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Isolation and identification of 4- α -rhamnosyloxy benzyl glucosinolate in *Noccaea caerulescens* showing intraspecific variation

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Abstract

Glucosinolates are secondary plant compounds typically found in members of the Brassicaceae and a few other plant families. Usually each plant species contains a specific subset of the ~130 different glucosinolates identified to date. However, intraspecific variation in glucosinolate profiles is commonly found. Sinalbin (4-hydroxybenzyl glucosinolate) so far has been identified as the main glucosinolate of the heavy metal accumulating plant species *Noccaea caerulescens* (Brassicaceae). However, a screening of 13 *N. caerulescens* populations revealed that in 10 populations a structurally related glucosinolate was found as the major component. Based on Nuclear Magnetic Resonance (NMR) and mass spectrometry analyses of the intact glucosinolate as well as of the products formed after enzymatic conversion by sulfatase or myrosinase, this compound was identified as 4- α -rhamnosyloxy benzyl glucosinolate (glucomoringin). So far, glucomoringin had only been reported as the main glucosinolate of *Moringa* spp. (Moringaceae) which are tropical tree species. There was no apparent relation between the level of soil pollution at the location of origin, and the presence of glucomoringin. The isothiocyanate that is formed after conversion of glucomoringin is a potent antimicrobial and antitumor agent. It has yet to be established whether glucomoringin or its breakdown product have an added benefit to the plant in its natural habitat.

Keywords: glucosinolates, intraspecific variation, chemotypes, *Thlaspi*, *Noccaea caerulescens*, populations, isothiocyanate

1. Introduction

Glucosinolates are a large class of sulfur and nitrogen containing plant secondary metabolites that are produced by most plant species belonging to the order *Brassicales* (Agerbirk and Olsen, 2012; Fahey, 2005). The core of every glucosinolate molecule is formed by a thiohydroximate group carrying two residues; an S-linked beta-glucopyranosyl moiety, and an O-linked sulfate residue (Agerbirk and Olsen, 2012). Additionally, each glucosinolate is characterized by a variable R-group. This R-group is synthesized from different amino-acids, such as leucine, valine, tryptophan and phenylalanine, and is used to further sub-divide glucosinolates into different structural groups, often referred to as aliphatic, indole and aromatic, or benzylic, glucosinolates (Agerbirk and Olsen, 2012; Fahey et al., 2001). In a recent review Agerbirk and Olsen (2012) reported that around 132 different natural glucosinolates are identified to date. For a part, the distribution of different types of glucosinolates is phylogenetically constrained due to the presence or absence of certain biosynthetic genes in the different branches of the Brassicales' evolutionary tree. For example, it was stated that glucosinolates with glycosylated R-groups appear to be limited to the Resedaceae and Moringaceae (Fahey et al., 2001). Also within plant families there is substantial variation in the ability to produce certain glucosinolates, due to evolutionary events including deletions and small-scale or whole genome duplications contributing to loss or gain of biosynthetic genes (Bekaert et al., 2012). As a consequence, each plant species has its own typical glucosinolate profile which may contain up to 37 different glucosinolates (Kliebenstein et al., 2001).

The glucosinolates, together with the β -thioglucosidase myrosinase, form a specific defence system against herbivores (Ahuja et al., 2010; Hopkins et al., 2009). In intact plants, myrosinase enzymes and glucosinolates are stored separately. As soon as plants are damaged by herbivore feeding or by artificial wounding, myrosinase and glucosinolates are mixed and react. Depending on the pH, and on the presence or absence of specific modifier proteins, such as nitrile specifier or epithionitrile specifier proteins, the glucosinolates are quickly converted to nitriles, epithionitriles or isothiocyanates (Agerbirk and Olsen, 2012; Kissen and Bones, 2009; Kissen et al., 2009; Wittstock and Halkier, 2002). This two-component

defence system has been coined as the “the mustard oil bomb” (Kissen et al., 2009; Ratzka et al., 2002). Especially the isothiocyanates, which also give cabbages and mustards their pungent flavour, are toxic or deterrent to a wide range of herbivores and pathogens (Brown and Morra, 1997; Hopkins et al., 2009; Park et al., 2013). Another property of isothiocyanates is that they contribute to human health. They can function as cancer preventing agents and can inhibit growth of bacteria such as *Helicobacter pylori*, which is the causal agent of gastritis (Fahey et al., 2002; Halkier and Gershenzon, 2006). Especially sulforaphane, the breakdown product of glucoraphanin (4-methylsulfinylbutyl glucosinolate), which is commonly found in high concentrations in broccoli, has been extensively studied and promoted because of its beneficial health effects (Fahey et al., 2002; Verkerk et al., 2009). Similarly, in many tropical countries extracts of the leaves, seeds and roots of *Moringa* tree species, belonging to the family *Moringaceae*, are used for a large range of medical uses (Eilert et al., 1981; Kumar et al., 2010). The main glucosinolate found in this tree is 4-(α -L-rhamnosyloxy) benzyl glucosinolate or glucomoringin, a rhamnose derivative of sinalbin (Amaglo et al., 2010; Bennett et al., 2003; Gueyrard et al., 2010; Mekonnen and Drager, 2003). The isothiocyanate of glucomoringin is a biologically very active compound that is reported to have a beneficial effect on a broad spectrum of human diseases, ranging from bacterial infections to cancer (Fahey, 2005; Faizi et al., 1994; Ragasa et al., 2012).

Until now glucomoringin and structurally related *O*-glycosylated glucosinolates had only been identified in members of the *Moringaceae* and *Resedaceae*. Here we report the identification of glucomoringin and its isothiocyanate breakdown product as isolated from *Noccaea caerulescens* (J.Presl & C.Presl) F.K.Mey., synonym. *Thlaspi caerulescens* J.Presl & C.Presl, which is a member of the *Brassicaceae*. *N. caerulescens* is extensively studied because of its capacity to accumulate large quantities of heavy metals, such as cadmium and zinc, when grown on polluted soils (Assuncao et al., 2003; Leitenmaier and Kupper, 2013). For this reason, the species is used as a phytoremediator to clean soils contaminated with heavy metals due to mining or industrial activities. Several studies were performed to assess the influence of cadmium (Cd) and zinc (Zn) exposure and accumulation on the glucosinolate content of metal hyperaccumulators. Sinalbin and sinigrin, plus several other aliphatic, benzylic and indoyle glucosinolates

were identified in *N. caerulescens* (Tolra et al., 2000) or in related species (*Noccaea praecox*, *Thlaspi arvense*) (Pongrac et al., 2008; Tolra et al., 2006). However, glucomoringin was never reported to be present (Asad et al., 2013; Tolra et al., 2000). For this reason, we isolated the main glucosinolate peak of our *N. caerulescens* extracts and applied advanced NMR and mass spectrometry analyses (Agerbirk and Olsen, 2012; Bennett et al., 2006) to accurately identify the molecular structure of this glucosinolate. Furthermore, we analysed seeds and leaves from 13 different populations in Europe to study natural variation in glucosinolate profiles in this species.

2. Results and Discussion

Desulfoglucosinolate extracts of *N. caerulea* were first analysed by HPLC-PDA (229 nm) following standard procedures (see section 4. Experimental). In addition to sinalbin ($R_t = 9.3$ minutes), an unknown glucosinolate with a similar UV absorption spectrum but eluting at 10.4 minutes was detected in several of the plant populations that were analysed (Figure S1). The unknown desulfoglucosinolate as well as its intact parent glucosinolate were isolated from the respective *N. caerulea* extracts and subjected to comprehensive NMR and mass spectral analyses to obtain the molecular structure. ^1H and ^{13}C NMR spectra were recorded both from the intact and the desulfated glucosinolate, resulting in the spectra as shown in Table 1 (numbering of the carbon atoms in as in Figure 1). As expected, the ^1H and ^{13}C NMR spectra of the desulfo- and the intact glucosinolate were similar.

In addition, the corresponding isothiocyanate (Figure 1, structure 3) of the unknown glucosinolate was produced by myrosinase treatment and purified on HPLC. ^1H and ^{13}C NMR spectra were recorded from the resulting product and identified as 4- α -rhamnosyloxy benzyl isothiocyanate (C- atom numbering as in Figure 1, results Table 1). The α -rhamnose conformation of the sugar moiety was confirmed according to the J-couplings of the protons on the 1'' to 5'' position. As a reference, the ^1H and ^{13}C NMR spectra of sinalbin (Figure 1, structure 4) were recorded. This allowed us to distinguish the NMR peaks belonging to the glucose (present in both glucosinolates) and rhamnose (only in the unknown glucosinolate) moieties in the unknown glucosinolate (Figure 1, Table1).

Furthermore g_2DHSQCAD and 2D COSY analyses were performed on sinalbin as well as the unknown glucosinolate to study proton-carbon and proton-proton coupling (Figures S1, S2, S3). Based on these data, the unknown glucosinolate in *N. caerulea* was identified as 4- α -rhamnosyloxy benzyl glucosinolate.

The identification was further confirmed by high resolution mass spectrometry of the desulfated glucosinolate. LC-MS ESI analyses (positive mode) yielded an m/z 491 which corresponding to the mass

of desulfated glucomoringin (Figure 1, structure 2). Using high resolution mass spectrometry with ESI detection in positive mode, an accurate mass of 514.13597 Dalton was obtained. This mass deviates only 0.07 mmu from the calculated monoisotopic mass of $C_{20}H_{29}O_{11}NSNa$ (m/z 514.13590), a sodium adduct of desulfated glucomoringin. In these systems, where the full extract is directly inserted in the TOF-MS and not purified over a column, sodium salts are commonly observed (De Nicola et al., 2012).

Thus, the mass spectral data in combination with the NMR data unequivocally identify the isolated unknown glucosinolate from *N. caerulescens* as 4-(α -rhamnosyloxy) benzyl glucosinolate or glucomoringin. Even though our analytical platforms do not allow to unequivocally discriminate between the L- and D-form, it is highly likely that we have identified 4-(α -L-rhamnosyloxy) benzyl glucosinolate, or glucomoringin. We based this conclusion on the high similarity of our glucosinolate with the glucomoringin extracted from *M. oleifera* and the fact that L-rhamnose is the natural configuration of this deoxy sugar in plants (Giraud and Naismith 2000). Interestingly, glucosinolates with glycosylated R-groups were believed to occur only in members of the Moringaceae or Resedaceae families (Fahey et al., 2001). Papers reporting the effect of metal pollution on glucosinolates always have reported sinalbin and sinigrin as the main glucosinolates in *N. caerulescens* (Asad et al. 2013, Tolra et al 2000). For this reason, we screened different accessions of *N. caerulescens* for the presence of glucomoringin. Of the 13 different accessions that we investigated, only three, all originating from Zn mining or smelting sites in the North East of Belgium, lacked glucomoringin and indeed contained sinalbin as the main glucosinolate (Figure 2, Table 2). All remaining accessions contained high levels of glucomoringin as well as low levels of sinalbin, which suggests that the latter may be the precursor for glucomoringin. Accessions from which we analysed both seeds and leaves showed that the presence or absence of glucomoringin is consistent among these samples: the Lellingen and Ganges accessions contained glucomoringin as a major compound in both seeds and leaves, whereas sinalbin was the major benzylic glucosinolate in La Calamine and Prayon leaves and seeds (Table 2). Interestingly, this congruency between seed and leaf profiles does not exist for sinigrin: this aliphatic glucosinolate was present in all leaves samples but was

not detected in seeds (Table 2). We found no correlation between soil pollution and the presence or absence of glucomoringin. Instead, only the three closely related accessions from La Calamine, Plombières and Prayon lack glucomoringin, which suggests this glucosinolate polymorphism reflects a regional genetic variation rather than a relation with heavy metal exposure. The conversion of sinalbin to glucomoringin could very well be performed by a single enzyme and a single loss-of-function mutation of the gene encoding it would already render plants unable to make glucomoringin. Recently the transcriptome of *N. caerulescens* accession Ganges has been published (Lin et al., 2014), which could provide a way to identify the rhamnosyltransferase gene involved. In contrast to what has been found in *Moringa* spp., we could not identify the acetylated form of glucomoringin in *N. caerulescens* (Bennett et al., 2003).

Given the predominance of glucomoringin in the populations we analysed, it is remarkable that this compound has never been detected before in *N. caerulescens* (e.g. in Asad et al 2013, Tolra et al. 2000). At present, we can only speculate about the reason. Possibly, the genotypes that were used for the previous experiments by chance did not contain glucomoringin. However, it may also be possible that on HPLC-PDA glucomoringin has been misidentified as the UV spectra of sinalbin and glucomoringin are very similar. A broader screening effort would be needed to elucidate how widespread the presence of glucomoringin in *N. caerulescens* populations is. Because the ability to produce glucomoringin is also present in populations from heavily contaminated soils and because of the potent medical properties of glucomoringin and its isothiocyanate (Eilert et al., 1981; Fahey, 2005; Faizi et al., 1994; Kumar et al., 2010; Ragasa et al., 2012) this plant species thus may be used for phytoremediation as well as the production of medicinal phytochemicals.

It has been shown before that small changes in the side-chain structure of a glucosinolate may have severe effects on the resistance to herbivores (van Leur et al., 2008). It is as yet unknown how the ability to produce glucomoringin may affect the preference and performance of herbivores feeding on this species. Theoretically, the presence of the rhamnose on the side-chain will make the isothiocyanate that is formed

more water soluble than comparable isothiocyanates without a sugar moiety. As a consequence, the isothiocyanate may be better taken up by herbivores or be transported over longer distances in the soil, thereby potentiating the effect of glucomoringin conversion products on pathogens and herbivores in the plant's environment. Current attempts to synthesize different glycosylated forms of sinalbin, including glucomoringin, may help to assess the role of side-chain glycosylation on the biological effect (Gueyrard et al., 2010).

3. Conclusions

This is the first unequivocal report of the presence of glucomoringin in *N. caerulesens*, and likely also the first well-founded report on the presence of glucosinolates with a glycosylated side chain in the Brassicaceae. It has yet to be assessed if there is a specific ecological function for this type of glucosinolates, e.g. as a more potent resistance mechanism against insects. Our finding opens the possibility to use this plant species for the production of the pharmaceutically interesting glucosinolate glucomoringin in temperate climate zones

4. Experimental

4.1 Chemicals

If not mentioned otherwise, all chemicals used in the procedures were of ACS quality or higher. Reference sinalbin (K-salt) was obtained from Phytoflan (Heidelberg, Germany).

4.2 Plant materials

Seeds of 13 *N. caerulea* accessions were collected from different locations in Europe (Table 2). Of several of these accessions, plants were grown in a climate controlled greenhouse. Leaf samples of these plants were directly frozen in liquid nitrogen, freeze dried and stored at -80°C until processing. *Moringa oleifera* leaf samples collected in New Delhi, India, were kindly provided by Dr Vartika Mathur, Sri Venkateswara College, University of Delhi, India. The air-dried samples from India were frozen upon arrival, freeze dried, ground and extracted like the other leaf samples.

4.3 Isolation and analysis of the desulfated glucosinolates

The dry plant material or seeds were ground with a Retsch MM 300 mixer mill at 30 Hz for 1 minute. Aliquots of 50.0 mg leaf or 2.5 mg seed material were weighed in 2.0 ml Eppendorf tubes after which 1.0 ml of a 70% methanol solution in water was added to the samples. The tubes were capped, vortexed and immediately placed in a water bath at 90°C for 5 minutes to stop myrosinase activity. After boiling, the tubes were placed in an ultrasonic bath for 15 minutes and centrifuged at 4500 r.p.m. for 10 minutes. The supernatant was collected and the remaining pellet was extracted once more with 1 ml 70% MeOH starting with the ultrasonic bath. Both supernatants were combined and added to a DEAE-Sephadex A-25 column (5x10mm). After the extracts were added, the columns were washed twice with 1 ml 70% MeOH, once with 1 ml MilliQ water and twice with 1 ml 20 mM NaOAc buffer (pH 5.5). Thereafter, 20 µL of aryl sulfatase solution (Sigma type H-1 of *Helix pomatia*) was added to the columns and flushed down with 50 µL NaOAc buffer (pH 5.5). The columns were covered with aluminium foil and incubated

overnight at room temperature. Thereafter, the resulting desulfoglucosinolates were eluted from the columns with two times 0.75 ml MilliQ water. The extract was freeze-dried, the residue redissolved in 1.0 ml of MilliQ water and stored at -20°C until further analysis. Quantitative HPLC analyses of the desulfated glucosinolates were performed on a Dionex Ultimate 3000 HPLC (Dionex, Sunnyvale, CA, USA) equipped with a C-18 column (Alltima C-18, 150 x 4.6 mm, 3µm; Alltech, Deerfield, IL, USA) held at 40°C and eluted with a linear water-acetonitrile gradient (2%-35% HPLC-grade acetonitrile in MilliQ water in 30 minutes). Peak detection was performed by a photodiodearray detector (PDA) with 229 nm as integration wavelength. Sinigrin (63, 188, 375, 500 and 625 µM) was used as an external standard. To calculate the concentrations of sinalbin and glucomoringin based on the sinigrin reference curve, we used 0.4 as the relative response factor (see Brown et al 2003).

Isolation of the unknown ds-glucosinolate peak detected at 10.4 minutes was performed on an Agilent 1100 HPLC (Agilent, Santa Clara CA, USA) equipped with a fraction collector (G1315B). Peaks were separated on a Merck LiChrospher 100 RP-18 column (250mm x 4.6mm, 5 µm; Merck, Darmstad, Germany). A 6% acetonitrile solution in water (1.0 ml/min) was used for isocratic elution of the intact glucosinolates, which were detected by PDA at 229 nm. The isolated peak had a retention time of 5.1 minutes (fraction collected 4.5-5.9 minutes).

4.4 Isolation of the intact glucosinolate by HPLC:

The intact glucosinolate was isolated using a modification of the method by (Agerbirk and Olsen, 2012). A total of 1.25 g ground, freeze-dried leaf sample was equally divided over eight 2-ml Eppendorf tubes. To each tube 1 ml 70% methanol in water was added, vortexed, followed by extraction in an ultrasonic bath for 15 minutes at 30°C, after which the tubes were incubated in a 90°C water bath for 12 minutes. Thereafter the samples were centrifuged for 10 minutes at 5000 r.p.m. in a table centrifuge. The supernatant was removed and the remaining pellet was extracted once more with 1 ml 70% methanol. The combined supernatants of all 8 tubes were transferred to a DEAE Sephadex A-25 column (30 mm x 6 mm) that was equilibrated with MilliQ. Subsequently the column was washed with 4 ml 70% methanol

followed by 2 ml MilliQ. The intact glucosinolate was eluted with 15 ml 5% ethanol in 0.5M KHSO₄. The eluent was evaporated to dryness with a Rotavapor and the residue was dissolved in 5 ml MilliQ. The isolated intact glucosinolate was purified and desalted on HPLC. Aliquots of 100 µl were injected on an Agilent 1100 HPLC equipped with a Merck LiChrospher 100 RP-18 reversed phase column (250 mm x 4.6 mm; 5 µm). An isocratic program with 100% MilliQ flow 1 ml/min was used for separation and a broad peak with a retention-time between 8.5 and 14.5 minutes was collected. The volume of the combined fractions was reduced using a Rotary evaporator (Büchi) and further dried over NaOH in a vacuum desiccators.

4.5 Preparation and isolation of the isothiocyanate

The unknown intact glucosinolate (about 13 mg) was dissolved in 2 ml buffer (126 mM Na₂HPO₄·2H₂O, 37 mM citric acid monohydrate, 1 mM L-Ascorbic acid, pH = 6). To this solution, 40 µl thioglucosidase (1 u, Sigma-Aldrich E.C. 3.2.1.47 isolated from *Sinapis alba*) was added and incubated at room temperature for 2 hours. Aliquots (100 µl) of the solution were injected on an Agilent 1100 HPLC equipped with a Merck LiChrospher 100 RP-18 column (see above) kept at 40°C and eluted as follows: flow rate 1 ml/minute, 2 minutes isocratic 15% acetonitrile, followed by a gradient to 80% acetonitrile in 18 minutes. The fraction from 15-18 minutes was collected using an automated fraction collector. The volume of the combined fractions was reduced using a Rotary evaporator (Büchi) and further dried over NaOH in a vacuum desiccator.

4.6 NMR analyses:

All NMR spectra were recorded with a Varian Inova 400 MHz NMR (1H) (100 MHz for ¹³C) equipped with a 5 mm PFG Indirect Detection Probe. About 1 mg of the unknown compound was dissolved in D₂O (Aldrich 99.9%D) or CDCl₃ (Aldrich, 99.8%D) in the case of the isothiocyanate, final concentration <0.1M. Proton spectra were recorded in 2-4 minutes, ¹³C and 2D g HMBC spectra in about 6 hours.

4.7 Mass spectrometry analysis

LC-MS analysis of the desulphoglucosinolates was performed on a Thermo Finnigan LCQ Fleet ion trap mass spectrometer (Thermo Fisher Scientific Inc., Waltham, MA USA) equipped with the same C18 column (Altima) and a similar eluent program as used for the analytical separations of the desulfated glucosinolates. Peaks were detected using ESI in positive mode.

The exact mass of the unknown isolated desulphoglucosinolate was determined on a JEOL AccuTOF CS JMS-T100CS mass spectrometer (JEOL Ltd. Tokyo, Japan) with ESI detection in positive mode for high resolution mass spectrometry. Samples were analysed by infusion in methanol (MeOH Absolute ULC/MS, Biosolve BV, Valkenswaard, Netherlands) containing 0.1% formic acid (for MS ~98%, Fluka, Germany) and 50 nM PPG425 (polypropyleneglycol average Mw = 425, Sigma Aldrich Chemie, Steinheim, Germany, as an Internal Mass Drift Compensator) at a flow of 100µl/min using an Agilent 1100 HPLC system without column installed.

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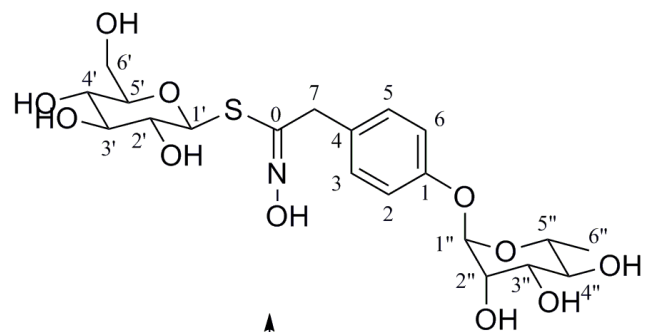
Table 1NMR Spectroscopic Data for desulphated Glucomoringin (**2**) in D₂O, Sinalbin (**4**) in D₂O and for the isothiocyanate of glucomoringin (**3**) in CDCl₃.

| | 2 | | 4 | | 3 | |
|-----------------|------------|----------------------|------------|----------------------|------------|----------------------|
| | δ_C | δ_H (J in Hz) | δ_C | δ_H (J in Hz) | δ_C | δ_H (J in Hz) |
| Glucose moiety | | | | | | |
| 1' | 80.9 | 4.57, d, (9.0) | 81.2 | 4.55, d, (9.7) | | |
| 2' | 71.9 | 3.17-3.30, m | 71.7 | 3.05-3.24, m | | |
| 3' | 77.0 | 3.17-3.30, m | 76.9 | 3.05-3.24, m | | |
| 4' | 68.8 | 3.17-3.30, m | 68.6 | 3.05-3.24, m | | |
| 5' | 79.7 | 3.09, dt, (9.7, 4.0) | 79.7 | 3.05-3.24, m | | |
| 6' | 60.2 | 3.50, m | 60.1 | 3.48, m | | |
| Benzylic moiety | | | | | | |
| 1 | 154.4 | | 154.7 | | 156.0 | |
| 2 | 117.4 | 7.00, d, (8.7) | 115.8 | 6.75, d, (8.5) | 116.6 | 7.07, d, (8.7) |
| 3 | 129.3 | 7.18, d, (8.7) | 129.3 | 7.11, d, (8.5) | 128.4 | 7.25, d, (8.7) |
| 4 | 130.4 | | 126.6 | | 128.2 | |
| 5 | 129.3 | 7.18, d, (8.7) | 129.3 | 7.11, d, (8.5) | 128.4 | 7.25, d, (8.7) |
| 6 | 117.4 | 7.00, d, (8.7) | 115.8 | 6.75, d, (8.5) | 116.6 | 7.07, d, (8.7) |
| 7 | 37.2 | 3.86, s | 37.4 | 3.89, s | 48.2 | 4.66, s |
| others | | | | | | |
| 0 | 154.6 | | 162.8 | | 128.3 | |
| Rhamnose moiety | | | | | | |
| 1'' | 98.1 | 5.41, d, (1.7) | | | 97.5 | 5.53, d, (1.7) |
| 2'' | 69.9 | 4.03, dd, (1.7, 3.5) | | | 70.6 | 4.16, dd (1.7, 3.5) |
| 3'' | 70.0 | 3.86, dd, (3.5, 7.0) | | | 71.5 | 3.99, dd (3.5, 9.3) |
| 4'' | 70.2 | 3.38, t, (9.7) | | | 73.4 | 3.56, t, (9.4) |
| 5'' | 69.3 | 3.67, m | | | 68.6 | 3.77, dq (6.3, 9.4) |
| 6'' | 16.6 | 1.09, d, (6.3) | | | 17.5 | 1.29, d, (6.3) |

Table 2Concentration of glucosinolates in seeds and/or leaves of different *Noccaea caerulea* accessions.

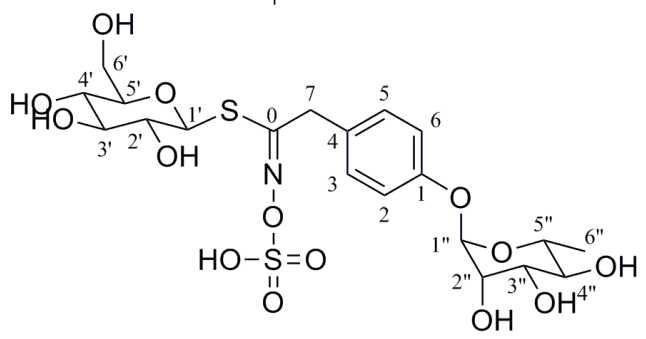
| accession | location | conc. Glucosinolates in seeds ($\mu\text{mol/g}$ dry weight) | | | conc. Glucosinolates in leaves ($\mu\text{mol/g}$ dry weight) | | |
|---------------------------------------|-----------------------------|---|-------------|--------------|--|-----------|------------|
| | | sinigrin | sinalbin | moringin | sinigrin | sinalbin | moringin |
| VDV: Valle de Varrados (France) | 42°46'34.56"N 0°49'39.09"E | 68.4 (12.7) | 11.4 (2.32) | 111.0 (12.8) | | | |
| PON: Pontaut (Spain) | 42°50' 3.05"N 0°44'0.23"E | 62.7 (1.7) | 8.9 (0.5) | 70.5 (4.1) | | | |
| LE: Lellingen (Luxembourg) | 49°59'28.63"N 6°0'10.76"E | 67.3 (3.7) | 17.7 (1.2) | 51.9 (5.7) | <0.1 | 4.7 | 34.6 |
| NAV: Navacelles (France) | 43°53'15.1"N 3°30'32.8"E | 27.9 (1.6) | 6.2 (0.7) | 67.3 (2.5) | | | |
| LAN: Lanastosa (Spain) | 43°13'51.86"N 3°25'57.69"W | 44.2 (5.8) | 6.7 (1.6) | 13.3 (4.0) | | | |
| Ardeche (France) | 44°48'28.00"N 04°20'44.80"E | 59.2 (8.2) | 19.1 (3.4) | 41.5 (8.2) | | | |
| GA: Ganges (France) | 43°56'11.2"N 3°40'17.2"E | 42.4 (5.7) | 8.9 (1.5) | 24.1 (2.9) | <0.1 | 2.1 (0.6) | 12.8 (1.9) |
| SF: Saint Felix de Pallières (France) | 44°2'40.03"N 3°56'18.05"E | | | | <0.1 | 0.4 (0.1) | 6.7 (1.7) |
| CDM: Col du Mas de l'Ayre (France) | 44°25'47.52"N 3°59'14.07"E | | | | <0.1 | 0.4 (0.1) | 6.7 (1.5) |
| MP: Monte Prinzerza (Italy) | 44°38'32.56"N 10° 5'4.87"E | | | | <0.1 | 0.1 | 0.8 |
| PB: Plombières (Belgium) | 50°44'4.31"N 5°57'49.52"E | | | | <0.1 | 4.1 | <0.1 |
| LC: La Calamine (Belgium) | 50°42'38.78"N 6° 0'37.42"E | 43.9 (6.1) | 78.4 (6.3) | <0.1 | <0.1 | 4.8 | <0.1 |
| PR: Prayon (Belgium) | 50°34'52.54"N 5°40'0.64"E | 44.9 (1.7) | 83.0 (12.3) | <0.1 | <0.1 | 15.1 | <0.1 |
| Moringa (india/lab) | 28°58.9'N 77°16.8'E | | | | <0.1 | <0.1 | 5.6 |

2



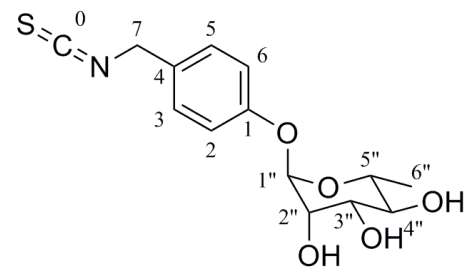
sulfatase treatment

1



myrosinase treatment

3



4

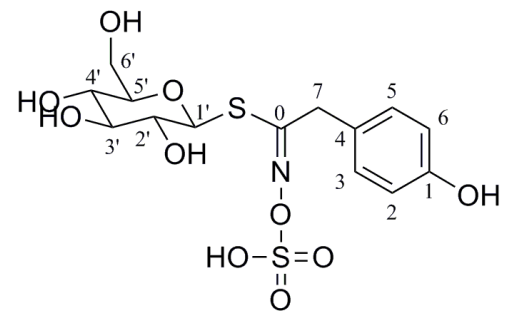


Figure 1: Schematic drawing of the reaction products formed after incubation of the 4- α -rhamnosyloxy benzyl glucosinolate (glucomoringin) with sulfatase or myrosinase. **1**: unknown glucosinolate (now identified as 4- α -rhamnosyloxy benzyl glucosinolate) **2**: desulfated glucosinolate (with carbon atom numbering) **3**: corresponding isothiocyanate, **4**. Sinalbin (sulfated).



Figure 2: Map of the locations of the analysed *Nocca caerulea* accessions. The accessions indicated with a yellow pin contain glucomoringin, those indicated with a red pin do not contain glucomoringin.

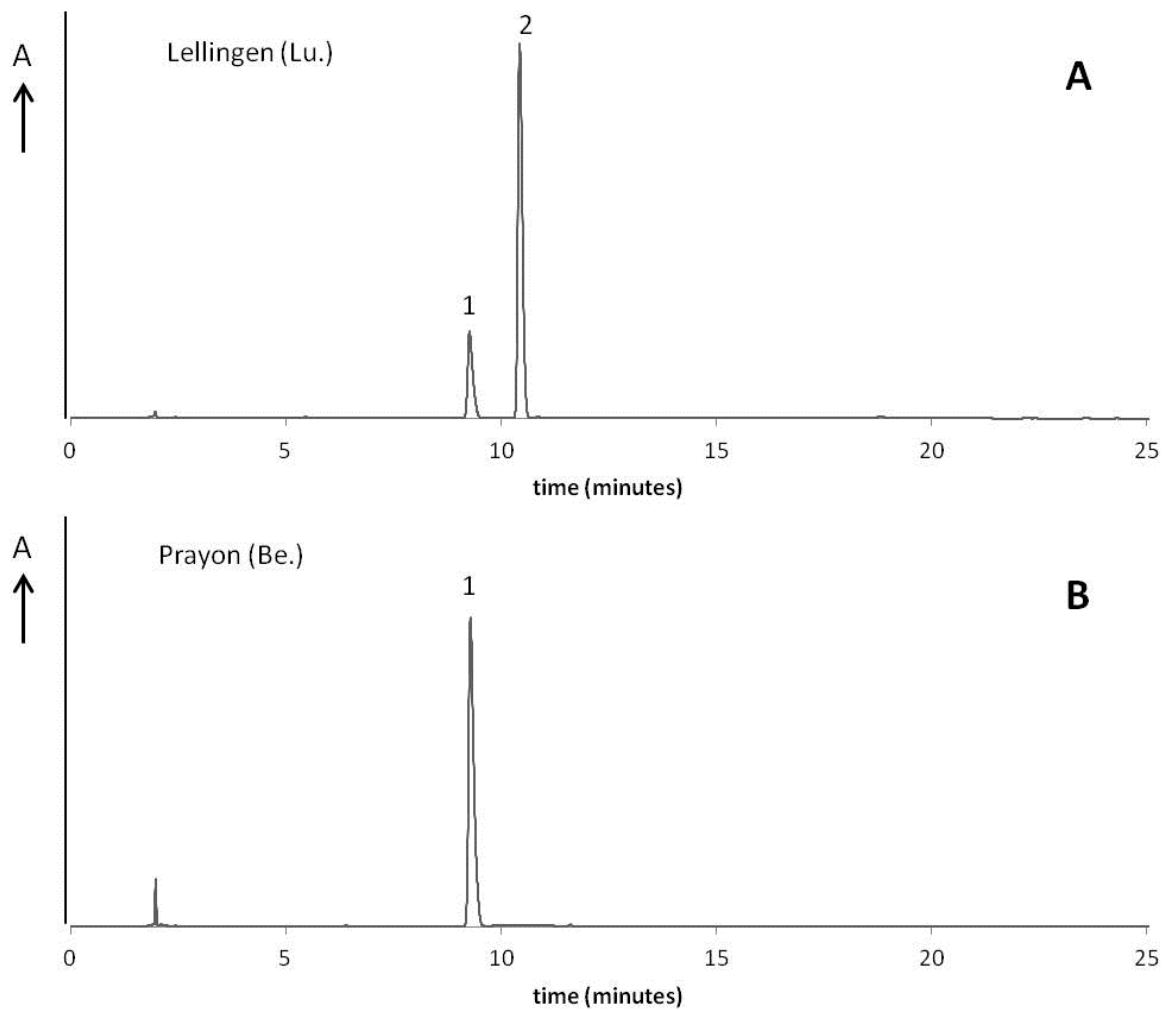


Figure S1: Selected HPLC chromatograms (229 nm) of *Noccaea caerulea* glucosinolate extracts of leaves of plants originating from two different populations; A. Lellingen, B. Prayon. Peaks: 1 = desulfosinalbin, 2 = desulfoglucomoringin.

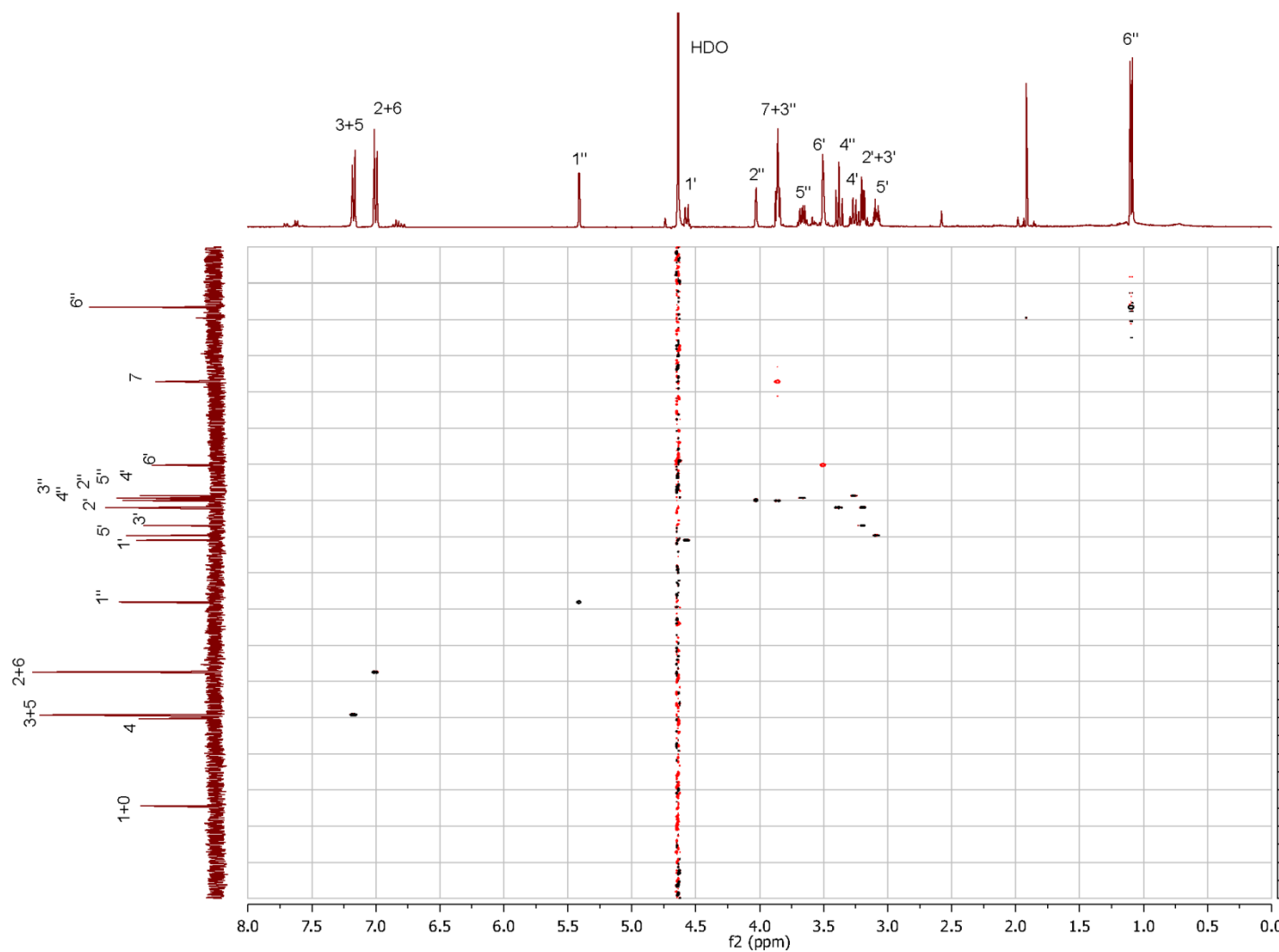


Figure S2: gHSQCAD spectrum of the unknown glucosinolate (here identified as glucomoringin).

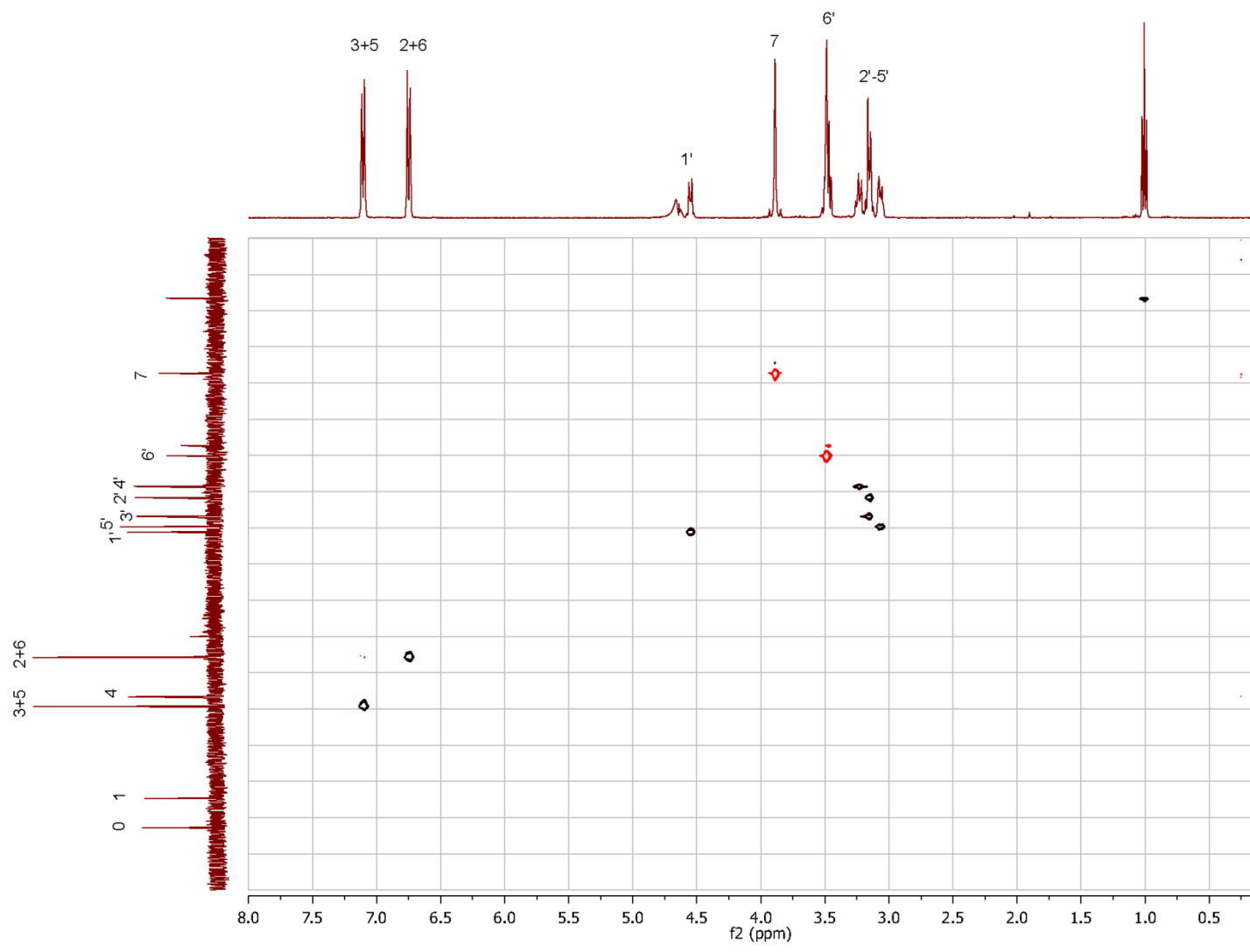


Figure S3: gHSQCAD spectrum of Sinalbin

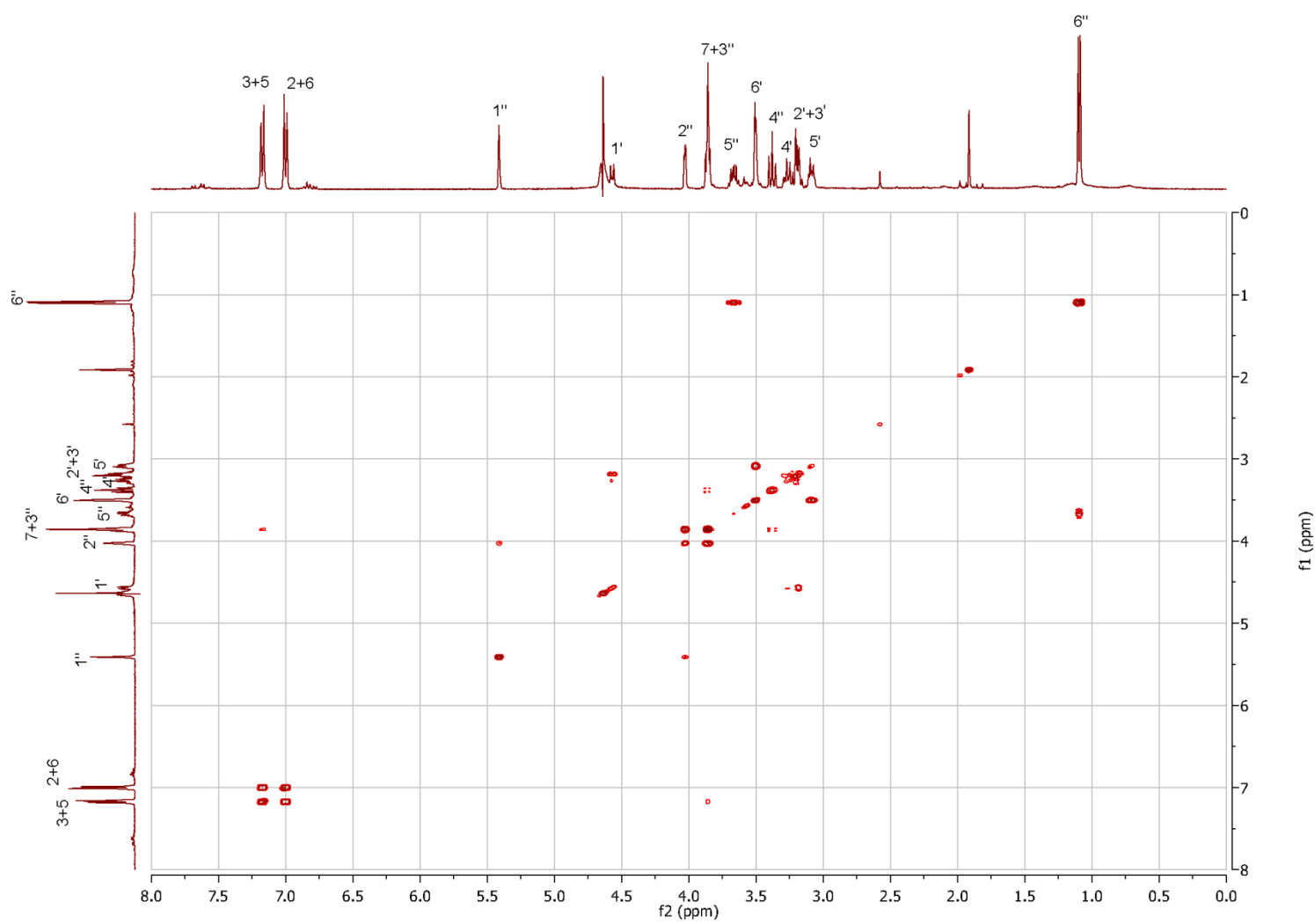


Figure S4: gCOSY NMR spectrum of the unknown glucosinolate (here identified as glucomoringin).