraction

1. Follow the procedure of intact GLS step A1-8.
2. Transfer 500 μ l of the extract on the top of a 20 mg of Sephadex DEAE-A 25 (shortened Pasteur pipet) in the formate form.
3. Wash the column twice with 1 ml of deionized water
4. Add 100 μ l of sulfatase type H-1 diluted 1 to 2.5 with water to achieve the desulfation and incubate overnight at 39 °C (minimum 16 h) (Möllers *et al.*, 1999)
5. Elute desulfatated GLS with 3 x 500 μ l deionized water and collect in 15 ml polypropylene tubes.
6. Vortex the sample for 20 sec.
7. Filter through 0.22 μ m nylon filter and transfer into 2 ml sample vials.

B. GLS detection by HPLC

1. Inject 25 μ l of each sample in HPLC system equipped with Nucleodur C₁₈ column, 125 mm x 3 mm.
Settings:
Flow rate: 0.6 ml/min
Column temperature: 35 °C
UV detection: 229
Solvent gradient for chromatographic separation:

Time (min)	% Solvent A	% Solvent B
	Water	ACN
0	99	1
20	80	20
25	80	20
22	99	1

In these conditions GLS elute with a retention time as reported in the following table:

Chemical name	t_R	Relative response factor ^a
3-(Methylsulfinyl)propyl-GLS	2.8	1.07
2-Propenyl-GLS	4.3	1.00
3-Butenyl-GLS	7.3	1.11
1-Methylpropyl-GLS and/or	8.9	1.00 ^b
2-Methylpropyl-GLS	8.9	1.00 ^b
2(S)-Hydroxy-2-phenylethyl-GLS and/or 2(R)-Hydroxy-2-phenylethyl-GLS	10.1	0.98 ^c
4-Pentenyl-GLS	10.9	1.15
Benzyl-GLS	11.8	0.95
Indol-3-ylmethyl-GLS	13.9	0.29
2-Phenylethyl-GLS	16.1	0.95
4-Methoxyindol-3-ylmethyl-GLS	16.5	0.25

^aThe response factors values used for the desulfated glucosinolates quantification as reported in EN ISO 9167-1 (1992), with except of following desulfated molecules: 1-methylpropyl-GLS, 2-methylpropyl-GLS and 2(S)-hydroxy-2-phenylethyl-GLS and/or 2(R)-Hydroxy-2-phenylethyl-GLS

^bResponse factor of 2-propenyl-GLS as desulfated molecules used for the relative quantification

^cExperimental response factor determinate by using the 2(S)-hydroxy-2-phenylethyl-GLS in comparison to 2-propenyl-GLS as desulfated molecules (see Note 2)

Notes

1. Plant tissues were collected from a field collection of the Institute of Plant Genetics, National Research Council, Thematic Centre for the Preservation of Mediterranean Biodiversity, located in Policoro (MT) (40° 17' 30" N, 16° 65' 16" E), where many accessions of horseradish, previously collected from various villages of the internal areas of the Basilicata region, are maintained and vegetatively propagated (for details see Sarli *et al.*, 2012).
2. It is possible to use the extracts to quantify desulfo GLS by preparing a calibration curve for each GLS by using a series of dilution and calculating the area under the peak of each compound.

It is also possible to quantify desulfo GLS by using the response factors reported in the table above, if you add an internal standard after step A3 (Procedure A), and by applying the following formula:

$$\frac{\mu\text{mol}}{\text{g}} = \frac{\text{Area}_{\text{dg}} \times \text{RF}_{\text{dg}} \times n}{\text{Area}_{\text{st}} \times \text{RF}_{\text{st}} \times m}$$

Wherein:

Area_{dg} is the peak area of the desulfoglucosinolate, Area_{st} is the peak area of internal standard, RF_{dg} is the response factor of the corresponding desulfoglucosinolate, RF_{st} is the response factor of the internal standard, n is the quantity, in μmol, of the internal standard added to the sample, m is the mass, in g, of the sample.

Recipes

1. 70% MeOH
 - 70 ml of MeOH
 - 30 ml of Milli-Q water
2. 0.1% Formic acid (HCOOH)
 - 1 ml of Formic acid
 - 999 ml of Milli-Q water
3. Sulfatase type H-1 (1 to 2.5)
 - 1 ml of sulfatase type H-1
 - 1.5 ml of Milli-Q water
4. DEAE-Sephadex A-25 (formiate form)

Convert the chloride form to the formiate form by adding for each column 500 µl of 6 mM imidazole formiate (dissolve 40 g imidazole in 100 ml 30% formic acid) and then rinsing the column with water.

References

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