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British Journal of Nutrition / Volume 95 / Issue 01 / January 2006, pp 51 - 58  
DOI: 10.1079/BJN20051596, Published online: 08 March 2007

**Link to this article:** [http://journals.cambridge.org/abstract\\_S0007114506000079](http://journals.cambridge.org/abstract_S0007114506000079)

### How to cite this article:

Manal Abd El Mohsen, Joanne Marks, Gunter Kuhnle, Kevin Moore, Edward Debnam, S. Kaila Srail, Catherine Rice-Evans and Jeremy P. E. Spencer (2006). Absorption, tissue distribution and excretion of pelargonidin and its metabolites following oral administration to rats. *British Journal of Nutrition*, 95, pp 51-58 doi:10.1079/BJN20051596

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## Absorption, tissue distribution and excretion of pelargonidin and its metabolites following oral administration to rats

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(Received 21 April 2005 – Revised 23 June 2005 – Accepted 4 August 2005)

Recent reports have demonstrated various cardiovascular and neurological benefits associated with the consumption of foods rich in anthocyanidins. However, information regarding absorption, metabolism, and especially, tissue distribution are only beginning to accumulate. In the present study, we investigated the occurrence and the kinetics of various circulating pelargonidin metabolites, and we aimed at providing initial information with regard to tissue distribution. Based on HPLC and LC-MS analyses we demonstrate that pelargonidin is absorbed and present in plasma following oral gavage to rats. In addition, the main structurally related pelargonidin metabolite identified in plasma and urine was pelargonidin glucuronide. Furthermore, *p*-hydroxybenzoic acid, a ring fission product of pelargonidin, was detected in plasma and urine samples obtained at 2 and 18 h after ingestion. At 2 h post-gavage, pelargonidin glucuronide was the major metabolite detected in kidney and liver, with levels reaching 0.5 and 0.15 nmol pelargonidin equivalents/g tissue, respectively. Brain and lung tissues contained detectable levels of the aglycone, with the glucuronide also present in the lungs. Other tissues, including spleen and heart, did not contain detectable levels of pelargonidin or ensuing metabolites. At 18 h post-gavage, tissue analyses did not reveal detectable levels of the aglycone nor of pelargonidin glucuronides. Taken together, our results demonstrate that the overall uptake of the administered pelargonidin was 18% after 2 h, with the majority of the detected levels located in the stomach. However, the amounts recovered dropped to 1.2% only 18 h post-gavage, with the urine and faecal content constituting almost 90% of the total recovered pelargonidin.

### Anthocyanidin: Pelargonidin: Absorption: Distribution: Metabolism

As part of the flavonoid family, anthocyanins are polyphenolic compounds that are widely distributed among pigmented fruit and vegetables. Anthocyanidins are particularly abundant in berries, such as blueberry and blackcurrant, as well as in red wine, thus their presence in the diet can be significant. As potential components of the human diet, previous research on anthocyanidins has concentrated on their biological activities, and included the assessment of possible health benefits. For example, previous dietary interventions in man and animals have demonstrated that the consumption of anthocyanin-rich red wine (Renaud & de Lorgeril, 1992) or blueberry extract (Joseph *et al.* 1999) may exert beneficial effects by lowering the risk of CHD, and by retarding age-related declines in neurological function, respectively.

To evaluate the health benefits of anthocyanidins in man it is important to understand their absorption, metabolism, tissue distribution and excretion. Several studies have shown that anthocyanins are absorbed as glycosides that could be

recovered from plasma and urine after oral administration (Cao & Prior, 1999; Miyazawa *et al.* 1999). However, the bioavailability of anthocyanins is very limited and their metabolism is still not fully understood. In addition, only a few studies (Tsuda *et al.* 1999; Matsumoto *et al.* 2001) have quantitatively evaluated the absorption of anthocyanins using purified compounds (directly administered using a defined dose) as opposed to feeding enriched foods or food extracts. To date, there are few data available with regard to anthocyanidin distribution among various body tissues where they could exert their biological effects through antioxidant potential, by association with the cell membrane or through direct intracellular mechanisms. Anthocyanidins are known to be stable under acidic conditions but rapidly broken down under neutral conditions (Brouillard, 1988) such as those encountered *in vivo*. It is thus necessary to characterise and to quantify anthocyanin metabolites, including ring-fission products, as well as the ingested form of anthocyanidins. In the present study, we

**Abbreviation:** TFA, trifluoroacetic acid.

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investigated metabolism and distribution of the strawberry anthocyanin, pelargonidin, after gavage into rats, and assessed the effect of time on the nature and the circulating levels of its metabolites.

## Experimental methods

### Animals

Male Sprague-Dawley rats (250 g) were obtained from the Comparative Biology Unit at the Royal Free and University College Medical School, and maintained on standard rat chow (Diet RM1; SDS Ltd, Witham, UK). Two groups of rats ( $n = 6$ ) were administered pelargonidin (Extrasynthese, purity 99.82%), dissolved in 50% aqueous ethanol, in a dose of 50 mg/kg body weight and killed 2 or 18 h post-gavage. Three animals served as control group and received only 50% aqueous ethanol. In all groups, animals were fasted for 4 h before pelargonidin administration.

At the selected times, animals were anaesthetised by intraperitoneal injection of pentobarbitone sodium (90 mg/kg; Sagital; Rhone-Merieux, Harlow, UK). Heparin (0.3 ml, 5000 IU/ml; CP Pharmaceuticals Ltd, Wrexham, UK) was injected via a tail vein injection to reduce blood clotting followed by collection of 2 ml blood by cardiac puncture into heparinised tubes. Animals were exsanguinated with 200 ml ice-cold heparinised 0.9% (w/v) saline and the blood-free organs were dissected out immediately. Plasma samples were separated by centrifugation at 600 g for 30 min and immediately acidified with 0.44 M-trifluoroacetic acid (TFA) in the ratio 5:1. The Hb concentrations in the cardiac blood sample and exsanguinated fluid was measured using a Hb assay kit (Sigma, Poole, UK) to allow calculation of blood volume. The removed tissues, liver, kidneys, spleen, heart, small intestine, large intestine, lung and brain, were weighed and then homogenised in TFA (10 volumes/g tissue) using a Ultra Turrax homogeniser (Janke & Kunkel, Staufen, Germany) for 40 s. Urine and faeces from rats were collected in metabolic cages containing 1.5 ml TFA to prevent the degradation of pelargonidin. Intestinal contents were drained by finger pressure, flushed with TFA, and contents kept for analysis.

### Sample preparation

Plasma (150  $\mu$ l) was de-proteinised with 30  $\mu$ l 20% (w/v) TCA then extracted with 200  $\mu$ l ice-cold methanol. HPLC analysis of re-extracted pellet showed no detectable retained material. Solvents contained in the extract obtained after centrifugation were evaporated under N and the residue was dissolved in 250  $\mu$ l 20% (v/v) methanol in 0.44 M-TFA for quantitative HPLC analysis. An additional aliquot of plasma was loaded on to an ODS solid extraction cartridge (Sep-Pak C18; Waters, Milford, MA, USA) after the column was washed with 3 ml methanol and equilibrated with 6 ml water-methanol-acetic acid (94:5:1, by vol.) prior to the application of the sample. Following the sample application, the cartridge was washed with the same equilibrating mixture and the anthocyanins were eluted from the cartridge material with 2 ml methanol. The extract was dried under N, and the residue dissolved in 250  $\mu$ l 20% (v/v) methanol in 0.44 M-TFA.

Urine samples were filtered through a micropore filter (Millex-GP, 0.22 mm filter unit; Millipore, Billerica, MA, USA). Extraction of pelargonidin and its metabolites from rat tissues was performed by mixing the homogenates with an equal volume of ice-cold methanol followed by vigorous vortexing for 1 min. The methanolic extract obtained after centrifugation was dried under vacuum at 30°C by rotary evaporation and then the extracts were treated as described earlier.

Quantitative determination of pelargonidin and its metabolites was based on external standards. An internal standard, rutin, was added to the samples before extraction and to the external standards to a final concentration of 5  $\mu$ M. Calibration curves were conducted over the range of 0.5–20  $\mu$ mol/l. Peak area ratios (compound/internal standard) were plotted against concentration of the compound/metabolite. The concentration of pelargonidin glucuronide was calculated using peak area of standard pelargonidin aglycone and expressed as pelargonidin equivalents as previously used for calculation of anthocyanidin metabolites (Felgines *et al.* 2002; Wu *et al.* 2002). Calibration curves obtained were linear over the entire range with correlation coefficient values  $\geq 0.995$ .

### Methods of analysis

HPLC analysis was undertaken using a Waters system (Milford, MA, USA) consisting of controller 600, auto sampler 717 plus, photodiode array detector 996 and on-line degasser. Samples were analysed on a Zorbax SB-C18 column (4.6  $\times$  250 mm, with 5  $\mu$ m particle size and a guard column of the same material, 4.6  $\times$  15 mm). Column temperature was set at 30°C. Mobile phase A consisted of 5% formic acid aqueous solution and mobile phase B of pure methanol. The gradient applied was as follows: from 0 to 5 min 95% A and 5% B, from 5 to 40 min to 100% B, from 40 to 45 min 100% B, and from 45.1 min 90% A and 10% B. Run time was 60 min followed by a 10 min delay prior to the next injection. Components were identified according to retention times and UV/visible spectra. Detection of pelargonidin was undertaken at 520 nm, internal standard at 300 nm and phenolic acids at 260 nm. LC-MS analysis was employed using a Finnigan LCQ Deca XP quadrupole ion trap mass spectrometer (San Jose, CA, USA). Separation was performed using a C18 column (50  $\times$  2.5 mm; Phenomenex, Schlieren, Switzerland) with the following gradient (phase A: 0.1% formic acid in water, phase B: 50% acetonitrile in water, 0.1% formic acid). The LC-MS gradient was 0–5 min 100% A, 5–40 min from 100% A to 50% A, 40–60 min to 0% A, 60–65 min 0% A. Compounds were detected using a full ion scan and identified by performing product ion scans on selected ions. GC-MS analysis was used as described previously (Rechner *et al.* 2002) to confirm the identity of phenolic acids detected by HPLC.

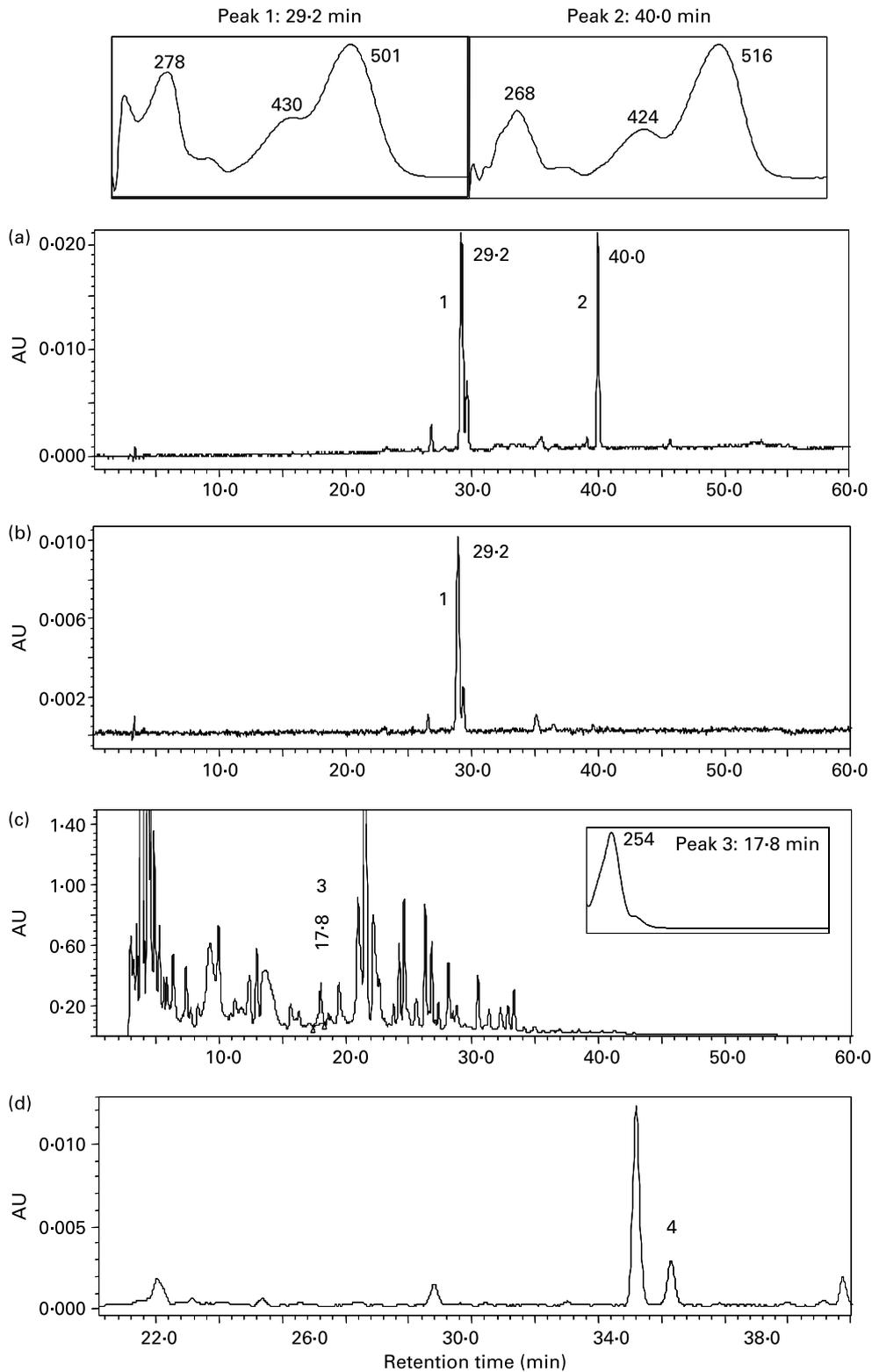
### Calculations and statistics

Values are given as means with their standard errors of the mean. Statistical analysis was performed using an unpaired *t* test (Instat) and considered significant at  $P < 0.05$ .

## Results

The present study investigated the metabolism of pelargonidin after oral administration to rats. HPLC analysis with

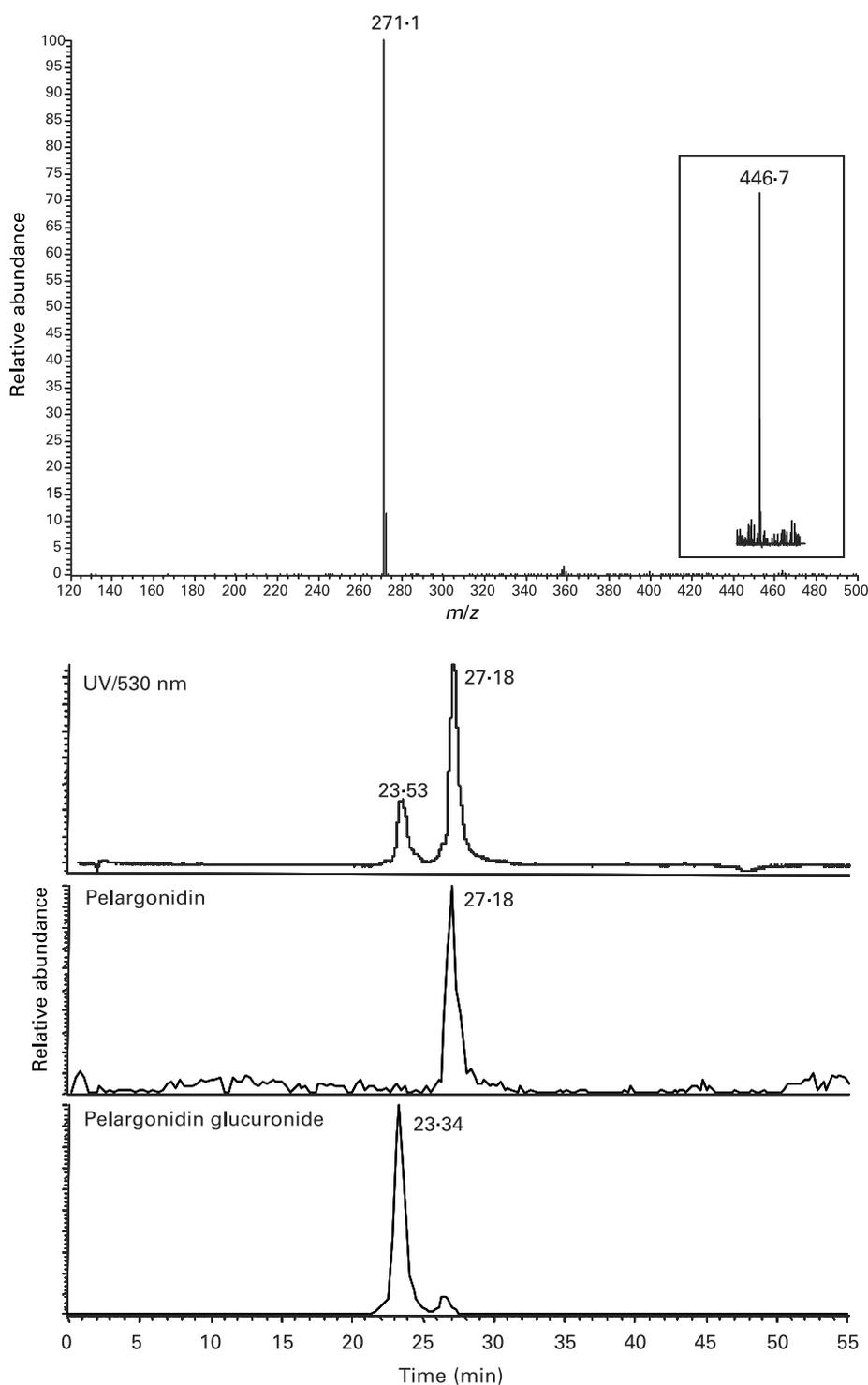
photodiode array detection was employed to identify the nature of metabolites (Fig. 1). Whereas no anthocyanins were detected in untreated controls, the oral administration of pelargonidin to rats led to the detection of a circulating



**Fig. 1.** Representative chromatograms of tissues obtained from pelargonidin-fed rats. (a), Small intestine extract at 520 nm: peak 1, pelargonidin-O- $\beta$ -D-glucuronides; peak 2, pelargonidin. (b), Plasma extract at 520 nm. (c), Urine at 260 nm: peak 3, *p*-hydroxybenzoic acid. (d), Plasma at 260 nm: peak 4, unidentified. AU, absorbance units.

anthocyanidin at 2 h post-ingestion. This compound was detected by HPLC, and eluted at a retention time of 40.0 min (peak 2), which corresponds to the retention time of free pelargonidin aglycone. In addition, a major peak at 29.2 min (peak 1), which partially co-eluted with a minor peak, was also detected in the gastrointestinal tract, plasma

and urine. Similarities regarding their absorbance spectra as well as the difficulties we experienced in attempting to completely separate these two peaks using various HPLC gradient conditions may indicate that these compounds are structurally related. Both compounds were more hydrophilic than pelargonidin, and thus were predicted to represent glucuronidated



**Fig. 2.** MS/MS trace of small intestine extract obtained from pelargonidin-gavaged rats. The trace shows pelargonidin ( $m/z = 271$ ) eluting at 27.18 min and pelargonidin-*O*-glucuronide ( $m/z = 447$ ) with a retention time of 23.34 min. The spectrum shows a single fragment at  $m/z = 271$ , the  $M^+$  ion of the aglycone, indicating the neutral loss of glucuronic acid. The insert shows the spectrum of the base peak ( $M^+$ ).

derivatives of pelargonidin. Thus, a subsequent LC-MS/MS analysis of the isolated peaks was undertaken to confirm the molecular mass for pelargonidin glucuronides (Fig. 2). Two major peaks are detected at 530 nm, one for pelargonidin with  $m/z = 271$  (retention time 27.18 min), preceded by a more polar glucuronide signal  $m/z = 447$  (retention time 23.53 min) (Fig. 2). The product ion spectrum of the major pelargonidin-glucuronide shows the neutral loss of glucuronic acid, and therefore the major peak is the  $M^+$  ion of the aglycone.

HPLC chromatograms of urine samples collected for 18 h post-gavage at 260 nm (Fig. 1(c)) also revealed the presence of peak 3, eluting at 18 min. This peak was identified as *p*-hydroxybenzoic acid by comparison with an authentic standard compound, and based on retention times and UV spectra as well as by GC-MS analysis following derivatisation (Fig. 3). This phenolic acid was also detected in plasma samples obtained at 2 or 18 h post-ingestion. Fig. 4 shows two modes of the C-ring fission of pelargonidin, which yield ultimately the detected product, *p*-hydroxybenzoic acid. The dotted line shows breakage of the C-ring between C8a-O1 and C2-C3 leading to formation of *p*-hydroxybenzoic acid directly. However, as shown by the dashed line, breakage of the C-ring can also occur between O1-C2 and C4-C4a, leading to the formation of *p*-hydroxyphenyl propionic acid, which could be subjected to  $\beta$ -oxidation leading to formation of *p*-hydroxybenzoic acid, as previously described by Scheline (1991).

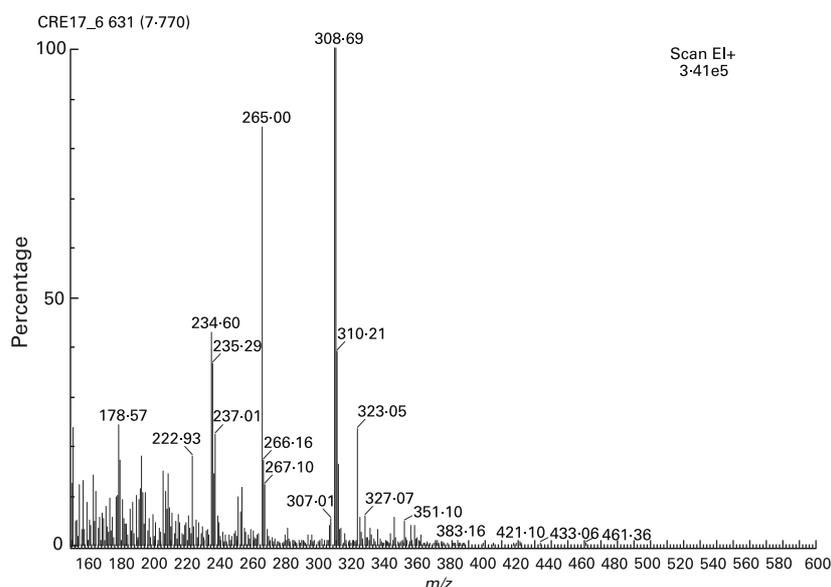
In addition, 2 h plasma samples revealed the presence of an additional peak eluting at 35.2 min (Fig. 1(d)), which was found to be present at lower levels after 18 h, but which was not detectable in any of the control samples. This finding may be indicative for the presence of another metabolic product resulting from pelargonidin intake. However, we were not able to structurally identify this compound as a pelargonidin metabolite.

Levels of pelargonidin metabolites at 2 and 18 h were calculated in plasma and tissues and expressed as nmol

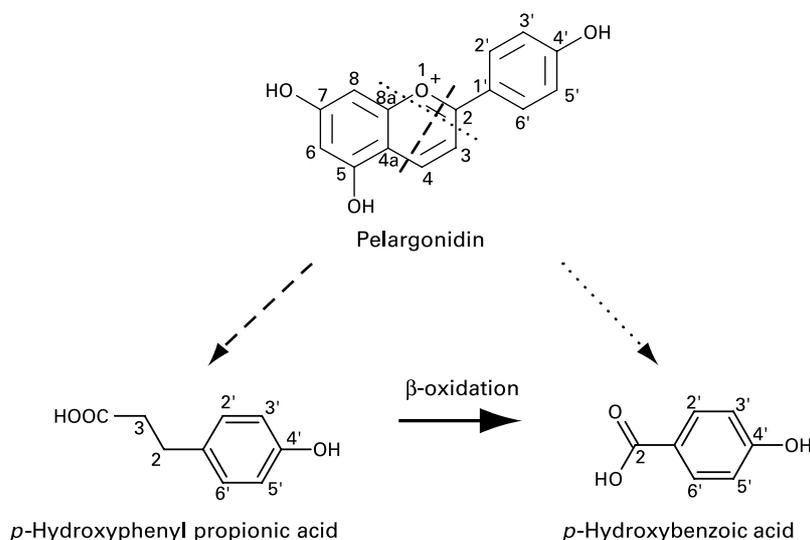
pelargonidin equivalents/ml plasma or g tissue (Fig. 5). At 2 h post-gavage, pelargonidin glucuronide was the major metabolite detected in kidney and liver, with levels reaching 0.5 and 0.15 nmol pelargonidin equivalents/g tissue, respectively (Fig. 5(a)). Whereas the brain and the lungs showed the aglycone as the main metabolite with similar levels reaching 0.16 nmol/g, in other tissues, mainly the spleen and heart, no compounds were detected. At 18 h post-gavage, our tissue analyses did not reveal the presence of any detectable levels of pelargonidin aglycone or its glucuronides. However, the colonic metabolite *p*-hydroxybenzoic acid was present in plasma, reaching 44% of total metabolites detected at 2 h post-gavage, and being the only metabolite detected in plasma at 18 h post-gavage (Fig. 5(b)).

Table 1 shows the levels of metabolites detected in different parts of the gastrointestinal tract at 2 and 18 h following the administration of pelargonidin to rats. Analyses of the stomach content following the administration of pelargonidin demonstrated the presence of *p*-hydroxybenzoic acid, which accounts for approximately 11% of the detected compounds. High levels of glucuronides were detected in the small intestine tissue extract 2 h post-gavage. At 18 h post-gavage, these metabolites disappeared completely from the small intestine, which would be consistent with the time-dependency of passage through the gastrointestinal tract. However, *p*-hydroxybenzoic acid was detected in the large intestine, both at 2 and 18 h post-gavage (Table 1).

Analysis of urine samples obtained by aspirating the urine directly from the rat bladder 2 h post-gavage demonstrated the presence of pelargonidin glucuronide as the only metabolite excreted into the urine at this time. However, *p*-hydroxybenzoic acid constituted approximately 50% of metabolites found in urine collected for 18 h post-gavage (Table 2). In contrast to the urine, the glucuronide conjugates were not found in any of the faecal samples, with *p*-hydroxybenzoic acid as the major product. The results reveal that the total recovery of the administered pelargonidin was only 18%

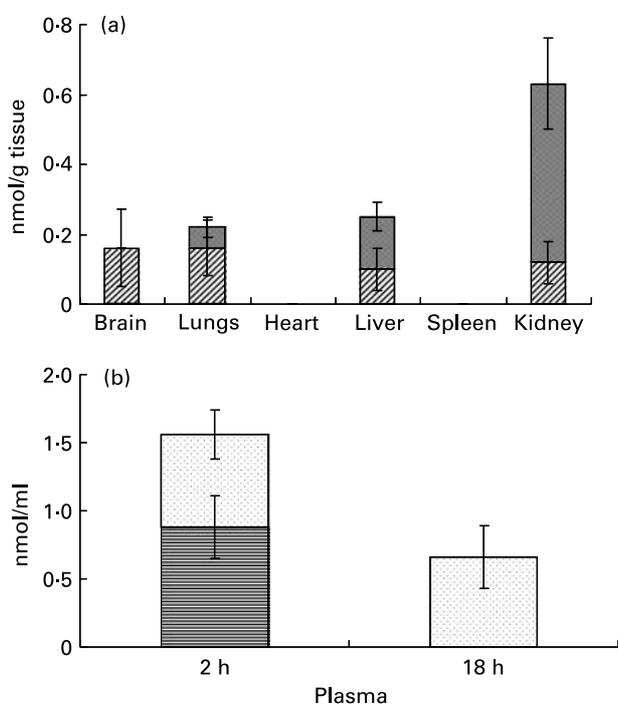


**Fig. 3.** Mass spectra of the derivatised derivatives of *p*-hydroxybenzoic acid as detected in urine. The spectra shows a characteristic fragment at  $m/z = 309$ , resulting from the loss of a *t*-butyl group of the derivatised compound. The identification was done by comparison with the spectra of the purified standard.



**Fig. 4.** Possible mechanism for degradation of pelargonidin leading to formation of *p*-hydroxybenzoic acid.  $\cdots \blacktriangleright$ , breakage of C-ring between C8a-O1 and C2-C3 leading to formation of *p*-hydroxybenzoic acid.  $-\cdots \blacktriangleright$ , breakage between O1-C2 and C4-C4a leading to the formation of *p*-hydroxyphenyl propionic acid.

after 2 h, where the majority of the detected levels were located in the stomach; however, the amounts recovered dropped to 1.2% only 18 h post-gavage, with the urine and faecal content constituting almost 90% of the total recovered pelargonidin.



**Fig. 5.** Concentrations of pelargonidin and its metabolites (calculated as nmol pelargonidin equivalents) in rat tissues obtained 2 h (a) and plasma obtained at 2 and 18 h (b) after oral administration of 50 mg/kg body weight of pelargonidin. For details of procedures, see p. 52. (a), Pelargonidin in free (■) and conjugated forms (▨); (b), pelargonidin in conjugated forms (▨) and *p*-hydroxybenzoic acid (▤). Values are means with their standard errors depicted by vertical bars (six animals per group).

## Discussion

Recently, much attention has been given to the biological functions of flavonoids. Foods that are rich in anthocyanins have been shown to exert multiple beneficial health effects. In this context, the bioavailability of anthocyanins represents a timely question that we were aiming to address by conducting the present study on the absorption, metabolism, distribution and excretion of pelargonidin.

Our results demonstrated that pelargonidin was absorbed from the gastrointestinal tract and it appeared in the plasma and urine mainly as a glucuronidated conjugate. It has been well documented previously that anthocyanidins are absorbed, in both man and animals, as their intact glycosidic forms present in fruit (Miyazawa *et al.* 1999; Tsuda *et al.* 1999). Despite this, various glucuronide conjugates of cyanidin (Wu *et al.* 2002) as well as pelargonidin (Felgines *et al.* 2003) have been detected in human plasma previously following the intake of elderberries and strawberries. The mechanism of glucuronide formation from the anthocyanidin glucoside was reported to involve the conversion of the glucoside directly into its corresponding glucuronide (Wu *et al.* 2002) by the action of UDP-glucose dehydrogenase. Furthermore, anthocyanidin glucosides can be hydrolysed yielding the aglycone, which can be subject to glucuronidation in the intestine and liver. Interestingly, the presence of cyanidin aglycone has been previously reported in the rat jejunum after ingestion of cyanidin-3-glucoside (Tsuda *et al.* 1999). The ingestion of pelargonidin aglycone and the ensuing detection of circulating pelargonidin glucuronides, in the present study, provides evidence that the flavylum cation structure does not impart resistance to enzymatic conversion by UDP-glucuronosyl transferase, as previously assumed by Miyazawa *et al.* (1999).

In spite of detecting pelargonidin and its glucuronide in rat plasma and tissues 2 h after administration, neither the native compound nor its metabolites were found 18 h after ingestion. A pharmacokinetic study regarding the absorption of anthocyanins has previously demonstrated that these compounds

**Table 1.** Total amounts of metabolites (nmol) detected in the gastrointestinal content, 2 and 18 h following the oral administration of 50 mg/kg body weight of pelargonidin to rats\*

Mean values with their standard errors (six animals per group)

	2 h						18 h					
	Pelargonidin aglycone		Pelargonidin glucuronide		p-HBA		Pelargonidin aglycone		Pelargonidin glucuronide		p-HBA	
	Mean	SEM	Mean	SEM	Mean	SEM	Mean	SEM	Mean	SEM	Mean	SEM
Stomach	6131	995	ND	ND	790	209	ND	ND	ND	ND	ND	ND
Small intestine	195	62.1	189	41.6	232	65.5	ND	ND	ND	ND	ND	ND
Large Intestine	8.7	6.1	0.25	0.17	39.7	35.1	ND	ND	ND	ND	52.5	31.5

ND, non-detectable; p-HBA, *p*-hydroxybenzoic acid.

\* For details of procedures, see p. 52.

are rapidly absorbed; reaching maximum plasma concentrations of 3.49  $\mu$ M at 15 min after a single oral dose was administered to rats. Thereafter the plasma concentration fell rapidly reaching 0.14  $\mu$ M at 240 min, and thus suggesting a relatively short half-life (Miyazawa *et al.* 1999). Rapid absorption could be attributed to the involvement of stomach-dependent mechanisms, possibly through a bilitranslocase-mediated mechanism as indicated following administration of grape anthocyanins to rats (Passamonti *et al.* 2003). The high instability of anthocyanidins at physiological pH and their rapid decomposition to ring-opened forms and/or degradation products is probably responsible for the relatively short half-life in biological fluids. Indeed, it was recently shown that the flavylium cation form is not regenerated by acidification following incubation of aglycones with control urine or plasma (Felgines *et al.* 2002).

One of the major findings of the present study was the identification of *p*-hydroxybenzoic acid as an important metabolite of pelargonidin in the rat plasma and gastrointestinal tissues. Despite the fact that this metabolite could also be derived from the metabolism of other compounds such as aromatic amino acids, the high levels detected post-gavage of pelargonidin, as compared with levels detected in control animals, support the notion that this phenolic acid commonly arises from the metabolism of pelargonidin. Indeed, a structurally similar metabolite, namely protocatechuic acid, was detected following an oral cyanidin glucoside administration to rats, with plasma levels reaching eight times that of the anthocyanidin (Tsuda *et al.* 1999). The detection of *p*-hydroxybenzoic acid in stomach extract and plasma 2 h after gavage indicates that its formation likely results due to the instability and degradation of the anthocyanidin. However, phenolic acid derivatives may also be formed by the colonic

**Table 2.** Total amounts of pelargonidin and its metabolites (nmol) excreted in rat urine and faeces collected for 18 h post-gavage\*

Mean values with their standard errors (six animals per group)

	Urine		Faeces	
	Mean	SEM	Mean	SEM
Pelargonidin	ND	–	182	14.7
Pelargonidin glucuronide	158	14.8	ND	–
<i>p</i> -Hydroxybenzoic acid	155	31.3	161	15.7

ND, non-detectable.

\* For details of procedures, see p. 52.

microflora. A comparison of the metabolites formed after oral and intraperitoneal administration of catechin, and the suppression of their formation after administration of antibiotics, confirms the catabolising activities of colonic microflora (Das & Griffiths, 1968). In addition, it was shown recently that *in vitro* incubation of anthocyanins with human faecal suspension yields phenolic acids as major products (Fleschhut *et al.* 2005). Our results, demonstrating that a similar pathway exists *in vivo*, clearly point to the importance of colonic metabolism of flavonoids. Indeed, these metabolites might be equally as important since anticancer properties of aromatic acids have been previously documented (Samid *et al.* 1997; Thibout *et al.* 1999).

Considering all pelargonidin metabolites detected, the percentages of products excreted over 18 h accounted only for 1.2% of the gavaged dose. Similar findings were also reported where total urinary excretion of blackberry anthocyanidins in rats (Felgines *et al.* 2002) as well as strawberry anthocyanin metabolites in human subjects (Felgines *et al.* 2003) corresponded to only 0.93 and 1.8% of the anthocyanidins ingested, respectively, pointing to the existence of other non-detectable metabolites.

In conclusion, the present results suggest the importance of investigating the biological activities of the degradation products, which could be formed at a later stage, possibly in the colon, when evaluating the properties of diet-derived anthocyanidins.

## Acknowledgements

The authors acknowledge the Food Standards Agency and the Biotechnology and Biological Sciences Research Council (grant no. BB/C518222/1) for financial support on this project.

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