

High-Throughput Antioxidant Profiling in Vegetables by Fourier-Transform Ion Cyclotron Resonance Mass Spectrometry

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ABSTRACT

Vegetables are an important source of dietary antioxidants. Electrospray ionisation Fourier-transform ion cyclotron resonance mass spectrometry was applied to study the antioxidant profiles of beans, broccoli, cabbage, endive, peas and 3 varieties of lettuces. Metabolite extracts from vegetables were analysed in positive and negative ion mode in the m/z range <1000. Mass peaks were assigned by comparing measured and theoretical molecular masses with an accuracy of 0.5 mDa using a metabolite reference database. More than 300 mass peaks were identified, and 100 of them were assigned as potential antioxidant compounds. Antioxidants were grouped into 7 metabolic categories. Phenolic compounds were the most numerous group of antioxidants (45-62%), followed by organic acids (12-24%), amino acids and amines (11-17%), pyridines and purines (5-10%), organosulphur compounds, terpenoids, and alkaloids (<5%). Phenolic compounds included flavonoids, phenolic acids, lignans, coumarins, stilbenes, phenylpropanoid glycosides and quinones. Flavonoids and phenolic acids accounted for more than half of all phenolic compounds. About 20% of antioxidants were found in all analysed vegetables, while 16% of compounds were variety-specific. Hierarchical cluster analysis revealed similarities in antioxidant profiles between cabbage and broccoli, beans and peas and lettuces and endive. With respect to occurrence in vegetable species, antioxidants can be classified as ubiquitous (appearance frequency >90%), common (appearance frequency between 10% and 90%), and rare (appearance frequency <10%). The results of this study demonstrate that ubiquitous antioxidants include primary metabolites and key compounds in secondary metabolic pathways, while rare antioxidants represent the end products of secondary metabolism.

Keywords: bean, broccoli, cabbage, endive, FTICRMS, lettuce, metabolic profiling, pea **Abbreviation: FTICRMS**, Fourier-transform ion cyclotron resonance mass spectrometry

INTRODUCTION

Dietary antioxidants are defined as food compounds which significantly decrease the adverse effects of reactive oxygen species, reactive nitrogen species, or both on the normal physiological function in humans (SEDRI 1998). Vegetables are a rich source of dietary antioxidants, and links between vegetable consumption and a lower risk of coronary heart diseases, diabetes, cataract, degenerative diseases and cancer have been extensively reported (Goldberg 2003; Valko et al. 2007). Antioxidants are the products of primary and secondary plant metabolism. Primary metabolites participate in growth, respiration, photosynthesis, hormone and protein synthesis. Primary metabolites with antioxidant properties include many vitamins and organic acids. Secondary metabolites determine the colour of vegetables, protect plants against herbivores, microorganisms and UV radiation, as well as attract pollinators and seed-dispersing animals. Secondary metabolites with antioxidant properties include phenolic compounds (e.g. flavonoids and phenolic acids), terpenoids (e.g. carotenoids and tocopherols), and alkaloids (e.g. caffeine) (Crozier et al. 2006; Hounsome et al. 2008).

The most characterised antioxidants with respect to their biological activity and occurrence in vegetables are vitamins, carotenoids and flavonoids (Kaur and Kapoor 2001). Vegetables contain vitamins C, E and B. Ascorbic acid is the major form of vitamin C in vegetables, while dehydroascorbic acid represents less than 10% of total vitamin C content (Wills *et al.* 1984). The vitamin B complex of plants includes vitamins B1 (thiamine and its phosphates), B2 (riboflavin and riboflavin 5-phosphate), B3

(nicotinic acid and nicotinamide), B5 (pantothenic acid), B6 (pyridoxal, pyridoxamine, pyridoxine and their 5'-phos-phates), B7 (biotin) and B9 (folic acid). Tocopherols and tocotrienols, known as vitamin E, have been reported in many leafy vegetables, tomatoes, and turnip. Plant carotenoids (α -carotenes, β -carotenes, xanthophylls, lycopene) are orange, yellow, and red lipid-soluble pigments, found in high amounts in tomatoes, carrots and sweet peppers. Carotenoids are essential precursors of vitamin A, which can be produced within the human body from α -carotene and β carotene (Granado et al. 1992; Crozier et al. 2006). Flavonoids include ~4,000 phenolic compounds, such as flavonols, flavones, isoflavones, chalcones and anthocyanidins. Flavonols quercetin, kaempferol and isorhamnetin have been found in onions, leeks, endives and broccoli; flavones apigenin, luteolin and chrysoeriol in parsley, thyme and celery; isoflavones in soy; chalcones and dihydrochalcones in tomatoes. Anthocyanidins cyanidin, delphinidin and malvidin have been identified in red cabbage, radish and red lettuces (Hollman and Katan 1999; Yao et al. 2004). In addition to the above compounds, antioxidants have been found among other groups of plant metabolites, such as organic acids, amino acids, polyamines, pyridines and organosulphur compounds (see **Table 1** for references).

Over the past decade, a significant amount of information has been accumulated on the identification and characterization of plant metabolites. Developments in mass spectrometry, nuclear magnetic resonance and infrared spectrometry enabled the characterization of plant metabolomes. Statistical tools (e.g. hierarchical cluster analysis, principal component analysis and self-organizing map analysis) were applied to compare large data sets. This analysis allowed the identification of compounds (or groups of compounds), which determine the differences between genotypes or phenotypes. Using this approach, the metabolic profiles of potato, tomato, and lettuce have been characterized (Roessner *et al.* 2001; Garratt *et al.* 2005; Schauer *et al.* 2005).

In this study, we applied Fourier-transform ion cyclotron resonance mass spectrometry (FTICRMS) to study antioxidant profiles in beans, peas, broccoli, cabbage, endive and lettuces. A major advantage of FTICRMS over other techniques is its high mass resolution and accuracy, which allows identification of organic compounds without their preliminary separation. The combination of FTICRMS with the low-fragmentation electrospray ionization technique provides an efficient research tool for the characterisation of small biomolecules. Recently, this technique has been successfully applied for metabolic profiling in *Arabidopsis thaliana* seedlings (Ohta *et al.* 2007; Takahashi *et al.* 2008). Development of software for FTICRMS data analysis and metabolite reference database (Shinbo *et al.* 2006) enabled characterisation of metabolic profiles in plants under dark/ light transitions and herbicide treatments (Oikawa *et al.* 2006; Ohta *et al.* 2007).

In our recent study we applied FTICRMS to characterise antioxidant compounds in white cabbage during 6 months of commercial storage (Hounsome *et al.* 2009). Identified compounds included 8 vitamins, 15 flavonoids and 11 phenolic acids. We suggested that changes in antioxidant compounds in cabbage during long-term storage reflect several ongoing processes, such as postharvest senescence, biennial cycle and response to fungal infection.

In this study we demonstrate the application of the electrospray ionization FTICRMS for screening antioxidant compounds in 8 vegetable varieties. We compared antioxidant profiles in beans, broccoli, cabbage, endive, peas and 3 varieties of lettuces, and classified antioxidants with respect to their occurrence in vegetables. This project was conducted as a part of the Research Councils' Rural Economy and Land Use Programme (http://www.relu.ac.uk/) to investigate different aspects of vegetable consumption in the UK from a consumer perspective. Samples were kindly provided by 3 major UK commercial growers who supply vegetables to the UK supermarket chains.

MATERIALS AND METHODS

Plant material and sample preparation

Analysed vegetables included beans (Phaseolus vulgaris L. cv. 'Fontana'), broccoli (Brassica oleracea L. var. italica cv. 'Monaco'), white cabbage (Brassica oleracea L. var. capitata L. alba DC. cv. 'Bison'), endive (Chicorium endivia L. var. crispum cv. 'Glory'), peas (Pisum sativum L. cv. 'Ambassador') and three varieties of lettuces (Lactuca sativa L.): cos (var. longifolia cv. 'Frisco'), iceberg (var. capitata cv. 'Diamond') and red Batavia (cv. 'Sovereign'). Due to the pragmatic nature of this project, plant samples were obtained from 3 major UK commercial growers. Names of companies are not mentioned to maintain commercial confidentiality. Vegetables were grown in a conventional agricultural system during the 2006 growth season in different fields and collected at different times of the year as part of commercial operations. Vegetables were randomly sampled from different batches shortly before their dispatch to supermarkets. Broccoli, cabbage, endive and lettuces were sampled as whole vegetables (broccoli n=4, cabbage n=10, endive n=4, cos lettuce n=4, iceberg lettuce n=4, red Batavia lettuce n=4). Beans and peas were sampled in pods, each sample batch (beans n=6, peas n=4) was between 200-300 g. Vegetables were transported refrigerated to the laboratory and processed on the same day upon arrival. Edible parts of vegetables were used for the analysis. As whole lettuces, cabbages and broccoli florets were too large to process, a standard sampling procedure was established to ensure compatibility between samples. Vegetable tissues (100-150 g) were washed with deionised water and homogenised in liquid nitrogen. Frozen powder (0.25 g) was resuspended in 1 mL 50% methanol solution. Samples were centrifuged for 10 min at $2000 \times g$. Supernatant was diluted 1: 100 using solutions containing 50% methanol and 0.1% formic acid for analysis in positive ion mode and 50% acetonitrile for analysis in negative ion mode. Acid hydrolysis was performed as described by Hertog and others (1982) to convert glycosylated forms of vitamins and flavonoids. For this purpose vegetable extracts were incubated at 80°C for 2 h in 1.2 M HCl and 50% methanol. Processed samples were stored at -20°C and analysed within 1-2 weeks. No significant changes in mass spectra were observed during 6 months of sample storage.

FTICRMS measurements

Mass spectra were obtained using the Bruker Apex IV FT-ICR mass spectrometer (Bruker Daltonics Inc. Billerica, MA, USA), equipped with 160 mm bore 4.7 Tesla actively shielded magnet (Magnex Scientific Ltd, Oxford, UK) and an Apollo electrospray ionization source. Samples were directly injected into the mass spectrometer using a Cole Parmer series 74900 syringe pump (Cole Parmer Instrument Co., London, UK) at a flow rate of 120 µL/h. Ions were generated in positive and negative ion mode at a nebulizer N₂ gas pressure of 3.5×10^5 Pa. and a dry gas temperature of 250°C. Source voltages on the capillary entrance were -4500 V and 4300 V, on the spray shield -3900 V and 4000 V, and on the capillary exit 78 V and -70 V for positive and negative modes, respectively. Ions were accumulated for 0.3 seconds in the external hexapole ion guide. The potential on the front trapping plate was 0.84 V and -1 V, on the back plate 0.8 V and -1.1 V, and on the analyser entrance -4 V and 8 V for positive and negative modes, respectively. The data were acquired using apexControl 1.0 $(\text{Compass}^{\text{TM}} \text{ software suite, Bruker Daltonics Inc.}).$ The scan range was set to m/z 50-1000. One hundred scans were accumulated to obtain each mass spectrum. The mass scale was calibrated using external standards, containing 0.5 µg/mL caffeine, diphenhydramine, sulphametazine, oxybutynin, terfenadine and reserpine. Internal calibration was performed by the addition to the sample of 1 μ M ¹⁵N-labelled amino acids glycine and tyrosine (98% enrichment, Cambridge Isotope Laboratories, Inc. UK).

Data analysis

Mass spectra were analysed using DataAnalysis 3.3 (CompassTM software suite, Bruker Daltonics Inc.). Signal-to-noise threshold cut-off was set to 4. ¹³C peaks were excluded from mass analysis. Peak list files were screened through the KNApSAcK metabolite reference database (http://kanaya.naist.jp/KNApSAcK/), containing ~23,000 metabolites. Compounds were identified by comparing the measured and theoretical exact molecular masses with an accuracy of ± 0.5 mDa. Mass lists were screened for 3 ion species (M+H)⁺, (M+K)⁺ and (M+Na)⁺ in positive ion mode, and for one ion (M-H)⁻ in negative ion mode. Each list consisted of 1500 to 3000 ion masses. In total, 40 vegetables representing 8 varieties were analysed in 3 replications in both positive and negative ion mode. The complete dataset contained 240 peak lists (120 samples in 2 ion modes). Hierarchical cluster analysis was performed using GeneSight 3.5 (BioDiscovery Inc.) on identified masses.

RESULTS

Analysis of the FTICRMS spectra of vegetables in the m/z < 1000 revealed ~ 3000 mass peaks in positive ion mode and ~ 1500 in negative. A mass accuracy of < 2 ppm was routinely achieved under our experimental conditions. More than 300 mass peaks were assigned by comparing measured and theoretical molecular masses with a precision of 0.5 mDa. One hundred peaks were assigned as antioxidant compounds. Although the screening of mass spectra with a precision of 0.5 mDa in most cases resulted in just one empirical formula, the database search often produced several structurally similar antioxidant compounds. **Table 1** summarises the information on antioxidants found in beans, peas, broccoli, cabbage, endive and lettuces. Only metabolites with reported antioxidant activities were included in this study.

The major metabolic classes of antioxidants found in

Exact	Molecular	Metabolic	Assigned compound	Reference	Vegetables
nass	formula	category			
38.0160	$C_3H_4O_3$	organic acid	pyruvic acid	Mallet and Sun 2003	beans, cabbage
8.1000	$C_4H_{12}N_2$	polyamine	putrescine	Das and Misra 2004	beans, broccoli, cabbage, cos, endive, iceberg, red Batavia
0.0317	C ₃ H ₆ O ₃	organic acid	lactic acid	Groussard et al. 2000	beans, broccoli, cabbage
02.0317	$C_4H_6O_3$	organic acid	acetoacetic acid	Mallet and Sun 2003	beans, broccoli, cabbage, cos, endive, iceberg,
02.0317	0411603	organic dela		Wallet and Sull 2005	peas, red Batavia
10.0368	$C_6H_6O_2$	phenolic	catechol,	Arts et al. 2003	beans, cabbage, cos
110.0508	0.11002	phenone	hydroquinone,	7 H IS Ci ul. 2005	
			resorcinol		
18.0266	$C_4H_6O_4$	organic acid	succinic acid	Naylin et al. 2006	beans, cos
122.0368	$C_7H_6O_2$	phenolic	benzoic acid	Weitzman and Stossel 1982	
		1			peas, red Batavia
122.0480	C ₆ H ₆ N ₂ O	pyridine	nicotinamide (vitamin B ₃)	Crowley et al. 2000	cos, endive, iceberg, peas
23.0320	C ₆ H5NO ₂	pyridine	nicotinic acid (vitamin B ₃)	Crowley et al. 2000	beans, broccoli, cabbage, peas
24.0524	$C_7H_8O_2$	phenolic	guaiacol,	Dizhbite et al. 2004	beans, broccoli, cabbage, cos, endive, iceberg,
			orcinol,	Hladyszowski et al. 1998	peas, red Batavia
			p-hydroxybenzyl alcohol	Liu and Mori 1993	
30.1218	$C_5H_{14}N_4$	amine	agmatine	Lee et al. 2003	broccoli, cabbage, endive, peas
32.0575	C_9H_8O	phenolic	cinnamaldehyde	Kim et al. 2007	beans, broccoli, cabbage, cos, endive, iceberg,
					peas
134.0215	$C_4H_6O_5$	organic acid	malic acid	Puntel et al. 2007	beans, broccoli, cabbage, cos, endive, iceberg,
					peas, red Batavia
135.0354	$C_4H_9NO_2S$	amino acid	homocysteine	Meucci and Mele 1997	beans, cabbage, cos, endive, iceberg, red
					Batavia
136.0524	$C_8H_8O_2$	phenolic	phenylacetic acid	Kim and Lee 2004	beans, broccoli, cabbage, cos, endive, iceberg,
					peas, red Batavia
136.1252	$C_{10}H_{16}$	terpenoid	limonene	Gerhäuser et al. 2003	beans, broccoli, cabbage, cos, endive, iceberg,
					peas, red Batavia
137.0841	$C_8H_{11}NO$	amine	tyramine	Yen and Hsieh 1997	beans, cos
138.0317	C ₇ H ₆ O ₃	phenolic	hydroxybenzoic acid	Gadow <i>et al.</i> 1997	peas
145.1579	$C_7 H_{19} N_3$	polyamine	spermidine	Das and Misra 2004	beans, broccoli, cabbage, cos, endive, iceberg,
					peas, red Batavia
146.0224	$C_6H_{10}S_2$	organosulphur	diallyl disulfide	Koh <i>et al.</i> 2005	cabbage, peas
46.0368	$C_9H_6O_2$	phenolic	coumarin	Kostova 2006	beans, broccoli, cabbage, cos, endive, iceberg,
146.0570	C II O		- 4:-::4	Demodern seelen of all 2001	peas, red Batavia
146.0579	$C_6H_{10}O_4$	organic acid	adipic acid	Papadopoulos et al. 2001	beans, broccoli, cabbage, cos, endive, iceberg,
148.0524	СНО	phenolic	cinnamic acid	Kanski <i>et al</i> . 2002	peas, red Batavia beans, broccoli, cabbage, cos, endive, iceberg,
148.0324	$C_9H_8O_2$	phenolic	chinamic acid	Kanski <i>el al</i> . 2002	peas, red Batavia
140.0510	C ₅ H ₁₁ NO ₂ S	amino acid	methionine	Meucci and Mele 1997	beans, broccoli, cabbage, cos, iceberg, endive,
149.0510	$C_5\Pi_{11}NO_2S$	ammo aciu	methonne	Medeel and Mele 1997	peas
150.0164	C ₄ H ₆ O ₆	organic acid	tartaric acid	Papadopoulos et al. 2001	cos, endive, iceberg, red Batavia
150.0537	$C_{4}H_{6}O_{6}$ $C_{6}H_{14}S_{2}$		dipropyl disulfide	Munday <i>et al.</i> 2003	beans, broccoli, cabbage, cos, endive, iceberg,
150.0557	06111402	organosuiphur	alpropyr alsunae	Williady Cr ur. 2005	peas, red Batavia
150.0681	$C_9H_{10}O_2$	phenolic	coumaryl alcohol	Ly et al. 2003	beans, broccoli, cabbage, cos, endive, peas, rec
150.0001	0911002	phenone	countary aconor	Ly Ci ui. 2005	Batavia
152.0473	$C_8H_8O_3$	phenolic	vanillin,	Kamat et al. 2000	beans, broccoli, cabbage, cos, endive, iceberg,
	0,11,003	phonone	3,4- dihydroxyacetophenone		peas, red Batavia
154.0266	$C_7H_6O_4$	phenolic	2,3-dihydroxybenzoic acid,	Sha and Schacht 2000	beans, endive
151.0200	- /0 - 4	P	gentisic acid,	Ashidate <i>et al.</i> 2005	
			protocatechuic acid	Saito and Kawabata 2004	
154.1358	C10H18O	terpenoid	geraniol	Choi <i>et al.</i> 2000	cabbage, peas
64.0473	$C_9H_8O_3$	phenolic	coumaric acid	Kikuzaki et al. 2002	peas, red Batavia
64.0685	C ₆ H ₁₂ O ₅	phenolic	quercitol	Orthen et al. 1994	beans, broccoli, cabbage, peas, red Batavia
164.0837	$C_{10}H_{12}O_2$	phenolic	eugenol	Reddy and Lokesh 1992	beans, cabbage, cos, endive, iceberg, peas, red
		1	5	5	Batavia
166.0491	$C_6H_6N_4O_2$	purine	methylxanthine	Geraets et al. 2006	broccoli, cabbage, cos, peas, red Batavia
166.0630	$C_9H_{10}O_3$	phenolic	apocynin,	Ben-Shaul et al. 2001	broccoli, cabbage
		1	paeonol	Hsieh et al. 2006	<i>, e</i>
168.0423	$C_8H_8O_4$	phenolic	vanillic acid	Mansouri et al. 2005	cabbage, red Batavia
168.0899	$C_8H_{12}N_2O_2$	pyridine	pyridoxamine (vitamin B ₆)	Jain and Lim 2001	beans, cabbage, cos, endive, red Batavia
170.0215	$C_7H_6O_5$	phenolic	gallic acid	Yilmaz and Toledo 2004	cabbage, endive
174.0164	C ₆ H ₆ O ₆	organic acid	aconitic acid,	Sousa et al. 2007	beans, broccoli, cabbage, cos, endive, iceberg,
		-	dehydroascorbic acid	Otero et al. 1997	peas, red Batavia
174.0528	$C_{7}H_{10}O_{5}$	organic acid	shikimic acid	Sousa et al. 2007	cabbage, peas
176.0321	$C_6H_8O_6$	organic acid	ascorbic acid	Otero et al. 1997	broccoli, cabbage
177.0460	C ₆ H ₁₁ NO ₃ S	organosulphur	alliin	Chung 2006	beans, cabbage
179.0794	C ₆ H ₁₃ NO ₅	amine	glucosamine	Yan 2007	beans, broccoli, cabbage, cos, endive, iceberg,
					peas, red Batavia
180.0423	$C_9H_8O_4$	phenolic	caffeic acid	Gülçin 2006	beans, broccoli, cabbage, cos, endive, iceberg
					red Batavia

Table 1 (C) Exact mass	Molecular formula	Metabolic category	Assigned compound	Reference	Vegetables
180.0786	C ₁₀ H ₁₂ O ₃	phenolic	coniferyl alcohol	Barclay et al. 1997	beans, cabbage, cos, endive, iceberg, peas, red Batavia
181.0739	C ₉ H ₁₁ NO ₃	amino acid	tyrosine	Meucci and Mele 1997	beans, cabbage, endive, iceberg, red Batavia
182.0579	$C_9H_{10}O_4$	phenolic	dihydrocaffeic acid,	Huang <i>et al.</i> 2004	beans, cabbage, cos, endive, iceberg, peas,
102.0379	0911004	phenone	4-hydroxy-3,5- dimethoxybenzaldehyde	Zheng et al. 2008	red Batavia
192.0270	$C_6H_8O_7$	organic acid	citric acid, isocitrate	Papadopoulos et al. 2001	beans, broccoli, cabbage, cos, endive, iceberg, peas, red Batavia
192.0423	$C_{10}H_8O_4$	phenolic	scopoletin	Shaw <i>et al.</i> 2003	cabbage
192.0425	$C_8H_{10}N_4O_2$	alkaloid	caffeine	Devasagayam <i>et al.</i> 1996	beans, cabbage, cos, endive, iceberg, peas, red Batavia
196.0610	$C_9H_{10}O_4$	phenolic	veratric acid	Miyazawa et al. 2003	beans, cabbage,
198.0528	$C_9H_{10}O_5$	phenolic	syringic acid	Schmeda-Hirschmann <i>et al.</i> 2004	beans, cabbage, cos, iceberg
208.0372	$C_{10}H_8O_5$	phenolic	fraxetin	Fernandez-Puntero et al. 2001	broccoli, cos, red Batavia
212.0837	$C_{14}H_{12}O_2$	phenolic	pinosylvin	Stojanović and Brede 2002	beans, cabbage, cos, endive, iceberg, peas, red Batavia
214.0994	$C_{14}H_{14}O_2$	phenolic	dihydropinosylvin	Cai et al. 2003	broccoli, cabbage, cos, iceberg, red Batavia
216.0423	$C_{12}H_8O_4$	phenolic	bergapten,	Yu et al. 2005	beans, broccoli, cabbage, cos, endive,
			xanthotoxin,	Fylaktakidou et al. 2004	iceberg, peas, red Batavia
220.0372	$C_{11}H_8O_5$	phenolic	purpurogallin	Sugiyama et al. 1993	cabbage, cos, iceberg
222.0674	$C_7H_{14}N_2O_4S$	amino acid	cystationine	Wada et al. 1996	beans, cabbage, cos, endive, iceberg, peas, red Batavia
222.0892	$C_{12}H_{14}O_4$	terpenoid	apiole	Singh et al. 2005	cos, endive, red Batavia
226.0994	$C_{15}H_{14}O_2$	phenolic	pinosylvin methyl ether	Pietarinen et al. 2006	beans, cabbage, cos, endive, iceberg, peas, red Batavia
228.0423	$C_{13}H_8O_4$	phenolic	euxanthone	Lin et al. 2005	beans, peas
228.0786	C ₁₄ H ₁₂ O ₃	phenolic	resveratrol, 5,6-dehydrokawain	Cai <i>et al.</i> 2003 Habsah <i>et al.</i> 2003	beans, cos
228.1150	C ₁₅ H ₁₆ O ₂	phenolic	dihydropinosylvin-monomethyl ether		beans, endive
238.0630	C ₁₅ H ₁₀ O ₃	phenolic	3-hydroxyflavone, 5-hydroxyflavone	Lemańska <i>et al.</i> 2001	beans, cabbage, cos, endive, iceberg, peas, red Batavia
240.0786	$C_{15}H_{12}O_3$	phenolic	flavidin, 2,4-dihydroxychalcone, 7-hydroxyflavanone	Jayaprakasha <i>et al.</i> 2004 Calliste <i>et al.</i> 2001 Ávila <i>et al.</i> 2001	beans, cabbage, endive, iceberg, peas, red Batavia
242.0943	$C_{15}H_{14}O_3$	phenolic	equol	Arora et al. 1998	beans, cabbage, cos, endive, iceberg, peas, red Batavia
244.0372	$C_{13}H_8O_5$	phenolic	gentisein, 1,2,5-trihydroxyxanthone	Lin <i>et al.</i> 2005 Minami <i>et al.</i> 1994	beans, cos, endive, iceberg, red Batavia
244.0736	$C_{14}H_{12}O_4$	phenolic	oxyresveratrol, piceatannol	Lorenz <i>et al.</i> 2003 Ovesna <i>et al.</i> 2006	beans, cabbage, iceberg
249.0402	$C_8H_{12}NO_6P$	pyridine	pyridoxine 5-phosphate (vitamin B ₆)	Jain and Lim 2001	beans, cabbage, cos, endive, iceberg, red Batavia
250.1317	$C_{13}H_{18}N_2O_3$	phenolic	caffeoylputrescine	Drolet et al. 1986	cabbage, cos, peas
252.1150	C ₁₇ H ₁₆ O ₂	phenolic	hinokiresinol	Song et al. 2007	beans, cabbage, cos, endive, iceberg, peas, red Batavia
254.0579	$C_{15}H_{10}O_4$	phenolic	chrysin,	Harris et al. 2006	beans, cabbage, peas
			rubiadin	Tripathi et al. 1997	
256.0372	$C_{14}H_8O_5$	phenolic	purpurin	Watanabe et al. 1993	red Batavia
266.1307	$C_{18}H_{18}O_2$	phenolic	honokiol,	Park et al. 2003	beans, cabbage, endive, iceberg, peas, red
			magnolol	Chen et al. 2001	Batavia
271.0606	$C_{15}H_{11}O_5$	phenolic	pelargonidin	Pietta 2000	cabbage
272.0685	$C_{15}H_{12}O_5$	phenolic	butein, butin,	Chen <i>et al.</i> 2006 Zhang <i>et al.</i> 2008	beans, cabbage
272 0007	C II O	mh an a ^{1:} -	naringenin chalcone	Calliste <i>et al.</i> 2001	aakkaaa
272.0896 286.0477	$\begin{array}{c} C_{12}H_{16}O_{7} \\ C_{15}H_{10}O_{6} \end{array}$	phenolic phenolic	arbutin kaempferol, luteolin	Myagmar <i>et al.</i> 2004 Brown <i>et al.</i> 1998	cabbage peas
286.0841	C ₁₆ H ₁₄ O ₅	phenolic	brazilin	Choi et al. 2007	beans, cabbage
287.0556	$C_{15}H_{11}O_{6}$	phenolic	cyanidin	Meyer <i>et al.</i> 1998	cabbage
290.0790	$C_{15}H_{10}O_{6}$ $C_{15}H_{14}O_{6}$	phenolic	catechin, epicatechin	Yilmaz and Toledo 2004	peas
292.1099	$C_{19}H_{16}O_3$	phenolic	purpuritenin A	Maurya and Yadav 2005	beans, red Batavia
304.0583	$C_{15}H_{12}O_7$	phenolic	dihydroquercetin	Dok-Go <i>et al.</i> 2003	beans
306.0740	$C_{15}H_{14}O_7$	phenolic	gallocatechin,	Rice-Evans et al. 1997	cabbage
307.0838	$C_{10}H_{17}N_3O_6S$	organosulphu	epigallocatechin glutathione	Noctor and Foyer 1998	cabbage
326.1518	C ₂₀ H ₂₂ O ₄	r phenolic	licarin A	Yu <i>et al.</i> 2000	cos, iceberg
327.1471	$C_{20}H_{22}O_4$ $C_{19}H_{21}NO_4$	alkaloid	boldine	Jang <i>et al.</i> 2000	beans, cabbage, cos, iceberg

Table 1 (Cont.)							
Exact	Molecular	Metabolic	Assigned compound	Reference	Vegetables		
mass	formula	category					
328.0947	$C_{18}H_{16}O_{6}$	phenolic	salvigenin	Lu and Foo 2002	beans		
328.1311	C19H20O5	phenolic	decursin	Kang and Kim 2007	iceberg		
338.1154	$C_{20}H_{18}O_5$	phenolic	lupiwighteone, wighteone	Erasto et al. 2004	beans		
342.1315	$C_{16}H_{22}O_8$	phenolic	coniferin	Kayano et al. 2004	peas		
344.0896	$C_{18}H_{16}O_7$	phenolic	cirsilineol,	Kelm et al. 2000	cabbage		
			nevadensin,	Ganapaty et al. 2007			
			penduletin,	Hajdú <i>et al</i> . 2007			
			usnic acid	Odabasoglu et al. 2006			
360.1573	$C_{20}H_{24}O_6$	phenolic	lariciresinol	Pietarinen et al. 2006	beans, cabbage		
372.1420	C17H24O9	phenolic	syringin	Es-Safi et al. 2007	cos, iceberg		
388.0907	$C_{18}H_{16}N_2O_8$	phenolic	betanidin	Kanner et al. 2001	cabbage		
388.1158	$C_{20}H_{20}O_8$	phenolic	artemetin	Dugas et al. 2000	cabbage		
402.1315	$C_{21}H_{22}O_8$	phenolic	nobiletin	Murakami et al. 2000	cos, red Batavia		
420.1784	$C_{22}H_{28}O_8$	phenolic	lyoniresinol	Tomosaka et al. 2008	peas		
462.0798	$C_{21}H_{18}O_{12}$	phenolic	kaempferol glucuronide,	Plumb et al. 1999	endive		
			luteolin glucuronide,	Lu and Foo 2001			
			scutellarein glucuronide	Sanz et al. 1994			
578.1424	$C_{30}H_{26}O_{12}$	phenolic	apigenin 7-rutinoside,	Wang et al. 2003	beans		
			procyanidin B1-B8	Teissedre et al. 1996			

vegetables included phenolics, terpenoids, organic acids, amino acids and amines, pyridines and purines, alkaloids and organosulphur compounds. Phenolic compounds represented the most numerous group of antioxidants (45-62%), followed by organic acids (12-24%), amino acids and amines (11-17%), pyridines and purines (5-10%), organosulphur compounds, terpenoids and alkaloids (<5%) (Fig. 1). Phenolic compounds included flavonoids (e.g. naringenin, cyanidin, quercitol), phenolic acids (e.g. benzoic, cinnamic, caffeic), lignans (e.g. hinokiresinol, lariciresinol, lyoniresinol), coumarins (e.g. decursin, fraxetin, xanthotoxin), stilbenes (e.g. resveratrol, oxyresveratrol, pinosylvin), phenylpropanoid gycosides (syringin, coniferin), quinones (arbutin, purpurin), etc. (Table 1). Flavonoids, the most numerous group of phenolics, were presented flavanols (catechin, gallocatechin), flavanones (naringenin, butin), flavones (artemetin, chrysin), isoflavones (gentisein, equol), flavonols (kaempferol) and anthocyanidins (cyanidin, pelargonidin). Phenolic acids included benzoic acid and its derivatives (dimethoxybenzoic, syringic, gallic, vanillic, veratric acids), cinnamic acid and its derivatives (caffeic, coumaric, rosmarinic acids) and phenylacetic acid (Table 1). Flavonoids and phenolic acids accounted for more than half of all phenolic compounds. About 12% of phenolics (e.g. benzoic acid, cinnamic acid, coniferyl alcohol, coumarin and vanillin) were present in all tested vegetables.

Non-phenolic antioxidants represented 38-55% of total number of assigned compounds (**Fig. 1**). These antioxidants included organic acids (acetoacetic, aconitic, adipic, ascorbic, citric, lactic, malic, pyruvic, shikimic, succinic, tartaric), amino acids (cystationine, homocysteine, methionine, tyrosine), amines (agmatine, tyramine), polyamines (putrescine, spermidine), pyridines (nicotinamide, nicotinic acid, pyridoxamine, pyridoxine 5-phosphate), organosulphur compounds (alliin, diallyl disulfide, dipropyl disulfide, glutathione), terpenoids (apiole, geraniol, limonene) and alkaloids (boldine, caffeine) (**Table 1**). About 30% of non-phenolic antioxidants were present in all vegetables. These were organic acids (acetoacetic, adipic, citric, malic), amines (putrescine, spermidine, glucosamine), amino acids (cystationine, tyrosine) and pyridines (pyridoxine 5-phosphate and pyridoxamine).

To classify vegetables with respect to their antioxidant composition, we performed hierarchical cluster analysis on mass datasets (**Fig. 2**). Vegetables were clustered into 3 major subgroups: i) cabbage and broccoli, ii) beans and peas and iii) endive and lettuces. Analysis demonstrated higher similarity of antioxidant profiles in endive and lettuces (short dendrogram arms), which reflects the close systematic relationship between these species. Lower similarity

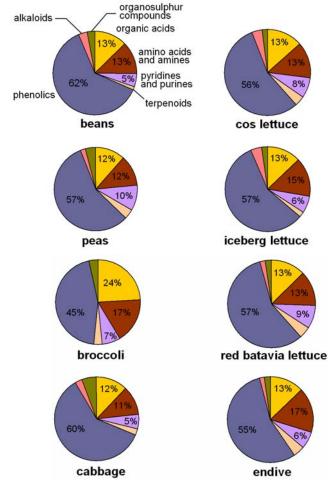


Fig. 1 Ratios between metabolic classes of antioxidants in vegetables. The numbers refer to percentage of compounds of each class.

(longer dendrogram arms) between beans and peas may reflect interspecies differences. Low similarity between cabbage and broccoli may result from different edible organs used for analysis (florets in broccoli and leaves in cabbage).

To characterise the distribution of antioxidants in vegetables we performed frequency analysis on all samples used in this study (**Fig. 3**). Analysing frequencies is important because of variation in the metabolite composition of samples taken from the same plants. Because different numbers

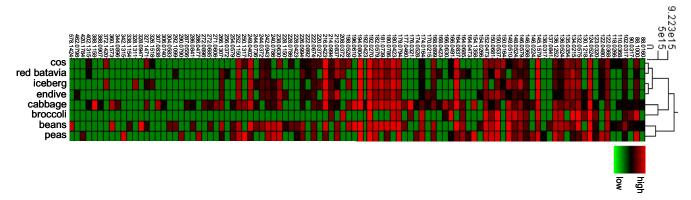


Fig. 2 Hierarchical cluster analysis of antioxidant profiles in vegetables. The dataset includes antioxidant compounds found in positive and negative ion modes (Table 1). Clustering was performed along the columns using the Euclidean distance and division cluster linkage method. The colours indicate global variation in relative peak intensities. Peak intensities were normalized with respect to internal calibrants and averaged for each variety to reduce the dataset size (originally 120 columns).

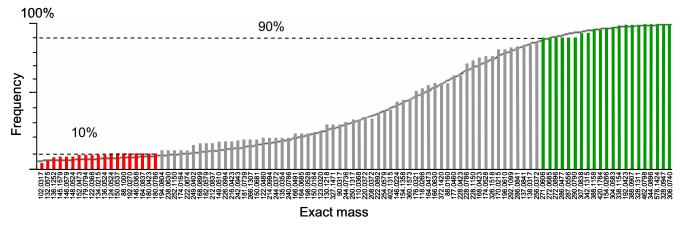


Fig. 3 Frequencies of antioxidant appearance in vegetables. Bars represent summed weighted frequencies of antioxidant compounds from 40 vegetable samples, fitted by sigmoidal function.

of samples were taken from different vegetables, the sums of weighted frequencies for each variety were calculated. Frequencies were plotted in ascending order and fitted by sigmoidal (Boltzmann) function. Three groups of compounds were derived from this analysis: 1) compounds appearing with frequency $\geq 90\%$; 2) compounds with frequency between 10 and 90%; and 3) compounds with frequency $\leq 10\%$. The first group of antioxidants, which we called ubiquitous, consists of 21% of all assigned antioxidants. It includes compounds from different metabolic classes, such as organic acids (acetoacetic, adipic, citric, malic), amines (putrescine, spermidine, glucosamine), pyridines (pyridoxine 5-phosphate and pyridoxamine), terpenoids (limonene), organosulfur compounds (dipropyl disulfide) and phenolic compounds (cinnamic acid, cinnamaldehyde, coumarin, vanillin, hinokiresinol, etc.). The second group, called common antioxidants, contains 60% of all compounds, representing all 7 metabolic categories. The third group (rare antioxidants) accounts for 19% of assigned compounds and contains only phenolic metabolites, such as flavonoids (e.g. gallocatechin, salvigenin, betanidin) and coumarins (decursin and scopoletin). Our analysis demonstrates that ubiquitous antioxidants include primary metabolites, as well as key compounds in secondary metabolic pathways, while rare antioxidants represent the end products of secondary metabolism.

DISCUSSION

Fourier-transform ion cyclotron resonance mass spectrometry has recently been proved to provide a powerful tool for plant metabolomics and metabolic profiling (Oikawa *et al.* 2006, Ohta *et al.* 2007; Takahashi *et al.* 2008). This mass instrument offers sufficient mass resolution and mass accuracy for identifying compounds, present in complex plant samples, which usually require preliminary chromatographic separation. Progress in the application of this technology was limited by the high running cost of FTICR mass spectrometer and the absence of information resources for identification of plant metabolites. The latter was successfully addressed by developing a comprehensive internetbased reference database KNApSAcK (Shinbo *et al.* 2006), containing information about ~23000 plant metabolites (http://kanaya.naist.jp/KNApSAcK/). Examples of FTICR-MS application for metabolomic analysis and metabolite profiling in plants are presented in papers by Aharoni *et al.* (2002), Mungur *et al.* (2005), Oikawa *et al.* (2006), Ohta *et al.* (2007), Takahashi *et al.* (2008).

The purpose of this study was to apply the FTICRMS for profiling antioxidant compounds in vegetables. Although FTICRMS provides a fast and efficient tool for screening plant metabolites, some limitations of this approach should be mentioned. 1. As shown in Table 1, the empirical formula derived from mass analysis can be assigned to several structurally similar antioxidants. 2. Although quantitative measurements of metabolites using FTICRMS have been previously demonstrated (Bristow et al. 2005), quantification of all compounds in complex metabolite mixtures is not yet possible. This problem is thought to arise from several effects, such as ion suppression and different ionization of molecules. It has been shown that ion suppression can be minimised by diluting the samples and accumulating a large number of mass scans (Kujawinski et al. 2002). Therefore, in our study we used very diluted samples and hundreds of mass scans to produce one spectrum. It should be mentioned that since many antioxidants can be oxidised during sample processing, factors such as temperature and pH should be carefully considered when selecting the metabolite isolation procedure.

Antioxidants are chemical compounds which delay the start or slow the rate of free radical formation. These compounds are characterised by the ability to donate the hydrogen atom or electron/proton, or ability to chelate metal ions, involved in formation of reactive oxygen species, or both (donating and chelating). In most cases, antioxidant action involves a combination of different mechanisms, therefore antioxidant properties cannot be attributed to a certain class of chemical compounds or to certain functional groups in these compounds. Our study demonstrates that antioxidants occur among different classes of plant metabolites such as organic acids, phenolic acids, sulphur-containing amino acids, polyamines, pyridines, flavonoids, lignans, couma-rins, stilbenes and alkaloids. Although ascorbic acid, carotenoids and tocopherols are often cited as the 'major' antioxidants in plants, the largest group of antioxidant compounds in vegetables are represented by phenolics (flavonoids, phenolic acids, lignans, coumarins, stilbenes, phenylpropanoid gycosides and quinines). The second large group of antioxidants in vegetables are organic acids (Fig. 1). Ascorbic acid, which belongs to this group, is, however, less abundant in vegetables than adipic, citric and malic acids (Table 1). Although some antioxidants, such as ascorbic acid and glutathione, are ubiquitous in growing plants (Noctor and Foyer 1998), these compounds seem to be unstable during the postharvest storage of vegetables. For example, ascorbic acid content in broccoli was shown to decrease by up to 70 %, depending on storage duration, temperature and packaging (Podşedek 2007). Loss of antioxidant compounds in vegetables is associated with postharvest senescence induced by wounding and disruption of energy, nutrient and hormone supply. This involves production of ethylene and free radicals, which trigger lipid peroxidation and degradation of chlorophyll and proteins. Antioxidants are involved in scavenging reactive oxygen species produced during senescence, therefore their levels in vegetables often decrease upon harvesting. Postharvest stability of antioxidants varies among plant species. Cruciferous vegetables retain ascorbic acid better than other vegetables (Lee and Kader, 2000). Our previous study has demonstrated that white cabbage retained about 50% of ascorbic acid after 6 months of storage (Hounsome et al. 2009). In agreement with literature reports, we found cabbage and broccoli to be a good source of ascorbic acid. The fact that ascorbic acid was not found in other vegetables most likely reflects the loss of this compound during the postharvest period, rather than problems with FTICRMS detection.

Some compounds, such as sulphur-containing amino acids and polyamines are usually not mentioned in literature as dietary antioxidants. However, these compounds are highly abundant in plant tissues, and play an important role in antioxidant defence system in humans (Løvaas 1997; Meucci and Mele 1997). Amino acids homocysteine, methionine, tyrosine and cystationine, polyamines putrescine and spermidine, and sugar amine glucosamine were found in almost all tested vegetables (Table 1). Another group of plant metabolites, organosulphur compounds, are mainly known due to glucosinolates, which do not possess antioxi-dant properties. In this study we found several organosulphur compounds with reported antioxidant activities: diallyl disulfide, dipropyl disulfide, alliin and glutathione (Table 1). In addition to organosulphurs, two other non-phenolic antioxidants, alkaloid caffeine and terpenoid limonene were found in all samples.

In recent years the focus of research has shifted from using antioxidant supplements towards a balanced combination of antioxidants provided by diet (Liu 2003). Despite the large amount of information on the beneficial effects of antioxidants, randomised clinical trials have failed to show any consistent benefit from the use of antioxidant supplements on cardiovascular disease or cancer risk, with some trials even suggesting possible harm in certain subgroups (Stanner *et al.* 2004). These trials have usually involved the administration of single antioxidant nutrients given at relatively high doses. A number of studies have demonstrated complex interactions between antioxidant compounds, known as synergistic effect. It has been shown that mixtures of antioxidants have higher antioxidant activity than individual antioxidants. For example, mixtures of carotenoids were more effective against oxidative damage than the single compounds (Stahl et al. 1998). Synergistic interactions have been also demonstrated between phenolic acids, beta-carotene, and ascorbic acid, as well between flavonoids and tocopherol (Trombino et al. 2004; Marinova et al. 2008). Although the mechanism of these interactions is not well understood, the data demonstrate the importance of complex assessment of antioxidant compounds in human foods. In this respect metabolite profiling can provide valuable information about antioxidant composition and distribution in vegetables. This study attempts to provide an overview of the whole range of antioxidants present in vegetables. FTICRMS is a powerful tool for highthroughput antioxidant profiling in plants. However, further development of metabolite identification and quantification is essential for comprehensive analysis of plant antioxidants.

CONCLUSIONS

In this study we analysed antioxidants profiles in 8 vegetables: beans, broccoli, cabbage, endive, peas and 3 varieties of lettuces using Fourier-transform ion cyclotron resonance mass spectrometry. Our data demonstrates that:

- 1. Electrospray ionisation FTICRMS provides a powerful tool for antioxidant profiling in plants;
- Antioxidant compounds are found among different classes of plant metabolites, such as phenolics, organic acids, amino acids and amines, pyridines and purines, terpenoids, alkaloids and organosulphur compounds;
- Phenolic compounds are the most numerous metabolic group, which accounts for 45-62% of all antioxidant compounds, followed by organic acids (12-24%), amino acids and amines (11-17%), pyridines and purines (5-10%), organosulphur compounds, terpenoids, and alkaloids (<5%)
- 4. With respect to occurrence in vegetables, antioxidants can be classified as ubiquitous (appearance frequency >90%), common (appearance frequency between 10% and 90%), and rare (appearance frequency <10%).

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