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^{1}H-^{13}C HSQC NMR Spectroscopy for Estimating Procyanidin/Prodelphinidin and *Cis*/*Trans*-Flavan-3-ol Ratios of Condensed Tannin Samples: Correlation with Thiolysis

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ABSTRACT: Studies with a diverse array of 22 purified condensed tannin (CT) samples from nine plant species demonstrated that procyanidin/prodelphinidin (PC/PD) and *cis/trans*-flavan-3-ol ratios can be appraised by ^{1}H-^{13}C HSQC NMR spectroscopy. The method was developed from samples containing 44 to ~100% CT, PC/PD ratios ranging from 0/100 to 99/1, and *cis/trans* ratios from 58/42 to 95/5 as determined by thiolysis with benzyl mercaptan. Integration of cross-peak contours of H/C-6′ signals from PC and of H/C-2′,6′ signals from PD yielded nuclei adjusted estimates that were highly correlated with PC/PD ratios obtained by thiolysis (R^2 = 0.99). *Cis/trans*-flavan-3-ol ratios, obtained by integration of the respective H/C-4 cross-peak contours, were also related to determinations made by thiolysis (R^2 = 0.89). Overall, ^{1}H-^{13}C HSQC NMR spectroscopy appears to be a viable alternative to thiolysis for estimating PC/PD and *cis*/*trans* ratios of CT, if precautions are taken to avoid integration of cross-peak contours of contaminants.

KEYWORDS: Condensed tannins, proanthocyanidins, procyanidins, prodelphinidins, nuclear magnetic resonance spectroscopy, NMR, thiolysis

INTRODUCTION

Condensed tannins (CTs) (also referred to as proanthocyanidins or PACs) represent a class of polyphenolic plant secondary metabolites that are composed of oligomers and polymers of flavan-3-ols.[1,2] These structures vary not only in flavan-3-ol subunit composition, but also in interflavan-3-ol bond connectivity and mean degree of polymerization (mDP). Condensed tannins are most commonly composed of procyanidin (PC) subunits derived from catechin and epicatechin and of prodelphinidin (PD) subunits derived from gallocatechin and epigallocatechin. Substituents at C-2 and C-3 in the C-ring of epicatechin and epigallocatechin have a *cis* configuration while catechin and gallocatechin possess a *trans* stereochemical orientation (Figure 1). These subunits are typically interconnected by C4-C8 interflavan-3-ol linkages (classified as a B-type linkage, Figure 1), but other less common interunit linkages such as the C4-C6 also occur in CTs.

A major point of interest in CTs stems from the potential positive impact they could bring to the agricultural industry because of their ability to modulate proteolysis during forage conservation and ruminal digestion,[3-7] to prevent bloat,[8] reduce intestinal parasite burdens[9] and lessen methane emissions from ruminants.[10,11] It is thought that the CT composition may play a role in how effectively they impart their biological effects on each of these outcomes, improving both the economical and environmental sustainability of ruminant farm operations. Thus, results from *in vitro* and *in vivo* experiments where CT content is known and the composition is well-defined should reveal CT types and levels that are required for optimizing ruminant health and productivity. Such information would help plant breeders with selection for CT content and structure and also help identify plant varieties that are good candidates for genetic modification.

Analytical techniques allowing for the rapid assessment of chemical structures of CT mixtures within and isolated from plant materials remain a high priority.[12] Development of robust analytical methods is required to gain a better understanding of how CTs affect the interdependency of CT/protein structure-activity relationships. Owing to the structural complexity of CTs, novel approaches are needed for their analysis, including new techniques to corroborate data from existing methods. These analytical techniques are needed for analyzing CT mixtures as these are relevant, and applicable to, nutritional and health research on CTs for both humans[13] and animals.[14]

A variety of analytical techniques have been developed for the characterization and analysis of condensed tannins. Thiolysis with benzyl mercaptan[15,16] is one of the most common methods to obtain compositional and structural data on *in situ* or isolated CT.[17] This method involves acid-catalyzed degradation of CT polymers into reactive monomeric cationic subunits which are subsequently trapped with nucleophiles, such as benzyl mercaptan, providing stable monomeric flavan-3-ol adducts. In this method, extension units are converted into stable C-4 thio ethers whereas terminal units of the polymers are liberated as intact flavan-3-ol monomers. HPLC analysis of the mixtures obtained from these depolymerization studies allows qualitative and quantitative assessment of CTs composition in terms of ratios of PC/PD and *cis*/*trans* subunits and overall mDP. It can thus be used to calculate the purity of isolated CT samples based on the total flavan-3-ol yield. Currently, thiolysis represents one of the most useful techniques available for the analysis of CT composition.

One dimensional (1D) NMR spectroscopic studies have been used previously to determine the compositional aspects of isolated condensed tannin samples by either solution state ^{13}C NMR spectroscopy [18-27] or cross-polarization magic angle spinning (CPMAS) solid state ^{13}C

NMR spectroscopy.[28-31] Solution state ^{13}C NMR spectroscopy has been utilized for determination of PC/PD [18-20,22-27] and *cis/trans* ratios,[18,19,22,24-27] estimations of mDP[18-20,22,24,26,27] and the identification of C4-C6 and C4-C8 linkages.[20,21] These NMR techniques, however, suffer from broad and often times unresolved signals, long acquisition times, and low signal-to-noise ratios which hamper an accurate assessment of CT composition. Solid phase studies of CT-containing plant material have been conducted using ^{13}C CPMAS NMR techniques.[28-31] Although this technique provides good signal-to-noise ratios, signals in the spectra are still broad and frequently overlap with non-CT signals. In addition, ^{13}C CPMAS requires the use of highly specialized equipment.

By contrast, common two-dimensional (2D) NMR techniques have not been extensively explored for assessing the composition of either purified CTs or CT present in whole plant materials.[32] Here we report the use of ^{1}H-^{13}C HSQC NMR spectroscopy as a means to determine PC/PD and *cis/trans* ratios of isolated CT samples.

MATERIALS AND METHODS

General Procedure for Purification and Characterization of Condensed Tannins.

Condensed tannins were purified from dried and milled plant material and analyzed for CT composition and purity as previously described.[15,16] Briefly, dried plant material was milled (typically using a cyclone mill) containing a 1 or 0.5 mm screen and the resulting ground material was extracted with 7:3 acetone/water (3 x 10 mL/g of dried material) and filtered. The combined filtrates were concentrated on a rotary evaporator (<40 ºC) to remove acetone and the resulting aqueous layer was extracted with one-half volume of dichloromethane (2 x) and was freeze-dried. The freeze-dried residue was purified in one of two ways. The first method

involved dissolving the freeze-dried residue in water and applying the resulting mixture to the top of a Sephadex LH-20 column pre-packed in water. The column was eluted with water, removing a majority of the carbohydrates present. Column elution was continued with 3:7 acetone/water (providing sample fraction 1) followed by elution of the column with 1:1 acetone/water to give sample fraction 2, which typically contained CTs of highest purity. Alternatively, the dried extraction residue is adsorbed onto Sephadex LH-20 as a 1:1 methanol/water solution to provide a mixture with the consistency of wet sand. This material is then placed in a Buchner funnel and consecutively rinsed with methanol/water (1:1) followed by a series of acetone/water mixtures (1:1, 7:3, 9:1) with each rinsing conducted three times with a 5 mL solvent per gram of Sephadex LH-20. The three rinse filtrates for each solvent were pooled, concentrated on a rotary evaporator (<40 ºC) to remove the volatile solvent and freeze-dried. In both purification methods, the freeze-dried samples were analyzed by ^{1}H-^{13}C HSQC NMR spectroscopy to assess relative purity and/or thiolysis to provide a numerical purity.

NMR Spectroscopy. ^{1}H, ^{13}C and ^{1}H-^{13}C HSQC NMR spectra were recorded at 27 °C on a BrukerBiospin DMX-500 (^{1}H 500.13 MHz, ^{13}C 125.76 MHz) instrument equipped with TopSpin 2.1 software and a cryogenically cooled 5-mm TXI ^{1}H/^{13}C/^{15}N gradient probe in inverse geometry. Spectra were recorded in DMSO-d_6/pyridine-d_5 (4:1) mixtures and were referenced to the residual signals of DMSO-d_6 (2.49 ppm for ^{1}H and 39.5 ppm for ^{13}C spectra). ^{13}C NMR spectra were obtained using 5K scans (acquisition time 4 h 30 min each). For ^{1}H–^{13}C HSQC experiments, spectra were obtained using 128 scans (acquisition time 18 h 30 min each) obtained using the standard Bruker pulse program (hsqcegtpsi) with the following parameters: Acquisition: TD 1024 (F2), 320 (F1); SW 10.0 ppm (F2), 160 ppm (F1); O1 2500.65 Hz; O2 11,318.20 Hz; D1 = 1.50 s; CNST2 = 145. Acquisition time: F2 channel, 102.55 ms, F1 channel

7.9511 ms. Processing: SI =1024 (F2, F1), WDW = QSINE, LB = 1.00 Hz (F2), 0.30 Hz (F1); PH_mod = pk; Baseline correction ABSG =5 (F2, F1), BCFW = 1.00 ppm, BC_mod = quad (F2), no (F1); Linear prediction = no (F2), LPfr (F1). Samples sizes used for these spectra ranged from 10-15 mg providing NMR sample solutions with concentrations of 20-30 mg/mL.

Calculating Procyanidin/Prodelphinidin (PC/PD) and *Cis/trans*-Flavan-3-ol Ratios. The percentage of PCs in the CT sample was calculated using the equation (1):

$$\% \text{ PC} = \text{PC-6}' / [\text{PD-2}'\text{6}'/2 + \text{PC-6}'] \times 100 \qquad \text{Equation (1)}$$

where PC-6′ is the integration of the contour for the H/C-6′cross-peak of the PC units and PD-2′6′ is the integration of the contour for the H/C-2′,6′cross-peak of the PD units. The PD-2′-6′ value is divided by 2 to account for the signal arising from two sets of correlated nuclei. The percentage of *cis* isomers present in the CT sample was calculated through integration of the respective H/C-4 *cis*- and *trans*-flavan-3-ol cross-peak contours centered around $^{1}H/^{13}C$ chemical shifts of 4.5-4.8/36.0 and 4.4-4.65/37.5 ppm, respectively, and used in equation (2):

% *cis*-flavan-3-ols = *cis*-flavan-3-ols/*cis*-flavan-3-ols + *trans*-flavan-3-ols] x 100 Equation (2)

Integrations of cross-peaks were performed in triplicate and the values were averaged. Integration of the peaks was performed using Topspin 2.1 software.

RESULTS AND DISCUSSION

We have recently shown that ^{1}H-^{13}C HSQC NMR spectroscopy can be a useful tool when assessing the presence of CT in forages and detection of CT left in residues after HCl-butanol treatment,[16] demonstrating the power of 2D NMR techniques. The current study included examining the ^{1}H-^{13}C HSQC NMR spectra of 22 purified CT samples prepared from nine different plant species. Based on thiolysis, the CT samples had PC/PD ratios ranging from 0/100

to 99/1, *cis/trans* ratios ranging from 58/42 to 95/5, and a CT content of 44 to ~100% as determined by thiolysis (Table 1). As an example, ^{1}H-^{13}C HSQC NMR spectrum of CT purified from *Lotus pedunculatus* (big trefoil, sample number 6, Table 1) is given in Figure 2A along with cross-peak assignments. The absence of significant cross-peak NMR signals from non-CT organic compounds in this spectrum also confirms a high degree of purity of this sample.

Determination of PC/PD Ratios. Quantification of signals arising from polymeric materials by ^{1}H-^{13}C HSQC NMR spectroscopy is often hampered by nuclei having differing *T1* and *T2* relaxation times and differences in coupling constants and resonance offset effects.[33] The presence of these effects results in skewing of cross-peak signal contour volumes and thus typically limits the utility of these contours for quantifying structural information. Usually these effects require special spectroscopic treatments, alterations in NMR acquisition parameters such as changes in pulse sequences or increased relaxation delays, before reliable quantification can be made.[34-36]

In the ^{1}H-^{13}C HSQC NMR spectra of these samples, a combination of the nuclei *T1* and *T2* relaxation and resonance offset effects can be observed for most cross-peak signals. The results of these effects lead to cross-peak contours in the spectra whose volumes are not proportional to the corresponding nuclei ratios. As a prime example, integration of the contours for signals arising from H/C-2′,5′ of PC units versus those from H/C-6′ of PC units would normally provide a ratio of 2:1 if none of the above mentioned effects were observed (Figure 2B). However, the integration ratios of H/C-2′,5′ versus H/C-6′ cross-peak contours in PC containing samples from this study showed wide variability with a range from 2.37:1 to 3.86:1 (n= 17, ave. = 3.15, SD ± 0.48). Most of the signals in the ^{1}H-^{13}C HSQC NMR spectra of these purified CT samples followed this trend. A comparison of integration values obtained from the

cross-peak contours could not be directly correlated with theoretical relative intensities of the nuclei giving rise to the signal. Similarly, in an attempt to assess the mean degrees of polymerization (mDP) of these samples, integration of the terminal methylene unit versus any of the other CT cross-peak signals in the spectra also led to no obvious correlation with the thiolysis data of this study. It is worth noting that even integrations of the C-4 methylene units of the flavan-3-ol monomers catechin, epicatechin and epigallocatechin under identical conditions only integrate, on average, to 72% of other signals present in the ^{1}H-^{13}C HSQC NMR spectrum.

However, integration ratios of H/C-6' cross-peak signals from PC units and the H/C-2', 6' cross-peak signal from PD units did show an extremely strong and unbiased relationship with PC/PD estimates from thiolysis determinations (Figure 3A). Thus, this is the first time that ^{1}H-^{13}C HSQC NMR data from purified CT samples have been corroborated with data from an alternative method (thiolysis) to quantify compositional characteristics of CTs. Separate NMR analyses conducted on a limited set of other purified CT samples at the University of Reading confirmed this method as providing reliable PC/PD ratios.

It is not clear how all of the parameters controlling contour intensities are interrelated: do the nuclei involved impart the same or similar *T1* and *T2* relaxation times, coupling constants and resonance offset effects, allowing for accurate comparison of the two contours, or is this simply a coincidence of cancellation of the effects? Answers to these questions remain to be determined.

To test for variability in sample to sample preparation and data acquisition, we prepared duplicate NMR solutions from the same CT samples and obtained NMR spectra of these preparations on different days. These results are given in Table 2. As shown, there is excellent

reproducibility of the method between these duplicate runs. In all, these experiments prove that this is a robust method for estimation of PC/PD ratios in purified CT samples.

Determination of *Cis/Trans* Flavan-3-ol Ratios. In order to assess *cis*- and *trans*-flavan-3-ol ratios (i.e. ratio of epicatechin and epigallocatechin versus catechin and gallocatechin) in these samples we focused on the H/C-4 cross-peak signal (Figure 2C). It has been reported[32] that this signal is segregated into two cross-peaks with $^{1}H/^{13}C$ chemical shifts of ~4.5-4.8/36.0 and ~4.4-4.65/37.5 ppm for the *cis*- and *trans*-flavan-3-ol subunits, respectively. The integration of cross-peak signals in ^{1}H-^{13}C HSQC NMR spectra of the same nuclei with the same connectivity in near identical electronic environments should be straight-forward as they should possess similar, if not identical, *T1* and *T2* relaxation times and pose little or no differences in coupling constants and resonance offset effects. Thus, we should be able to use the data obtained from these ^{1}H-^{13}C HSQC NMR spectra to directly measure this structural element of isolated CTs. The percentage of *cis* isomers present in the CT sample was calculated through integration of the respective H/C-4 *cis* and *trans* cross-peak contours (Figure 2C). Integration ratios from these contours provided strongly related but biased estimates of *cis/trans* ratios relative to thiolysis (Figure 3B). A literature search revealed that this segregation of the *cis* and *trans* signals of flavan-3-ol moieties is most likely not absolute and this could provide an explanation for the bias in *cis/trans* estimates relative to thiolysis. NMR spectroscopic data from epicatechin (*cis*) oligomers report ^{13}C chemical shift in the range of 37.5 ppm, overlapping into the previously designated "*trans*" signal region.[37,38] The lack of signal segregation is more pronounced in structures containing C4-C6 interflavanyl linkages.[38,39] Thus, overlapping of signals from *cis*- and *trans*-flavan-3-ol subunits is the most likely contributing factor for slightly larger discrepancies between the

thiolysis/NMR correlations for *cis/trans*-flavan-3-ol subunit assessments, and may also be responsible for the biased regression fit (Figure 3B).

Precautions. The first issue here, as with most analytical techniques, is to obtain a spectrum with strong signal to noise ratio before attempting to integrate the data. If sample size is limited, extended acquisition times need to be considered. When using this technique on samples of low purity it is imperative that the user be able to recognize any non-CT impurity signals present and avoid incorporating them into the integration values. For PC/PD ratio evaluations, we have found that the signals indicated in Figure 2B are the most common impurity signals which may interfere in obtaining reliable results. These signals most likely arise from trace amounts of non-CT polyphenols present in the sample. For the assessment of *cis/trans* ratios, the problem of integration of non-CT impurities does not seem to be an issue. The H/C-4 cross-peak signals appear, even in spectra of whole plant material, in an area void of other non-CT signals. The major issue in the *cis/trans* ratio assessment is the resolution of the two signals. In some cases these signals are not well resolved (Figure 3B) and care needs to be taken in selecting the integration areas.

In conclusion, the method developed now permits analytical assessment, via 2D ^{1}H-^{13}C HSQC NMR spectroscopy, of two specific chemical properties of purified CT samples: PC/PD and *cis/trans* ratios. Purified CT samples examined encompass the entire range of procyanidin/prodelphinidin ratios from 0/100 to 99/1 and a substantial range of *cis/trans*-flavan-3-ol ratios from 58:42 to 95.5:4.5. The observations outlined here also provide validation of thiolysis data for analysis of CT composition. In contrast to thiolysis, NMR spectroscopy represents a non-destructive analytical tool, which can be important when sample quantities are limited. Thiolysis requires ca 4 mg for a single determination, whereas NMR analysis requires

only 10 mg for an 18 h acquisition time using the described instrumentation. No additional straight-forward correlations were found upon examination of other cross-peak signals in these ^{1}H-^{13}C HSQC NMR spectra. Additional spectroscopic examination of these samples is warranted to investigate whether other significant structural information can be obtained using quantitative ^{1}H-^{13}C HSQC NMR data[34-36] or alternative NMR techniques.

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ABBREVIATIONS USED

^{1}H-^{13}C HSQC, proton-carbon-13 heteronuclear single quantum coherence ; NMR, nuclear magnetic resonance; PC, procyanidin; PD, prodelphinidin; *cis*, 2,3-*cis*; *trans*, 2,3-*trans*; CT, condensed tannins; mDP, mean degree of polymerization; ^{13}C, carbon-13; CPMAS, cross polarization magic angle spinning; 1D, one dimensional; 2D, two dimensional; 5K, five thousand; DMSO-d_6, perdeuterated dimethyl sulfoxide.

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Figure 1. Structures of common flavan-3-ol monomeric subunits found in condensed tannins (left). A condensed tannin tetramer (right) showing C4-C8 (B-Type) linkages, PC and PD extender units and a terminal unit.

Figure 2. (Panel A) Signal assignments for the ^{1}H-^{13}C HSQC NMR spectrum (500/125 MHz, DMSO-d_6/pyridine-d_5, 4:1) of purified condensed tannin sample (Table 1, Sample Number 6) from *Lotus pedunculatus* (big trefoil) leaves; (Panel B) B-Ring aromatic region cross-peak signals including H/C-2′,6′ PD signal and the H/C-2′,5′ and 6′ signals from procyanidin units; and (Panel C) H/C-4 *cis*- and *trans*-flavan-3-ol cross-peak signals. Contours were integrated as indicated by boxes. Non-tannin related signals arising from impurities are noted and are not included in the integration.

Figure 3. (Left Panel) Proportion of procyanidin subunits in 22 isolated condensed tannin samples as determined by thiolysis vs. ^{1}H-^{13}C HSQC NMR. (Right Panel) Proportion of *cis* subunits in 22 isolated condensed tannin samples as determined by thiolysis vs. ^{1}H-^{13}C HSQC NMR.

Table 1. Comparison of Data from Thiolysis and ^{1}H-^{13}C HSQC NMR Determinations for 22 Condensed Tannin (CT) Samples.

CT Sample Number	plant species	CT content (thiolysis) (%)*	SD	% PC (thiolysis)	SD	% PC (NMR)	SD	% *cis* (thiolysis)	SD	% *cis* (NMR)	SD
1	*Lespedeza cuneata*	96.3	0.08	5.9	0.06	4.9	0.06	79.2	0.26	73.2	1.19
2	*Lotus corniculatus*	92.5	0.03	54.0	0.62	55.6	1.17	93.3	0.58	86.8	0.59
3	*Lotus corniculatus*	78.1	0.40	68.0	0.35	70.9	0.19	87.5	0.15	88.7	1.17
4	*Lotus corniculatus*	75.3	0.01	57.1	0.12	60.8	0.32	91.3	0.13	87.0	1.58
5	*Lotus pedunculatus*	108.0	0.01	16.0	0.07	14.6	0.80	81.7	0.22	69.3	1.63
6	*Lotus pedunculatus*	91.3	0.35	25.9	0.29	23.7	0.28	78.7	0.23	75.4	1.22
7	*Lotus pedunculatus*	85.8	0.01	17.5	0.06	17.5	0.50	79.5	0.05	71.7	0.45
8	*Lotus pedunculatus*	80.3	0.41	28.1	0.17	29.0	1.02	74.4	0.15	71.0	1.79
9	*Onobrychis viciifolia*	102.2	8.13	37.3	0.29	39.1	3.23	82.9	0.27	84.4	5.05
10	*Onobrychis viciifolia*	93.7	4.55	19.2	0.06	19.1	0.28	83.3	0.21	77.9	1.96
11	*Onobrychis viciifolia*	82.4	1.10	51.7	0.32	56.7	0.62	83.5	0.10	79.7	0.90
12	*Onobrychis viciifolia*	44.3	0.17	57.3	0.07	59.0	1.45	68.7	0.00	64.9	1.10
13	*Securigera varia*	56.6	n=1	18.2	n=1	22.5	0.17	89.7	n =1	87.6	0.56
14	*Sorghum bicolor*	58.8	0.02	100.0	0.00	100.0	0.00	85.5	0.09	87.1	2.60
15	*Theobroma cacao*	63.8	n = 1	100.0	n=1	100.0	N	93.4	n = 1	100.0	N
16	*Theobroma cacao*	49.0	0.01	100.0	0.0	100.0	0.00	90.1	0.12	88.7	2.06
17	*Tilia sp.*	92.7	0.04	98.5	0.05	99.2	0.19	95.5	0.09	91.2	0.15
18	*Tilia sp.*	61.1	0.47	98.1	0.14	99.2	0.47	89.4	0.11	89.1	0.73
19	*Trifolium repens*	120.6	0.01	0.8	0.00	0.0	N	69.3	0.07	61.1	1.01
20	*Trifolium repens*	111.4	4.80	1.3	0.00	0.0	N	58.9	1.27	56.3	0.75
21	*Trifolium repens*	106.6	5.08	0.9	0.04	0.0	N	58.3	0.24	50.6	1.22
22	*Trifolium repens*	97.6	0.01	1.1	0.04	0.0	N	69.8	0.02	56.1	1.60

N = Not detected; *ND = not determined as based on single analyses* Note: % purity refers to g tannins/100 g fraction; % PD = 100 - % PC; % *trans* = 100 - % *cis*.

Table 2. Comparison of Duplicate NMR Data with Thiolysis Data Obtained from Condensed Tannin (CT) Samples (% PC = Percentage of Procyanidins in CT Sample; % *cis* = Percentage of *cis*-flavan-3-ols in CT Sample).

CT Sample Number	% PC (thiolysis)	SD	% PC (NMR)	SD	% *cis* (thiolysis)	SD	% *cis* (NMR)	SD
3	68.0	0.35	70.0	0.49	87.5	0.15	88.6	1.32
3			71.1	0.50			88.6	0.68
4	57.1	0.12	60.4	0.41	91.3	0.13	89.8	0.53
4			59.7	0.53			91.1	0.50
6	26.0	0.29	24.4	0.35	78.7	0.23	75.4	1.02
6			23.7	0.17			75.1	0.99
7	17.5	0.06	17.5	0.50	79.5	0.05	71.6	0.45
7			18.6	0.47			71.8	2.20
11	51.7	0.32	56.9	0.42	83.5	0.10	79.5	1.20
11			55.5	0.13			80.9	0.68

Note: Percentages for prodelphinidins (PD) and *trans* flavanols are not shown as % PD = 100 - % PC and % *trans* = 100 - % *cis*.

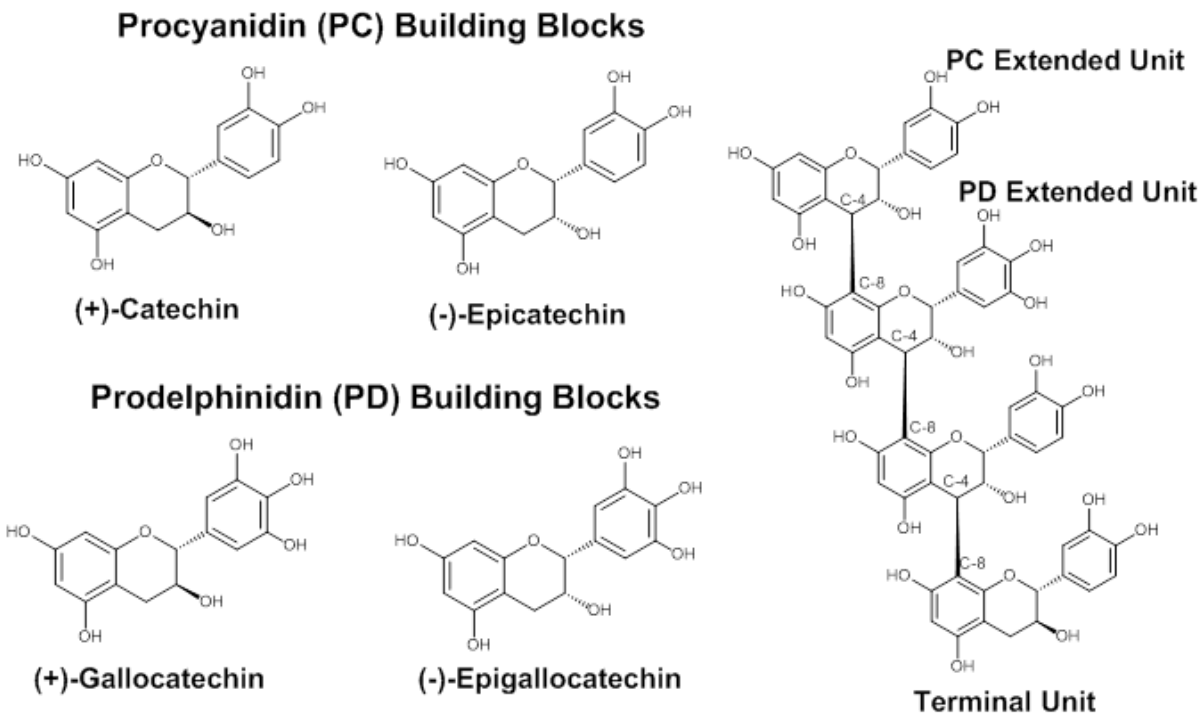
Procyanidin (PC) Building Blocks
(+)-Catechin
(-)-Epicatechin
Prodelphinidin (PD) Building Blocks
(+)-Gallocatechin
(-)-Epigallocatechin
PC Extended Unit
PD Extended Unit
Terminal Unit
C-4
C-8

Figure 1.

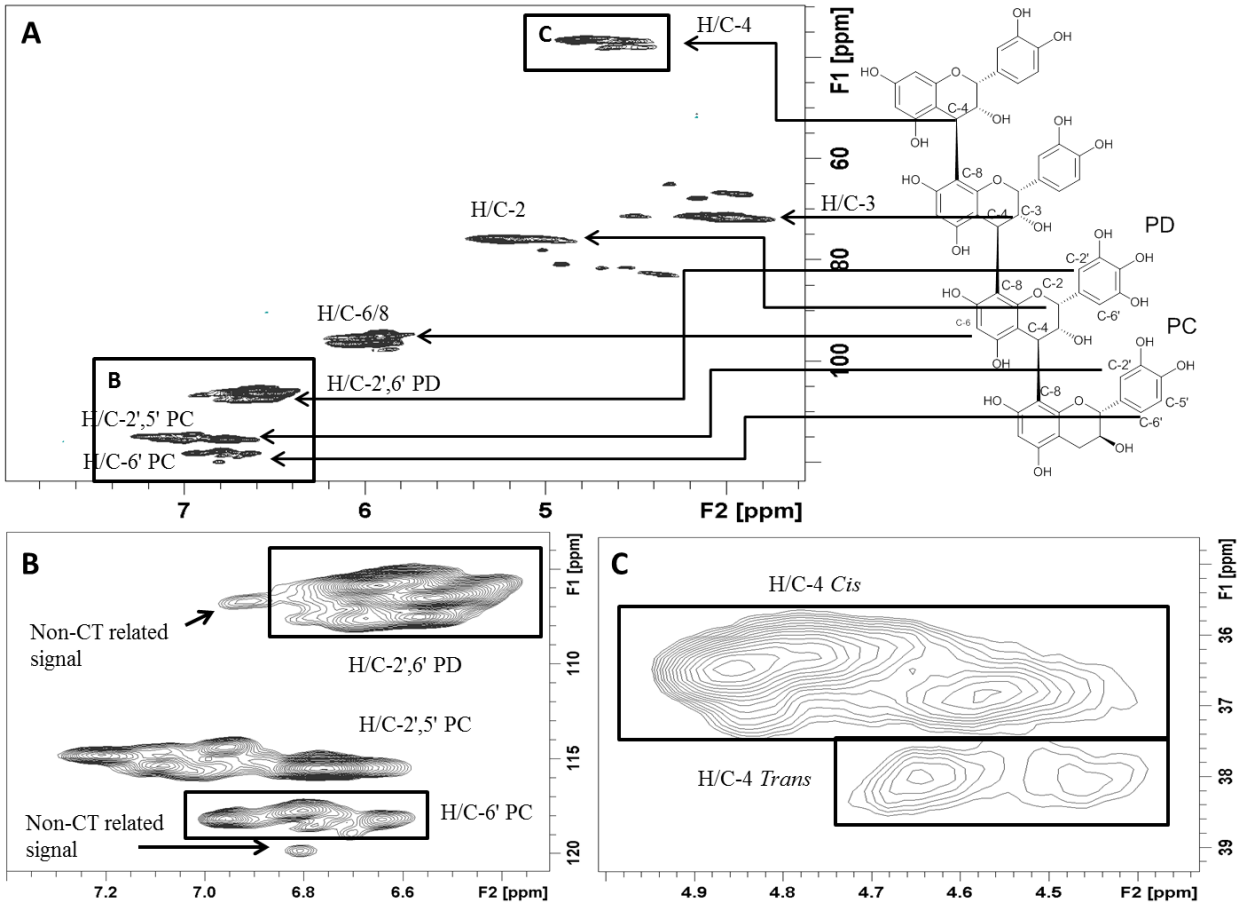
A
C
H/C-4
H/C-2
H/C-3
H/C-6/8
B
H/C-2',6' PD
H/C-2',5' PC
H/C-6' PC
F1 [ppm]
F2 [ppm]
PD
PC
B
Non-CT related signal
H/C-2',6' PD
H/C-2',5' PC
H/C-6' PC
Non-CT related signal
C
H/C-4 Cis
H/C-4 Trans

Figure 2.

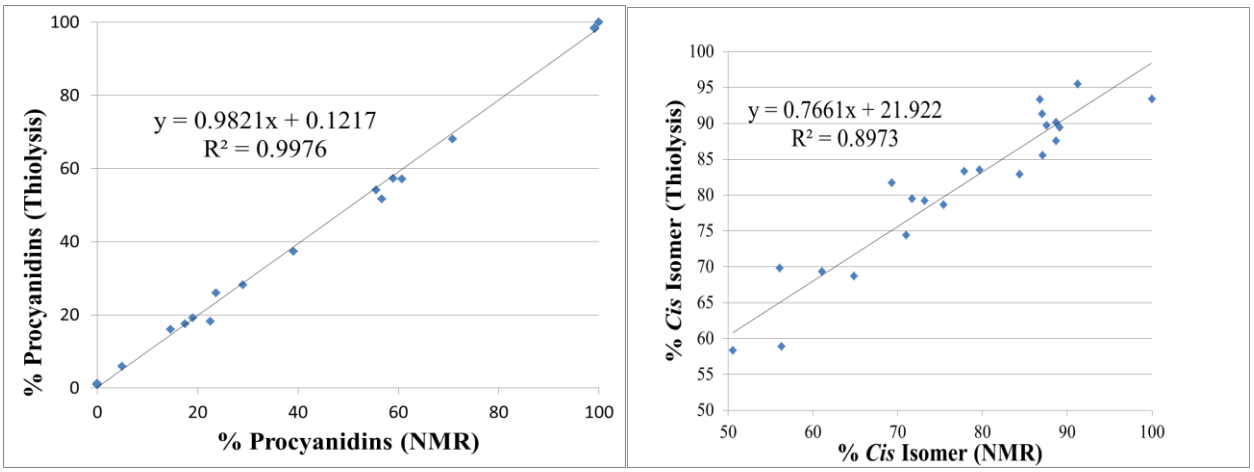
y = 0.9821x + 0.1217
R² = 0.9976
% Procyanidins (Thiolysis)
% Procyanidins (NMR)
y = 0.7661x + 21.922
R² = 0.8973
% Cis Isomer (Thiolysis)
% Cis Isomer (NMR)

Figure 3.

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