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Carbon-13 Cross-Polarization Magic-Angle Spinning Nuclear Magnetic Resonance (CPMAS NMR) for Measuring Proanthocyanidin Content and Procyanidin to Prodelphinidin Ratio in Sainfoin (Onobrychis vicifolia) Tissues

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ABSTRACT

A procedure based on $^{13}$C CPMAS NMR was developed to study procyanidins (PCs) and prodelphinidins (PDs) directly in milled sainfoin plant tissues. Blackcurrant and *Tilia* samples enabled reference spectra of purified proanthocyanidin (PA) fractions, crude extracts and milled plant tissues, with characteristic resonances at 155, 144 and 132 ppm. PC/PD ratios were estimated from the $I_{132}/I_{155}$ intensity ratio and differed by 2.5 to 5.9% compared to thiolysis data. Normalization to the 155 ppm signal intensity from reference spectra enabled analysis of PA contents with an error of ca 8 g PAs/100 g plant tissue. The procedure estimates the lignin contribution and allows for a correction of the PA content. In six sainfoin accessions, estimated PA contents, were 1.6- to 20.8-fold higher than the thiolysis and 1.4- to 2.6-fold higher than the HCl-butanol-acetone results. Method differences may reflect the presence of unextractable, possibly high molecular weight PAs in sainfoin.

KEYWORDS: condensed tannins; CPMAS NMR; thiolysis; HCl-butanol-acetone assay; extractable and non-extractable proanthocyanidins; sainfoin;
INTRODUCTION

Given the beneficial effects of proanthocyanidins (PAs) on human and animal nutrition and health,¹,² there is a need for analytical methods capable of providing information on the true PA content and their composition in plants. However, their analysis is challenging as many plants synthesize complex polymeric mixtures of PAs (Figure 1).³,⁴ Extractable PAs are typically analyzed after degradation in the presence of nucleophiles followed by chromatographic separation of the reaction products and mass spectrometric analysis; this gives data on PA content, flavan-3-ol subunit composition, mean degree of polymerization (mDP) and procyanidin/prodelphinidin (PC/PD) and cis/trans flavan-3-ol ratios.⁵⁻⁸ Extraction, however, is often incomplete as many plants also contain large quantities of unextractable PAs, which can lead to underestimation of total contents.⁹⁻¹¹ Therefore, in planta PA depolymerization with HCl-butanol-acetone, thiolysis or phloroglucinolysis has been used to estimate extractable plus unextractable PAs; but, incomplete reaction, side reactions or degradation-resistant PAs continue to pose challenges.³,⁴,¹²⁻¹⁴ Of particular concern is the fact that the HCl-butanol-acetone reagent often, but not always, gives higher PA contents than thiolytic degradation;¹⁵,¹⁶ although opposite results have also been found with galloylated PAs (unpublished results). Questions remain, therefore, about the true PA contents in plants and foods. Currently, little is known about the quantities and composition of unextractable PAs and alternative methods deserve evaluation.¹⁰,¹⁷

In contrast to the above degradation methods, nuclear magnetic resonance (NMR) has the potential to provide direct evidence and accurate results for total, i.e. extractable plus unextractable, PAs. Solution-state ¹³C NMR spectroscopy has been used to determine the relative purities,¹⁸,¹⁹ PC/PD¹⁸⁻²¹ and cis/trans flavan-3-ol ratios¹⁸,²⁰,²¹ and mDP values of extracted PAs.²⁰,²¹ Whilst two dimensional gel-state NMR proved useful for assessing the presence and composition of PAs in Lotus tissues,²² our proof-of-concept study showed that
gel NMR spectra of sainfoin (*Onobrychis viciifolia*) tissues had signal resolution and matrix interference problems.

Solid-state NMR experiments have been used for the direct analysis of plant materials and require neither extraction nor derivatization of PAs. Cross-polarization magic-angle spinning (CPMAS) NMR techniques have previously been used to evaluate the efficiency of PA extraction in pine barks, softwood and barks from quebracho and humus as well as to characterize PAs in dietary fiber samples. In addition, differences in fingerprint spectra were used to distinguish between PA transformations of leather tannings and during fungal degradation. However, significant problems with poor resolution and frequent overlaps of multiple signals prevented any in depth quantitative estimates of the PA content and PC/PD ratios.

Sainfoin is a PA-containing forage legume, which is of interest for its nutraceutical and environmental properties. It engenders high voluntary intakes, has excellent nutritive value, lowers *in vitro* greenhouse gas emissions, prevents bloat in ruminants, lowers gastrointestinal nematodes, and can enhance the unsaturated fatty acid composition of meat and milk products. The work presented here used six sainfoin accessions as a case study for complex PA mixtures. Such complexity may have been the main reason for the observed discrepancies between the thiolysis and HCl-butanol-acetone assay results and still requires detailed experimental confirmation. Therefore, the aim of this study was to develop a procedure based on a $^{13}$C CPMAS NMR technique for estimating PA content and composition directly, without chemical modification in different sainfoin accessions and to compare the results to thiolysis and HCl-butanol-acetone data.

**MATERIALS AND METHODS**
**Plant Samples.** Blackcurrant (*Ribes nigrum*) leaves were collected from Hildred PYO farm (Goring-on-Thames, UK). *Tilia* (*Tilia* spp.) flowers were purchased from Flos (Mokrsko, Poland) as described by Ropiak et al. Sainfoin (*Onobrychis vicifolia*) leaves and stems, from six accessions, were provided by the National Institute of Agricultural Botany (Cambridge, UK). Blackcurrant leaves and sainfoin plants were lyophilized or air-dried and subsequently ground to pass a 1 mm sieve and ball-milled. Fine powders of ball-milled blackcurrant leaves and *Tilia* flowers were also mixed on a weight basis (100/0, 70/30, 50/50, 30/70 and 0/100) for NMR analysis and the development of the procedure to elucidate PC/PD ratios.

**Chemicals and Reagents.** Hydrochloric acid (36%), formic acid, acetic acid, butan-1-ol, HPLC-grade acetone, HPLC-grade methanol, HPLC-grade dichloromethane, HPLC-grade hexane, HPLC-grade acetonitrile and ammonium chloride were purchased from Fisher Scientific Ltd (Loughborough, UK). (±) – Dihydroquercetin (98%) was from Apin Chemicals (Abingdon, UK). Benzyl mercaptan and acetone-\(d_6\) (99.9%) were supplied by Sigma-Aldrich (Poole, UK). Deuterium oxide (D\(_2\)O) was from CK Isotopes (Ibstock, UK). Sephadex LH-20 was purchased from GE Healthcare (Little Chalfont, UK). Deionized water was purified in a Milli-Q system (Millipore, Watford, UK).

**Proanthocyanidin Extraction and Purification.** Finely ground plant tissue (50 g) was weighed into a conical flask. Acetone/water (500 mL, 7:3, v/v) was added and the mixture was vigorously stirred for 1 h. The mixture was filtered under vacuum. The filtrate was further extracted with dichloromethane (250 mL) to remove lipids and chlorophyll. Polyphenols were concentrated in the aqueous phase with a rotary evaporator below 37.5 °C to yield the “crude extract”. 
This crude extract was diluted with deionized water (2 L) and filtered under vacuum. The filtrate was loaded on a 400 x 65 mm i.d., glass column which had been packed with 70 x 65 mm i.d., Sephadex LH-20 resin and was equipped with a sintered-glass frit. The column was then washed with deionized water (2 L) to elute sugars and low molecular mass phenolics. Gravity elution with acetone/water (3:7, 1:1 and 8:2, v/v; 1 L each) yielded three PA fractions from each plant source; the first 200 mL of each eluate was discarded and PA fractions 1, 2 and 3 were collected in the next 500, 300, and 300 mL, respectively. The organic solvent was removed on a rotary evaporator below 37.5 °C and the PA fractions were frozen, freeze-dried and stored at -20 °C. Fractions 2 that eluted with 1:1 acetone/water were designated as “purified PA fractions”. Various aliquots from the purified blackcurrant and Tilia PA fractions (152:0, 36:83, 45:38, 63:25, 0:157 mg) were also mixed, in order to provide a range of nominal PC/PD values of 0/100, 30/70, 50/50, 70/30 and 100/0, for NMR analysis. The calculated PC and PD contents within purified PA fraction (in g PCs or PDs/100 g purified PA fraction) were based on the PC and PD contents from thiolysis (Tilia: 97.4 g PCs/100 g of purified PA fraction and blackcurrant: 94.9 g PDs/100 g of purified PA fraction).

Thiolysis of Purified Proanthocyanidins. Thiolysis reactions on purified PA fractions that eluted with acetone/water (1:1 v/v) were performed according to Novobilský et al. Purified PA fractions (4 mg) were weighed into a 10 mL screw-capped vial, dissolved in methanol (1.5 mL) and acidified with HCl (0.5 mL, 3.3% in methanol, v/v). This was followed by addition of benzyl mercaptan (50 µL) and the reaction mixture was stirred in a water-bath for 1 h at 40 °C. The reaction was quenched by adding ultrapure water (2.5 mL) to the mixture at room temperature. The analysis and quantification of thiolysis reaction products were
performed with reverse-phase HPLC according to Gea et al. The operating conditions and parameters of HPLC analysis were described by Williams et al.

**In planta Thiolyis of Sainfoin Proanthocyanidins.** Milled freeze-dried sainfoin plant material (200 mg) was weighed into a 10 mL screw-capped vial, dissolved in methanol (2 mL) and acidified with HCl (1 mL, 3.3% in methanol, v/v). This was followed by addition of benzyl mercaptan (100 µL) and the mixture was stirred in a water-bath for 1 h at 40 °C. The reaction was stopped by transferring the vials into an ice-bath and 1% formic acid in ultrapure water (9 mL) was added at room temperature. The samples were centrifuged (3000 rpm, 5 min) and 1 mL of the mixtures added to HPLC vials for LC-MS analysis.

Samples (5 µL) were injected into an 1100 series HPLC system (Agilent Technologies, Stockport, UK) connected to a 150 x 3 mm i.d., 5 µm, ACE super C\textsubscript{18} column (Hichrom, Theale, UK) fitted to a 150 x 3 mm i.d., 5 µm, ACE EXCEL Ultra-Inert UHPLC compatible guard cartridge (Hichrom, Theale, UK). The column temperature was set at 60 °C. The HPLC system consisted of a G1379A degasser, a G1312A binary pump, a G1313A ALS autoinjector, a G1314A VWD UV detector and a G1316A column oven and an 1100 series MSD API-ES mass spectrometer (Agilent Technologies, Waldbronn, Germany). Data were acquired and processed with ChemStation software (version A 10.01 Rev. B.01.03).

The flow rate was 0.4 mL/min using formic acid (1%) in water containing 100 mg/L ammonium chloride (solvent A) and HPLC-grade acetonitrile (solvent B). The thiolyis reaction products eluted with the following gradient: 0-7 min, 2.5% B; 7-15 min, 2.5-5% B; 15-22 min, 5-10% B; 22-40 min, 10-40% B; 40-45 min, 40-100% B; 45-49, 100-2.5% B; 49-60, 2.5% B. Mass spectra were recorded in the negative ionization scan mode from m/z 100-1000. The mass spectrometer operating conditions were as follows: 3000 V for capillary voltage, nebulizer gas pressure at 35 psi, drying gas at 12 mL/min and dry heater temperature
at 350 °C. Flavan-3-ols and their benzyl mercaptan adducts were identified by their retention
times, ultraviolet-visible (UV) spectra recorded at 280 nm and molecular masses and they
were quantitated using published response factors against dihydroquercetin.\textsuperscript{5,8}

\textit{In planta Sainfoin Proanthocyanidin Analysis with the HCl-Butanol-Acetone Assay.} The
HCl-butanol-acetone assay followed the procedure of Grabber et al.\textsuperscript{22} with minor
modifications. Briefly, lyophilized sainfoin plant tissue was weighed (10 mg) into a 10 mL
screw-capped vial. The reagent mixture was prepared by mixing ammonium ferric sulfate
(150 mg) in ultrapure water (3.3 mL), hydrochloric acid (12 M, 5 mL), butan-1-ol (42 mL)
and acetone (50 mL). An aliquot of the reagent (10 mL) was added to the sainfoin sample.
The sample was left at room temperature (1 h) to check for the presence of flavan-4-ols or
flavan-3,4-diols as these generate false positives.\textsuperscript{43} Tubes were then stirred and heated at 70
°C for 2.5 h in the dark. Samples were left to cool to room temperature and centrifuged (3000
rpm, 1 min). Absorbance of the supernatant was recorded by scanning from 450-650 nm
using a V530 spectrophotometer (Jasco, Great Dunmow, UK). The HCl-butanol-acetone
reagent was used as a blank and all samples were run in duplicate. A sainfoin sample of
known PA content and composition was used for quality control purposes.

Analysis of Proanthocyanidins by \textsuperscript{13}C Cross-Polarization Magic-Angle Spinning

\textbf{Nuclear Magnetic Resonance.} The solid-state CPMAS NMR spectra were recorded on a
Bruker Avance III 500 MHz spectrometer (Bruker Biopsin, Rheinstetten, Germany) operating
at carbon-13 Larmor frequency of 125.78 MHz (11.75T). Crude extracts and purified PA
fractions from \textit{Tilia} and blackcurrant and ball-milled \textit{Tilia}, blackcurrant and sainfoin plant
tissue samples (~80 mg) were compressed into standard 4 mm zirconia rotors prior to
analysis. A standard bore Bruker MAS probe was used and rotors were spun at 10 kHz rate.
The proton 90° pulse width was 3.7 µs at a power level of 32 W. The variable amplitude CP
ramp (90-100) was used with a contact time of 1 ms. In total, 4096 signal transients were
averaged into each spectrum with a 6 s relaxation delay at ambient temperature. All spectra
were referenced to an external adamantane signal (at 38.5 ppm with respect to TMS) as a
secondary reference. Dipolar dephasing experiments were run with identical parameters to
$^{13}$C CPMAS NMR experiments and dephasing filter time (tdd) of 45 µs was optimized to
attenuate the signals of all protonated carbons.

To accurately extract intensity of all signals, each experimental spectrum was fitted
with a calculated spectrum that was constructed from the sum of signals with a Gaussian line
shape. Numbers and positions of signals in the calculated spectrum corresponded directly to
the number and positions of signals in the experimental spectrum. The width, amplitude and
position of each signal in the calculated spectrum was optimized by minimizing the
discrepancy between the experimental and calculated spectra using an “in house” written
Matlab routine based on the least square method. A number of initial parameters were tested
and the sets with the smallest discrepancy led to final results with a narrow range of the
fitting error. Once the width and the amplitude of the fitted signal at 155 ppm were known,
the intensity was calculated.

The PA content of sainfoin plant tissue samples was estimated by comparing the
signal intensity of the reference signal $I_R$ at 155 ppm (from *Tilia* with 95 g PAs/100 g of
purified PA fraction, determined by thiolysis) with the intensity $I_S$ of the 155 ppm signal in
the investigated sample.

The calculation of experimental error involved three factors: (i) the non-strictly
quantitative character of the solid-state NMR experiments because of the cross-polarization
signal enhancement (and possibly the miscalibration of the relaxation parameters); (ii) the
small variation in setting hardware parameters, such as tuning and matching of the probe; and
(iii) the numerical fitting of the spectra.

The recording of 2-3 spectra of the same sample with slightly different settings regarding the first two factors (including contact time and relaxation delay) resulted in an error of ca 2%. The discrepancies of the spectra acquired with different parameters introduced an additional error of ca 2% after the fitting process. This accounted for a total error of ca 4% for the intensity measurements. The PA content was calculated by comparing the reference signal intensity to the intensity of the sample of interest. Therefore, the total experimental error (ca 8% or 8 g PAs/100 g of plant tissue) derived from the sum of the intensity errors between reference and sample intensity.

The influence of the spinning sidebands on the signal intensity was assessed by comparing the 155 ppm signal intensity recorded at 10, 12 and 15 kHz spinning rates for both crude extract and sainfoin plant tissue samples. Only small differences of less than 1%, well within the experimental error were observed. At 10 kHz spinning rate, the spinning sidebands of the signal at 155 ppm were not fully removed and decreased the intensity by ca 12%. However, the identical anisotropy of the chemical shift of C5, C7 and C8a carbons in purified PA fraction and plant tissue PAs is expected (due to the same molecular structure) and by normalizing the intensity of the 155 ppm signal from the sample of interest to the high purity reference with 95 g PAs/100 g of purified PA fraction from *Tilia*, the contribution of these factors to the intensity error could be neglected.

However, it should be noted that $^{13}$C CPMAS NMR is not strictly quantitative and, therefore, it is unlikely but not impossible that the discrepancies could have exceeded to a small extent the range of the estimated experimental error.

RESULTS AND DISCUSSION
Assignment of Signals in the Solid-State NMR Spectrum. The $^{13}$C CPMAS NMR spectra of the purified PA fractions from Tilia and blackcurrant consisted of approximately ten resolved resonances each centered between 170 and 0 ppm and showed several PA fingerprint signals (Figure 2B). Characteristic resonances at 155, 144 and 132 ppm were clearly observed in the spectra of all three sample types, i.e. purified PA fractions, crude extracts and plant tissues (Figure 3). The signal at 155 ppm originated from the non-protonated carbons C5, C7, C8a of the flavan-3-ol structure (Figure 2A). The signal at 144 ppm was from the resonance of the non-protonated carbons C3’, C5’ of blackcurrant PAs and the non-protonated carbons C3’, C4’ of Tilia PAs. The most distinctive signal at 132 ppm originated from the non-protonated carbons C1’, C4’ in blackcurrant PAs but only from the non-protonated carbon C1’ in Tilia PAs.

In order to confirm these assignments, dipolar dephasing solid-state NMR experiments were performed. The experimental parameters were optimized to detect only signals from non-protonated carbons (Figure 2B). The resulting data confirmed our initial identification and were in good agreement with spectra in reports on pecan nutshell PDs and Photinia leaf PCs. Weak signals of not completely removed spinning sidebands were detected in the 80-40 ppm region. Some very weak contributions from the protonated carbons were also visible in the 40-20 ppm region and could have been caused by a small miscalibration of parameters for the dephasing filter.

The resonances from protonated carbons were spread across the 120-0 ppm region and showed line broadening and overlap, especially for the crude extracts and milled plant tissues (Figures 2B and 3). This was likely due to interferences from other plant components such as lignin, pectin, cutin, cellulose and hemicellulose as previously reported in crude extracts and milled plant samples. However, our signal assignments of the purified PA
fractions from *Tilia* and blackcurrant were consistent with the much better resolved $^1$H-$^{13}$C HSQC solution NMR spectra of the same samples.

**Estimation of Proanthocyanidin Content in the Presence of Lignin.** Accurate quantification of PAs in plant matrices is of great interest and several techniques have been used including solid-state NMR. The region between 160-120 ppm is understood to be the most suitable despite some possible lignin or lignin-PA complex contributions especially in milled plant samples.$^{25, 44}$ In particular, Wawer et al.$^{27}$ postulated that resonances at 155 and 144 ppm in solid-state NMR dipolar dephased spectra from *Aronia* and blackcurrant fiber powders could include responses from C3, 5-OMe and C4-OH carbons of lignin. However, this interference was on a smaller scale (i.e. there were no significant changes to the signal pattern) compared to resonances below 120 ppm (Figure 3) where the appearance changed completely between purified PA fractions and milled plant tissues.

The characteristic signals at 155, 144 and 132 ppm in the spectra of the purified PA fractions could only have derived from PAs as large polymeric lignin molecules were not expected to be soluble in acetone/water.$^{27, 46}$ Our strict, two step purification process also assured that the low molecular weight phenolics or lignin fragments which may have been present in small quantities in crude extracts were predominantly removed in the purified PA fractions. The intensities of the 132 ppm signal spinning side bands at 211 ppm (left spinning side band) and 53 ppm (right spinning side band) were exactly the same (0.02% difference) which proves that there is no, or only negligible, lignin contribution to the NMR spectra of the purified PA fractions. Figure 3 shows the comparison of the normalized amplitude of these characteristic signals between purified PA fractions, crude extracts and milled plant tissues. However, it was not possible to estimate the contribution of impurities or plant matrix components to the intensity because currently there is no technique that can provide
completely accurate data on PA concentrations in the plant tissues. However, a contribution of lignin in this region of the spectrum was expected between 155 and 144 ppm as postulated, and it should affect the width of the resonances due to overlap. Therefore, measuring the change of the signal width in purified PA fractions, crude extracts and milled plant tissues could provide an estimate of lignin interference. In our case, the estimated increase in the width of the 155 and 144 ppm signals was <15% for crude extracts and <25% for plant tissues, and corresponded well to the increased intensity of the lignin signal at 53 ppm, which was especially evident for the *Tilia* plant tissues (Figure 3B).

This simple estimate showed that the lignin interference had to be taken into account as a correction to the PA content in plant samples. To test the accuracy of this assumption, the content of extractable PAs in the crude extract from blackcurrant was estimated with $^{13}$C CPMAS NMR and compared to the thiolysis result. The PA content determined by NMR without correction was 39.7 g PAs/100 g crude extract and with the 15% reduction was 33.7 g PAs/100 g crude extract (with an error of ± 8 g PAs/100 g). The latter value was in excellent agreement with the thiolysis result of 29.2 g PAs/100 g crude extract. This demonstrates that both techniques can be used for the analysis of extractable PAs.

Although this simple calculation may take into account a lignin contribution, other factors could also influence the width of the resonance line, e.g. relaxation and mobility issues can arise from large polymeric molecules and small variation in the shimming quality. Further, one needs to be aware that the thiolysis and HCl-butanol-acetone assays have their own limitations, especially when applied directly to plant tissues.

Despite problems arising from possible plant matrix interferences in the milled plant tissues, it may be concluded that the 155 ppm signal of the purified PA fractions was a suitable marker to estimate PA contents in unknown milled plant tissues. Therefore, the PA content of sainfoin plant tissue samples was estimated by comparing the intensity of a
reference signal, $I_R$, of the purified PA fraction (95 g PAs/100 g as determined by thiolysis) from *Tilia* and the intensity of the $I_S$ signal of the investigated sample. The content error was estimated to be ca 8 g PAs/100 g plant tissue.

The PA content of selected sainfoin plant tissues determined by $^{13}$C CPMAS NMR (without and with a 25% reduction for the lignin correction) was then compared to results from the thiolysis and the HCl-butanol-acetone assays (Table 1). Selected spectra are shown in Figure 4. Clearly, solid-state NMR consistently gave the largest PA content in all sainfoin plant tissue samples and thiolysis gave the smallest. Most HCl-butanol-acetone values were higher than the thiolysis data and accounted for ca 50% of the NMR data after lignin correction. The results varied greatly between the sainfoin accessions for all methods, but it is worth pointing out that the three Zeus samples (S1, S2 and S6) gave consistent results with the NMR and HCl-butanol-acetone but not the thiolysis technique.

The above results illustrate the complexity of determining PA content directly in plants. Our NMR results can be regarded as an upper limit of PA content compared to the other techniques (Table 1) and is more likely to measure previously undetected unextractable PAs because NMR does not require any chemical modification of the sample. Solid-state NMR has already proved useful for estimating differences in PA contents within species, i.e. for samples with similar PA compositions. Our approach to account for a lignin contribution could improve the way NMR could be used for PA measurements more generally, including for the first time full quantitative elucidation. The validity of our approach is illustrated by the analysis of the amplitude of the lignin signal at 53 ppm for samples S1, S3 and S4 (Figure 4). The strongest lignin signal was detected in the S1 sample and the weakest and approximately equal signals in the S3 and S4 samples. Had lignin interference been the predominant factor in the measurement of PA content then S1 should have had more PAs than the S3 and S4 samples. As evident from the 155 and 145 ppm
signals this was not the case as the S3 sample with the smallest lignin contribution had the largest PA content.

There are several possible reasons for the differences between the NMR and the thiolysis or HCl-butanol-acetone data. Whilst our results account for a lignin contribution at 155 ppm on the basis of the reference *Tilia* and blackcurrant samples, they do not rule out contributions from other matrix components in the investigated sainfoin plant tissues.\textsuperscript{24, 27} Secondly, solid-state NMR experiments are not strictly quantitative because the efficiency of the cross-polarization depends on proton abundance in the neighboring environment, the molecular dynamics and the relaxation behavior for improving the $^{13}$C signal.\textsuperscript{25, 28} These factors are difficult to fully control in NMR experiments and are unlikely to be the same for all carbons, and will also vary between different plant matrices. Therefore, it is possible that the proton environment of the more complex and possibly larger sainfoin PAs (mDP of 12 to 27) could have resulted in a larger signal enhancement (and consequently PA content) compared to the purified *Tilia* PAs (mDP of 8), but it is unlikely that this factor alone increased the NMR signal by 100% (Table 1).

Another explanation could arise from the limitations of the thiolysis and HCl-butanol-acetone assays in quantitating PAs directly in plant tissues.\textsuperscript{3, 4} It has been shown that longer reaction times during thiolysis with benzyl mercaptan resulted in considerably higher concentrations.\textsuperscript{5} In addition, the existence of thiolysis-resistant PAs has also been reported.\textsuperscript{9} The previous versions of the HCl-butanol assay often underestimated total PA content when applied directly to plant or extracted residue samples, possibly because of incomplete interaction with bound PAs.\textsuperscript{12} However, the present HCl-butanol-acetone assay used a recent modification by Grabber et al.,\textsuperscript{22} who demonstrated that inclusion of acetone achieved complete degradation of unextractable PAs from two *Lotus* species. Nevertheless, it is possible that sainfoin may have more diverse PA mixtures than *Lotus*,\textsuperscript{48} which could explain
the observed differences between the NMR, thiolysis and HCl-butanol-acetone results.

Overall, the important point here is that the NMR results suggested that NMR allows accounting for all extractable and unextractable PAs, whereas the thiolysis and HCl-butanol-acetone assays may be underestimating PA contents.

**Estimation of Mean Degree of Polymerization.** The mean degree of polymerization (mDP) can be obtained by integrating the C3 signals from PA extension units at 73 ppm and the corresponding signals of PA terminal units at 67 ppm in solution NMR. However, significant signal overlap from various plant constituents in milled plant tissues and consequently poor resolution of that region in the $^{13}$C CPMAS NMR spectra (Figure 3) prevented estimation of mDP values.

**Estimation of procyanidin/prodelphinidin ratios.** The procedure to estimate PC to PD ratios within PAs was based on their different hydroxylation patterns, which were clearly distinguishable in the 140-100 ppm region of the $^{13}$C CPMAS NMR spectra (Figure 2B). The resonance at 132 ppm was assigned to C1´ and C4´ carbons of PDs from blackcurrant, but in *Tilia* PCs this resonance was assigned only to C1´ carbon and had only half the intensity of the PDs signal. This difference in the 132 ppm signal intensity directly reflected the PD and PC composition because the PA contents were comparable (i.e. blackcurrant 87 g PAs/100 g of purified PA fraction; and *Tilia* 95 g PAs/100 g of purified PA fraction); the molar percentages obtained by thiolysis were 94.9% PDs and 97.4% PCs for the purified PA fractions, from blackcurrant and *Tilia* respectively.

This observation was validated by thiolysis and $^1$H-$^{13}$C HSQC solution NMR experiments on artificial samples with the nominal PC/PD values equal to 100/0, 70/30,
50/50, 30/70 and 0/100, prepared by mixing purified PA fractions from *Tilia* and 
blackcurrant. 

However, the solution NMR methods, such as used by Czochanska et al.\textsuperscript{20} and Zeller 
et al.,\textsuperscript{23} to estimate PC/PD ratios, which rely on the high resolution of the solution NMR 
spectra could not be transferred directly to the solid-state NMR experiments. The 
experimental resolution of the recorded solid-state NMR spectra was in the order of 12.5 Hz 
(or 0.1 ppm) but the typical linewidth was ca 550 Hz (or 4.5 ppm), as expected for large 
polymeric molecules. In addition to that, the spectra of milled plant tissues show significant 
overlap with other signals from the plant matrix below 110 ppm, which prevents any 
meaningful PC/PD analysis (Figures 3 and 5A). Therefore, it was impossible to calculate 
PC/PD ratio on the basis of the C2’ - C6’ carbons as used in solution NMR experiments. 
The experiment on the artificial mixtures of purified PA fractions with nominal 
PC/PD values was repeated using the solid-state NMR technique. The intensity ratios 
between signals at 132 and 155 ppm were selected as a calibration standard (to minimize the 
experimental errors) because they were expected to vary linearly between a value of 1/3 or ca 
0.33 for pure PCs, and 2/3 or ca 0.67 for pure PDs reflecting the number of carbons 
contributing to each signal (Figures 2B and 5A). The resulting $I_{132}/I_{155}$ experimental ratios 
were within the theoretical limits (Figure 5B) and showed excellent linear dependence when 
plotted as a function of the nominal PC/PD values. 

However, to extend the procedure to estimate the unknown PC/PD ratios of milled 
sainfoin tissues, it was necessary to repeat the experiment on the artificial mixtures but this 
time these were prepared on a plant weight (mg) basis using the milled *Tilia* and blackcurrant 
tissues. This allowed us to assess and overcome several problems, such as imprecisions in the 
nominal PC/PD values and plant matrix interference, which varied between different plants 
and could obscure any direct calculation of the unknown PC/PD ratio using only milled *Tilia*
or blackcurrant plant tissues. It was assumed that the nominal PC/PD values (again equal to 100/0, 70/30, 50/50, 30/70 and 0/100) in these plant tissues would be the same as in the purified PA fraction mixtures. A similar linear dependence of the $I_{132}/I_{155}$ on the nominal PC/PD values was indeed observed for the milled plant tissues compared to the purified PA fraction mixtures (Figure 5B). The large offset between both curves shows that there is a possible interference of the plant matrix. However, both curves showed a similar slope, which confirmed that extraction and purification had no effect on the PC/PD ratio in *Tilia* and blackcurrant mixtures, and that the matrix effect on intensity was almost uniform across these PC/PD ratios. Therefore, the gradient of the $I_{132}/I_{155}$ intensity ratio as a function of nominal PC/PD values could be regarded as a universal tool for setting up a calibration curve to estimate unknown PC/PD ratios in other plant samples such as sainfoin.

Table 1 shows a comparison between the PC content in sainfoin samples from the $^{13}$C CPMAS NMR spectra and the thiolysis data. It can be seen that NMR estimated much higher PC contents in purified PA fractions than thiolysis in four of the eight samples, but had a rather low accuracy with an estimated experimental error of ±16 mg PCs/100 mg purified PA fraction. The biggest contributor to the estimated error was the uncertainty in the fitting of the 132 ppm signal, which overlapped with other signals from the plant matrices (Figures 4 and 6A). This was not surprising given the large offset already observed for the $I_{132}/I_{155}$ intensity ratio dependence on nominal PC/PD values between purified PA fractions and milled plant tissues (Figure 5B). However, the degree and pattern of overlap was much larger and variable in the sainfoin compared to the reference *Tilia* and blackcurrant plant tissues (Figure 5A). This problem was particularly evident when comparing the NMR spectra of the S3 and S1 tissues (Figure 4); and was also highlighted in Figure 6, which compared the NMR spectra of sainfoin leaves and stems. Obviously, it was only possible in the case of the fairly resolved structure to extract the correct intensity of the signal at 132 ppm from the spectrum.
Therefore, a better resolution of the NMR spectra will be required to improve the accuracy of
the estimated PC molar percentage or PC/PD ratio beyond a qualitative assessment.

However, the comparison of sainfoin leaf and stem spectra also demonstrated the
potential of the $^{13}$C CPMAS NMR approach to gain a rapid qualitative assessment (Figure 6).
The stem spectra showed a very different pattern of resonances in the region between 160 -
120 ppm. This was most likely due to poorer resolution of signals at 144 and 132 ppm in
stems, which tend to have higher lignin content than leaves, and also corresponded to lower
PA content in stems than leaves$^{37}$ and a greater contribution of plant matrix interferences.

This study reported a first attempt to develop a procedure based on $^{13}$C CPMAS NMR
technique for the direct screening of PAs in sainfoin plant tissues. By using contrasting
signature PA spectra from blackcurrant and *Tilia* reference samples, a calibration for PAs was
developed based on signal intensities at 155 and 132 ppm. There were surprisingly large
differences between the PA content obtained by the thiolysis or HCl-butanol-acetone assays
and $^{13}$C CPMAS NMR results. These differences could be due to interfering plant matrix
components, size and mobility differences in reference and sainfoin PAs, and/or presence of
unextractable PAs in sainfoin plant tissues. It is very unlikely that various sizes and mobility
differences would influence the cross-polarization enhancement to such an extent that it could
account for the observed PA content differences. A simple correction to account for the
matrix interference based on the linewidth difference between the purified PA fraction and
milled plant tissue has been used. However, even with the applied correction the differences
between the $^{13}$C CPMAS NMR and thiolysis or HCl-butanol assays indicated that the two
degradation methods may not detect the entire PA present in some plants. Therefore, only $^{13}$C
CPMAS NMR provided full information on both extractable and unextractable PAs in
sainfoin plant tissues.
In conclusion, this newly developed procedure based on \(^{13}\)C CPMAS NMR technique has proved to be a useful tool for estimating the upper limit of PA content directly in whole plants and could be of interest for probing the bioactivities of unextractable PAs.

**ABBREVIATIONS USED:** PA, proanthocyanidin; PC, procyanidin; PD, Prodelphinidin; \(^{13}\)C CPMAS NMR, carbon-13 cross-polarization magic-angle spinning nuclear magnetic resonance; mDP, mean degree of polymerization;

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**SUPPORTING INFORMATION DESCRIPTION:** \(^1\)H-\(^{13}\)C Heteronuclear Single Quantum Coherence (HSQC) Nuclear Magnetic Resonance Analysis of the Purified PA Fraction Mixtures; Chromatogram of the Thiolysis Reaction Products from the Blackcurrant Purified PA Fraction; Chromatogram of the Thiolysis Reaction Products from the *Tilia* Purified PA Fraction.
References


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procyanidin/prodelphinidin and cis/trans-flavan-3-ol ratios of condensed tannin samples:


(42) Williams, A. R.; Fryganas, C.; Ramsay, A.; Mueller-Harvey, I.; Thamsborg, S. M.


FIGURE CAPTIONS:

**Figure 1.** A generic proanthocyanidin structure depicting flavan-3-ol subunits.

**Figure 2.** (A) Schematic representation of a proanthocyanidin structure consisting of a prodelphinidin subunit (top) and a procyanidin subunit (bottom), and (B) $^{13}$C CPMAS NMR spectra (black solid line) of purified prodelphinidin fraction from blackcurrant leaves (upper) and purified procyanidin fraction from *Tilia* flowers (lower). The assignment of the resonances is consistent with the labelled carbon positions of the proanthocyanidin structure including resolved terminal (4t) and internal (4i) carbons. The grey lines depict the corresponding dipolar dephased solid-state NMR spectra, which were optimized to detect only non-protonated carbons. The asterisks mark the spinning side-bands.

**Figure 3.** Comparison of $^{13}$C CPMAS NMR spectra of milled plant tissues (black line), acetone/water crude extracts (green line) and purified proanthocyanidin fractions (blue line) from (A) blackcurrant leaves and (B) *Tilia* flowers. Black arrows depict the signals at 155 and 132 ppm. All spectra were normalized to the amplitude of the 155 ppm signal to evaluate interferences from other plant components in the area of interest (160-120 ppm, highlighted in grey color).

**Figure 4.** Comparison of $^{13}$C CPMAS NMR spectra of sainfoin plant (A) S3, (B) S1 and (C) S4 tissues. The insets show superimposed regions for comparison of the 155 and 145 ppm proanthocyanidin signals and the 53 ppm lignin signal.
Figure 5. (A) Comparison of the selected regions of $^{13}$C CPMAS NMR spectra recorded for mixtures with nominal PC/PD values of 100/0, 70/30, 50/50, 30/70 and 0/100, prepared by mixing purified PA fractions (bottom) and milled plant tissues (top) of Tilia (PC-rich) and blackcurrant (PD-rich). The red arrows reflect the change in the 132 ppm signal intensity as function of the nominal PC/PD value and a small change in chemical shift. The increase in the color intensity of lines (light grey to black) denotes the increase in the PC/PD value. The black arrows depict other signals that also reflect spectral differences but are less useful due to signal overlaps. All signals were normalized to the amplitude of the signal at 155 ppm (red dot). (B) The relation between the intensity ratio $I_{132}/I_{155}$ and nominal PC/PD values for the calibration mixtures of purified PA fractions and milled plant tissues. The dots denote the experimental points and the red lines show fitted linear dependence.

Figure 6. Fingerprint $^{13}$C CPMAS NMR spectra from sainfoin (A) leaves and (B) stems. Both spectra were fitted using the sum of Gaussian shape lines and the result of the fit is superimposed on the original spectrum (red dotted lines). The grey lines below show Gaussian lines used to fit the corresponding region of the spectrum. The signals from this region were used to estimate PA content and PC/PD ratios and are shown in blue.
Table 1. Proanthocyanidin (PA) Contents and Procyanidin (PC) Contents within Purified PA Fraction Determined by $^{13}$C CPMAS NMR, the HCl-Butanol-Acetone Assay and Thiolysis-HPLC Analysis.

<table>
<thead>
<tr>
<th>Sainfoin plant tissue</th>
<th>Accession</th>
<th>PA (g/100 g of plant tissue)</th>
<th>PC (mg/100 mg of purified PA fraction)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>NMR (– 25%)$^a$</td>
<td>HCl-butanol</td>
</tr>
<tr>
<td>S1</td>
<td>Zeus</td>
<td>16$^b$</td>
<td>12</td>
</tr>
<tr>
<td>S2</td>
<td>Zeus</td>
<td>16</td>
<td>12</td>
</tr>
<tr>
<td>S3</td>
<td>Cholderton</td>
<td>21</td>
<td>16</td>
</tr>
<tr>
<td>S4</td>
<td>Hampshire</td>
<td>14</td>
<td>10</td>
</tr>
<tr>
<td>S5</td>
<td>Ambra</td>
<td>18</td>
<td>14</td>
</tr>
<tr>
<td>S6</td>
<td>Zeus</td>
<td>15</td>
<td>11</td>
</tr>
<tr>
<td>S7</td>
<td>Perly</td>
<td>14</td>
<td>11</td>
</tr>
<tr>
<td>S8$^e$</td>
<td>Cotswold</td>
<td>19</td>
<td>14</td>
</tr>
<tr>
<td></td>
<td>Common</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

$^a$ Correction for lignin interference; $^b$ experimental error was ± 8 g/100 g of plant tissue for the PA content, and ± 16 mg PCs/100 mg of purified PA fraction for the PC content within purified PA fraction; $^c$ experimental error was ± 0.5 g/100 g of plant tissue for the PA content; $^d$ experimental error was ± 0.6 g/100 g of plant tissue for the PA content, and ± 5 mg PCs/100 mg of purified PA fraction for the PC content within purified PA fraction; $^e$ sample S8 was analyzed only once.
Figure 1.
Figure 2.
Figure 3.
Figure 4.
Figure 5.
Figure 6.
GRAPHIC FOR TABLE OF CONTENTS:

13C CPMAS NMR

procyanidins
prodelphinidins

δ13C / ppm

Purified PA
Crude extract
Plant tissue