



Attenuated-Total-Reflection Fourier-transformed spectroscopy as a rapid tool to reveal the molecular structure of insect powders as ingredients for animal feed



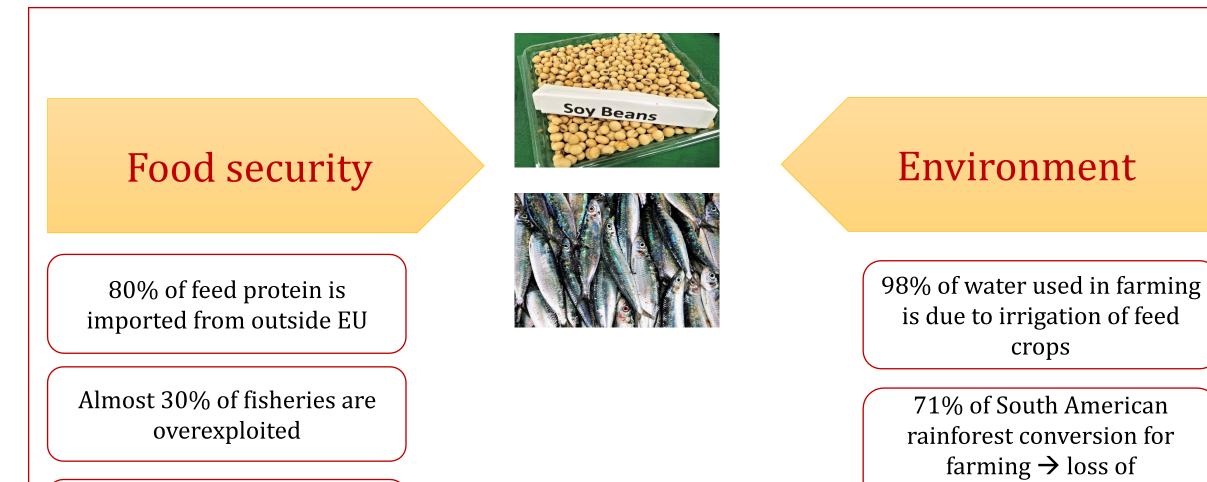
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80% of global arable land is dedicated to livestock

GMO-feeds social concern

High GHG emissions linked to fertilization, pesticides and transport

biodiversity



Protein Feed Source of the future

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Amazon rainforest's final frontier under threat from oil and soya



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Waitrose ends use of GM animal feed on its farms: Critics hail decision as 'beginning of the end' for use of the crops in the UK

• Waitrose meat, milk and eggs will not come from animals on GM feed





The Telegraph

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All seafood will run out in 2050, say scientists





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Monsanto ordered to pay \$289m damages in Roundup cancer trial

() 11 August 2018

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INSECTS ARE A COMMON INGREDIENT IN THE DIET OF MONOGASTRIC ANIMALS



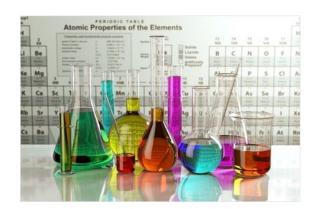






Traditional 'wet' analytical chemistry

- \checkmark Usually looks for a specific known component through homogenisation of the tissue.
- ✓ Information about the spatial origin and distribution of the component of interest is lost.
- ✓ Rely heavily on the use of harsh chemicals altering the native feed structures and possibly generating artifacts.
- $\checkmark~$ Require reasonable amounts of feed material.
- \checkmark Time consuming, expensive and prone to errors within and between laboratories.
- Studying the secondary structure of proteins leads to an understanding of the components that make up a whole protein.
- > Often vital to understanding its digestive behaviour, nutritive quality, utilisation and availability in animals.





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Contents lists available at ScienceDirect Spectrochimica Acta Part A: Molecular and **Biomolecular Spectroscopy**

Application Potential of ATR-FT/IR Molecular Spectroscopy in Animal Nutrition: Revelation of Protein Molecular Structures of Canola Meal and Presscake, As Affected by Heat-Processing Methods, in Relationship with Their Protein Digestive Behavior and Utilization for Dairy Cattle

Katerina Theodoridou and Peigiang Yu*

College of Agricultural and Bioresources, Department of Animal and Poultry Science, University of Saskatchewan, 51 Campus Drive, Saskatoon, SK S7N 5A8 Canada

Journal of Agricultural Science, Page 1 of 13. © Cambridge University Press 2013 doi:10.1017/S0021859613000452

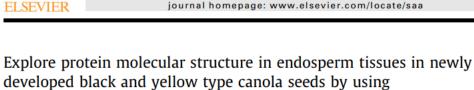
ANIMAL RESEARCH PAPER

Ruminal dry matter and nitrogen degradation in relation to condensed tannin and protein molecular structures in sainfoin (Onobrychis viciifolia) and lucerne (Medicago sativa)

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Magnitude Differences in Bioactive Compounds, Chemical Functional Groups, Fatty Acid Profiles, Nutrient Degradation and Digestion, Molecular Structure, and Metabolic Characteristics of Protein in Newly Developed Yellow-Seeded and Black-Seeded Canola Lines

Katerina Theodoridou,^{†,⊥} Xuewei Zhang,^{†,§} Sally Vail,[#] and Peigiang Yu^{*,†,§}



synchrotron-based Fourier transform infrared microspectroscopy



SPECTROCHIMICA ACTA



Article

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Katerina Theodoridou^a, Sally Vail^b, Peigiang Yu^{a,c,*}

FOOD CHEMISTRY

Synchrotron-Based Microspectroscopic Study on the Effects of Heat Treatments on Cotyledon Tissues in Yellow-Type Canola (Brassica) Seeds

Peiqiang Yu,*^{,†,‡} Katerina Theodoridou,[‡] Hangshu Xin,[‡] Pei-Yu Huang,[§] Yao-Chang Lee,[§] and Bayden R. Wood



CrossMark

Using vibrational infrared biomolecular spectroscopy to detect heat-induced changes of molecular structure in relation to nutrient availability of prairie whole oat grains on a molecular basis

M. D. Mostafizar Rahman^{1,2}, Katerina Theodoridou^{1,3} and Peigiang Yu¹



RESEARCH

<u>Objectives of the study</u>



Assess the potential of FTIR to reveal the nutrient molecular profile of four insect species (allowed to include in animal feed)



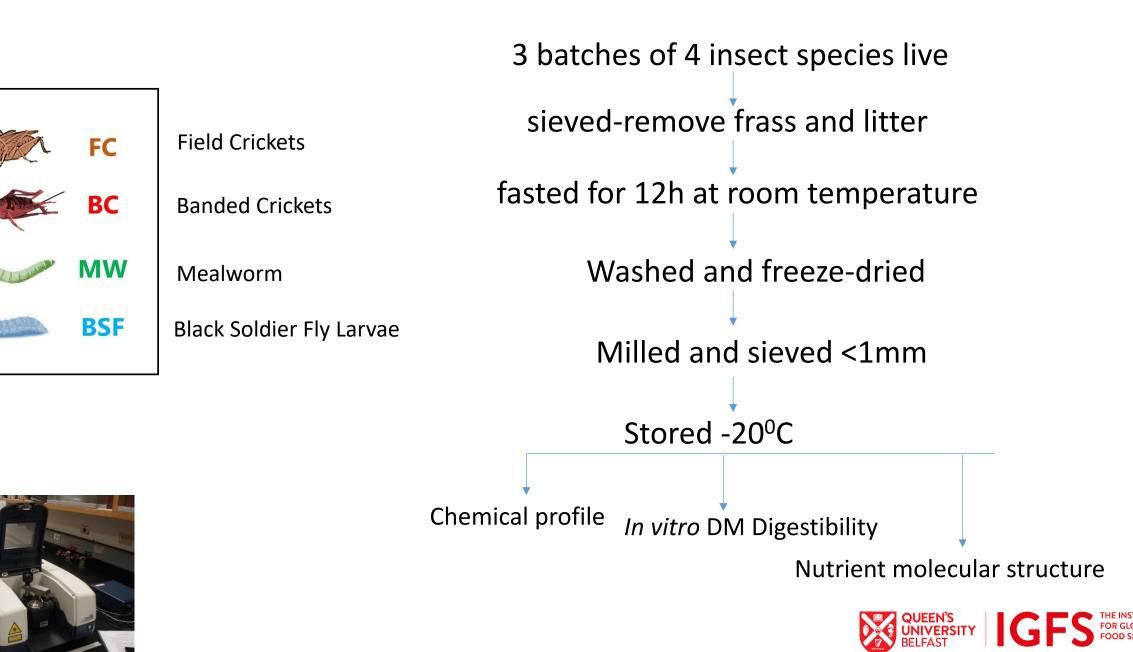
Explore the chemical profile and digestibility of those insect species



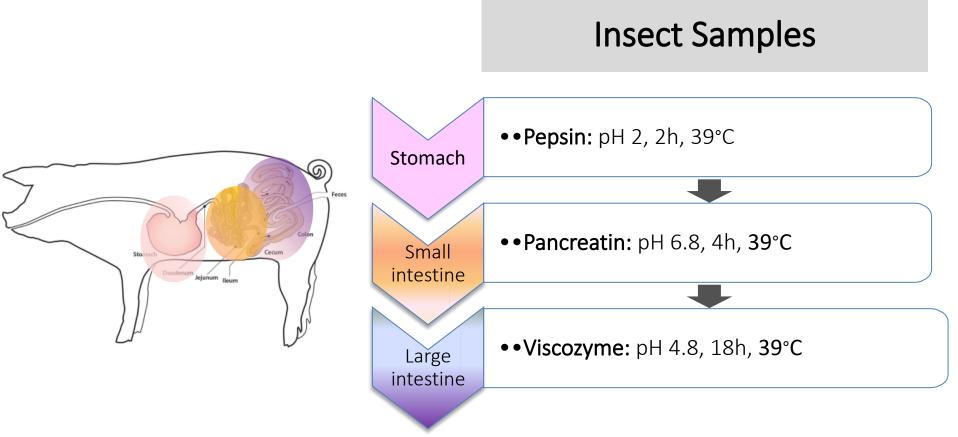
Study the relationship between the molecular structure and the nutritive value of those insect species



Experimental Design



In vitro DM digestibilty



Adapted and modified from Boisen and Fernandez 1997



Statistical Analysis

- One-way ANOVA was performed for the chemical composition molecular structure and digestibility using the statistical package SPSS 25.
- Significance was declared at P<0.10
- Correlation between the molecular structure profile and wet chemistry analysis in the insect samples were analyzed using Pearson's correlations
- The normality test of the residual data was conducted using the Shapiro-Wilk method

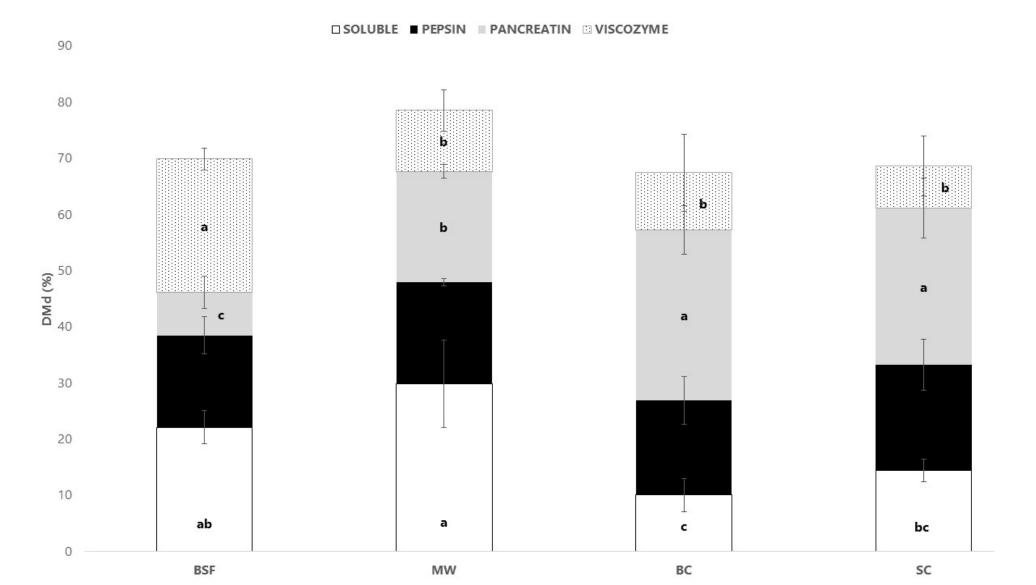
Chemical profile

No significant differences for CP, chitin, NDF/ADF, between cricket species. MW has a higher CP content compared to the rest species (P < 0.0001).

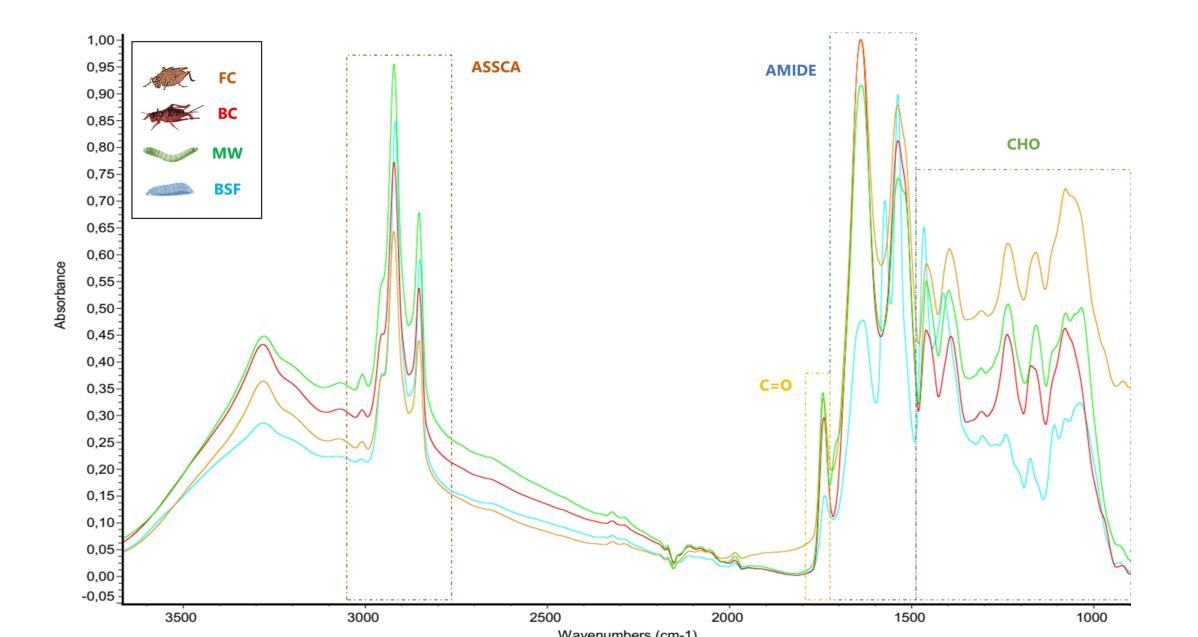
g/100g DM		Insect species									Р	-
	-	BSF		MW		FC		BC				
DM (g/100g)		35.2	а	38.5	а	28.6	b	26.9	b	1.59	* *	-
EE		31.3	ab	35.0	а	21.8	С	25.3	bc	1.82	*	
CP (Nx6.25)	<	38.9	С	50.1	b	66.8	а	62.0	> ^a	3.38	* * *	
ASH		13.0	а	4.69	С	8.58	b	7.06	b	1.050	* *	
NDF		13.2	b	11.7	b	35.2	а	32.3	a	3.31	* * *	
ADF		9.33	bc	8.36		15.6	а	12.1	ab	0.954	* *	
ADICP		4.06	b	4.38	b	9.81	а	5.75	ab	0.338	* *	
ADL (sa)		1.17		1.44		1.50		1.16		0.104	NS	* P < 0.05 ** P < 0.001
Chitin		5.83	b	4.86		7.70	а	7.41) a	0.506	*	*** P < 0.001

In vitro DM digestibilty

MW showed the highest (P<0.05) DMd compared to other species. Pancreatin DMd was higher in both cricket species compared to MW or BSF.

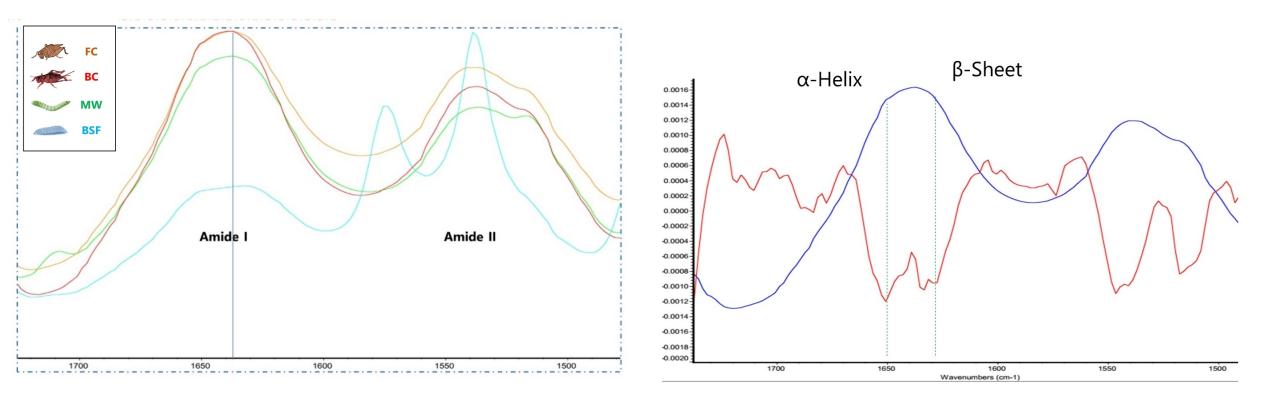


Mid-infrared spectra (3600-900 cm⁻¹)



Primary protein structure: Amide I and Amide II

Secondary protein structure: α-helix and β-sheet



Amide I band was lower (P < 0.0001) for BSF compared to the other species. Amide region of BSF presented an extra peak.

Pattern of α -helix and β -sheet: FC>BC> MW>BSF (P < 0.0001)

Correlations between spectra parameters and nutrition profiles

	AmideA	AM I	AM II	AM III	Aml/Amll	аН	bS	aH/bS
СР	0.96	0.95	-0.62	0.65	0.82	0.96	0.95	0.51
NDF	0.79	0.72	-0.28	0.29	0.52	0.73	0.72	0.40
ADF	0.70	0.65	-0.16	0.30	0.41	0.68	0.66	0.51
CHITIN	0.46	0.42	-0.10	0.12	0.26	0.46	0.42	0.58
PancDMd	0.88	0.91	-0.69	0.68	0.85	0.91	0.90	0.46
ViscDMd	-0.71	-0.76	0.75	-0.67	-0.80	-0.75	-0.74	-0.26

Strong positive correlation (r>0.95; P<0.001) was found between the height of the AmI, α H, β S, AmideA with the CP content.

No correlation (P>0.05) was found with the chitin content.

PancDMd positively correlated (P<0.01) with CP (r=0.91) and NDF (r=0.80), which was reflected in the significant correlations found between these chemical constituents and spectral features, especially AmI, α H, β S.



- Application of FTIR demonstrated the potential to identify nutrient molecular structural differences between insect species rapidly.
- ✓ Correlations between chemical constituents and spectral features, revealing the possibility of using FTIR for the proximal composition profiling of insect powders.
- ✓ FTIR is a rapid technique to predict the chemical composition and quality of insect powders intended to be included in animal feed formulations, as well as to prevent fraud and adulterations.
- ✓ Further analysis focusing on the effect of the insect's life stage, species, feed substrate, and rearing conditions would maximize the possibilities of ATR-FITR applicability.



<u>Future work:</u> 4-year project starts October 2023

'Insects for sustainable animal feed: Livestock farming in a climate change challenged world'





Biotechnology and Biological Sciences Research Council

Thank you for your attention







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