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Introduction

Sainfoin (*Onobrychis viciifolia*) is an excellent fodder legume with very high voluntary intakes by cattle, sheep and horses. Ruminants utilise sainfoin **protein** much more **efficiently** than lucerne or soy protein. More efficient nutrient utilisation from sainfoin leads to **less emission of nitrogen, and methane**, which is one of the most damaging greenhouse gases. Research also suggests that sainfoin possesses natural **anthelmintic properties**. It could therefore serve as an ideal fodder legume in more sustainable livestock farming systems.¹

This interdisciplinary research examines the hypothesis that the **condensed tannins** are responsible for the above mentioned benefits of sainfoin. However, there are contradictory reports in the literature concerning the **structures and molecular weights** of sainfoin tannins.² Therefore, detailed chemical analysis is required to identify its active, beneficial tannins. This **structure-activity relationship** study will be of use in future plant breeding/selection studies.

Tannin extraction & fractionation

Procedure:

- freeze dry *Onobrychis viciifolia* var. Cotswold Common (50% flowering, whole plants)
- extract with acetone/H₂O-7:3
yield = 21g/100g sainfoin (on dry matter basis)³
- separate into 2 fractions (with Sephadex LH20) resp. 9 fractions (with Toyopearl HW50F; first 3 fractions were eluted with water and were not analysed)

column packing	MeOH/H ₂ O-1:1* (A)			acetone/H ₂ O-7:3 (B)		
Sephadex LH20 ⁴	13.1%			4.1%		
Toyopearl HW50F	0.9% (A1)	10.5% (A2)	3.6% (A3)	0.1% (B1)	2.3% (B2)	1.2% (B3)

* is expressed asg extract/100g sainfoin (on dry matter basis)
* may contain some sugars

Table 1

Isothermal titration calorimetry

	sumac tannins (GT)	tara tannins (TA)	sainfoin tannins (CT)	chestnut tannins (ET)	myrabolan tannins (ET)
K_a (M ⁻¹)	1.7x10 ⁵	1.0x10 ⁴	6.6x10 ³	9.0x10 ²	7.0x10 ²
binding affinity					
ΔH_{obs} (kJ/mol)	-30.3	-33.0	-21.82	-39.8	-58.1
$T\Delta S$ (kJ/mol)	-0.45	-10.17	0.03	-22.94	-41.87
ΔG (kJ/mol)	-29.85	-22.83	-21.79	-16.86	-16.23
stability of tannin•BSA complex					
N	9.4	2.5	35.2	17.7	22.0

GT = gallotannins
CT = condensed tannins
ET = ellagitannins

K_a = equilibrium binding constant
N = tannin/protein binding stoichiometry

Table 2⁵

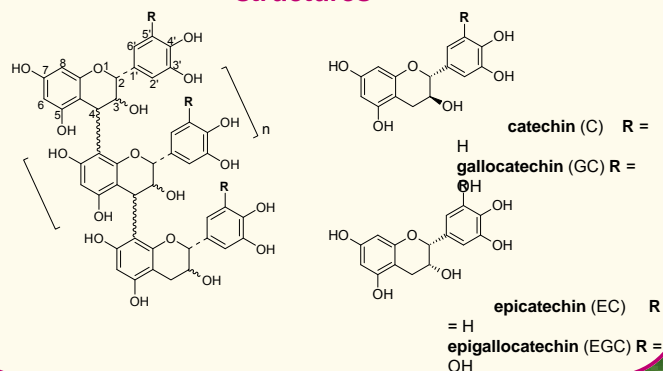
Experimental procedure:

A solution of 3mg/ml of fraction B (Table 1, entry 1) in a 50mM citric acid buffer (pH 6) was titrated into a 10 μ M BSA solution in 50 mM citric acid buffer (pH 6) at 298K. A CSC Nano ITC Series III calorimeter (Calorimetry Sciences Corp., Lindon, UT) was used for all experiments.

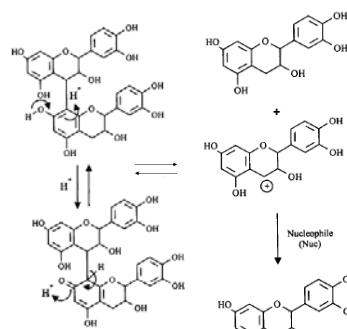
Conclusion

- Calorimetry shows that both the binding affinity and the stability of the sainfoin tannin-BSA complex falls between gallotannins and ellagitannins
- Toyopearl enabled the isolation of a fraction containing high molecular weight tannins
- The high molecular weight fraction has a higher prodelphinidin content, this confirms earlier studies
- Sainfoin contains a highly complex mixture of condensed tannins in terms of PC:PD ratios and molecular weights. Comparisons between the tannins from sainfoin, *Lotus pedunculatus* and *Lotus corniculatus* are ongoing in order to determine if this complexity is the key to their beneficial effects of sainfoin

Condensed tannin composition and structures



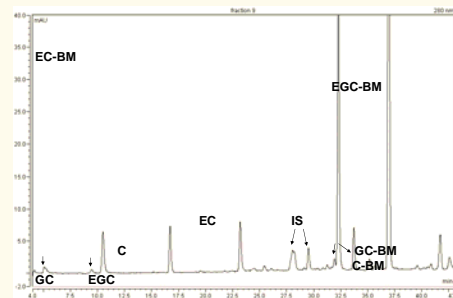
Tannin degradation & analysis



Degradation conditions:

- tannin (1mg), 100 μ l of 5% benzyl mercaptan/MeOH; 100 μ l of 3.3% HCl/MeOH, 50 °C, 20 min
- 40 mM NaOAc/H₂O, rt, 5 min

HPLC conditions: column: Intersil ODS-2, 250x4.6mm; S_μ; solvent A: 1%HOAc/H₂O; solvent B: MeOH; wavelength: 280 nm; internal standard = dihydroquercetin



fraction	mDP	PC: PD	cis:trans
A	8.2 (1.4)	35.0 (0.7) : 66.0 (0.7)	91.0 (0.2) : 9.0 (0.2)
B	14.2 (0.4)	40.0 (0.1) : 60.0 (0.1)	85.0 (0.2) : 15.0 (0.2)
B3	36.3 (5.7)	20.0 (1.4) : 80.0 (1.4)	92.0 (1.4) : 8.0 (1.4)

mDP = mean degree of polymerisation
PC:PD = ratio of procyanidin to prodelphinidin content in the tannins
cis:trans = ratio of cis and trans orientation of the benzyl group on C2 and the OH on C3; values are not corrected for C3-isomerisation
* value between brackets is the standard deviation (n = 2)

Table 3

References & acknowledgements

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