Bioavailability of procyanidin dimers and trimers and matrix food effects in \textit{in vitro} and \textit{in vivo} models

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Among procyanidins (PC), monomers, such as catechin and epicatechin, have been widely studied, whereas dimer and trimer oligomers have received much less attention, despite their abundance in our diet. Recent studies have showed that as dimers and trimers could be important in determining the biological effects of procyanidin-rich food, understanding their bioavailability and metabolism is fundamental. The purpose of the present work is to study the stability of PC under digestion conditions, the metabolism and the bioavailability by using a combination of \textit{in vitro} and \textit{in vivo} models. Simultaneously, the matrix effect of a carbohydrate-rich food on the digestibility and bioavailability of PC is investigated. The results show a high level of stability of PC under gastric and duodenal digestion conditions. However, the pharmacokinetic study revealed limited absorption. Free forms of dimers and trimers have been detected in rat plasma, reaching the maximum concentration 1 h after oral intake of a grape seed extract.

**Procyanidins: Dimers: Trimers: Digestion: Bioavailability**

Procyanidins (PC) are found in most plants and in a wide range of foods, such as red wine, cocoa, tea and fruits, and thus they are a part of the human diet. PC belong to the group of flavonoids, and are phenolic compounds mainly formed of (+)-catechin and (-)-epicatechin units with $C_8-C_8$ and/or $C_4-C_6$ bonds. PC are formed from the condensation of monomeric units, between two and five units for oligomers and over five units for polymers\textsuperscript{(1)}.

Grape seed procyanidin extracts (GSPE) have shown several bioactivities. They improve antioxidant cell defences\textsuperscript{(2)} and the plasma lipid profile\textsuperscript{(3,4)}, limit adipogenesis\textsuperscript{(5)} and function as insulinomimetic\textsuperscript{(6)} and anti-inflammatory\textsuperscript{(7)} agents. A preliminary study of an \textit{in vitro} model by the authors has also determined that dimeric and trimeric oligomers are the most powerful PC molecules that mimic the complete GSPE\textsuperscript{(8)}. Thus, to explain these health effects of PC and understand the mechanism by which PC act at the cellular level \textit{in vivo}, it is essential to determine PC stability during the digestive process as well as their bioavailability.

Despite it being clear that monomers are absorbed in human subjects and animals, there are controversies about the bioavailability of oligomeric forms. Different studies, following the ingestion of chocolate\textsuperscript{(9–12)}, black and green tea\textsuperscript{(13–16)}, red wine\textsuperscript{(17,18)} and grape seed extract\textsuperscript{(11)}, have shown that during digestion, the oligomers are fragmented into monomeric units of catechin and epicatechin. These are then absorbed, appearing in plasma and urine primarily as glucuronidated, methylated and sulphated metabolites. A study by Rios \textit{et al.}\textsuperscript{(19)} with six healthy subjects who consumed a rich PC cocoa beverage studied the depolymerisation of PC in the stomach and proved that they were remarkably stable in the stomach environment. Another study by Sano \textit{et al.}\textsuperscript{(20)} was the first work to detect procyanidin B1 in human serum after oral intake of a GSPE, the maximum concentration appearing 2 h after intake. As well as stability during digestion, the dose of intake could determine the primary site of phenol metabolism. Large doses are metabolised primarily in the liver, and small doses may be metabolised by the intestinal mucosa, with the liver playing a secondary role to further modify the polyphenol conjugates from the small intestine\textsuperscript{(21)}.

The bioavailability of polyphenols has been also reviewed, especially to focus on intestinal absorption and the influence of their chemical structure (for example glycosylation, esterification and polymerisation) and the effect of the food matrix composition\textsuperscript{(22)}. Different studies have been carried out into the interaction between polyphenols and a food matrix such as milk\textsuperscript{(23,24)}, olive oil\textsuperscript{(25)} or sugar\textsuperscript{(26)}. The evidence suggests that variations in polyphenol absorption also occur due to

**Abbreviations:** GSPE, grape seed procyanidin extract; IN, fraction found inside dialysis solution; OUT, fraction found outside dialysis solution; PC, procyanidins.

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interactions between polyphenols and other food components. For example, some authors suggest that the carbohydrate composition of the meal may influence flavonoid absorption through an effect on the motility of gastric secretions and hepatic blood flow. The polyphenol–protein interaction has also been studied, and shown to be similar to antigen–antibody interactions. The results of these studies suggest that interactions may occur between digestive proteins and monomers and dimers of flavanols from grape seed extract during pancreatic digestion.

Nevertheless, one of the most important limitations of the bioavailability studies in human subjects or animal models is related to inter-individual variability. In vitro digestion models are a very useful tool for studying the stability of food components and evaluating the potential effects of the food matrix on their bioavailability. Reviewing the literature, a wide range of in vitro digestion methods have been designed to simulate the effect of digestion on food or food components and study its consequences. In vitro digestion methods can be classified into two categories: static and dynamic gastrointestinal models. The static models simulate the transit of alimentary bolus through the human digestive tract by sequential exposure of food to simulated mouth, gastric and small intestinal conditions. In contrast, the dynamic gastrointestinal models mimic the gradual transit of ingested compounds through the digestive tract.

Other important limitation related to the study of PC bioavailability concerns the low concentration of their metabolites in biological samples. Therefore, measuring them requires a precise, robust, sensitive and selective analytical technique that allows them to be identified and quantified. Liquid chromatography with MS is a powerful tool for both confirmatory and quantitative analyses in complex matrices due to its high sensitivity and selectivity. Apart from the analytical technique, the use of an adequate extraction procedure to minimise the matrix effects is very important.

As PC dimers and trimers could be important in determining the biological effects of PC-rich food, understanding the bioavailability and metabolism of PC oligomers is fundamental. The objective of the present research is to study the stability of PC under digestion conditions, the metabolism and the bioavailability by using a combination of in vitro and in vivo models. Simultaneously, the matrix effect of a carbohydrate-rich food on digestibility and bioavailability of PC is investigated.

**Experimental methods**

**Chemicals and reagents**

A GSPE was used as the source of PC. The PC content of the extract was > 75% (monomers 22%, dimers 20% and trimers to pentamers 56%). The composition of commercial cereal-based food, used as carbohydrate-rich food, was as follows: starch (hydrolysed wheat, wheat, barley, maize and oats); malt extract; mineral salts (Ca, Fe and Zn); vitamins (C, niacin, E, thiamine, A, B6, B9 and D); flavouring (vanillin).

Internal standard catechol and the standards of (−)-epicatechin and (+)-catechin were purchased from Sigma Aldrich (St Louis, MO, USA); procyanidin B2 from Fluka Co. (Buchs, Switzerland) and trimer was isolated from phenolic extract obtained from cocoa nibs (32) using the technique described by Serra et al. (33). The cocoa nibs were kindly donated by La Morella Nuts, S.A. (Reus, Spain). Individual stock standard solutions and the internal standard were dissolved in methanol and stored inside a dark-glass flask at −18°C. Standard mixtures, at different concentrations, were prepared by appropriate dilution of the stock solutions, and were stored in dark-glass flasks at 4°C before chromatographic analysis.

Acetonitrile (HPLC grade), methanol (HPLC grade), acetone (HPLC grade) and glacial acetic acid (≥ 99.8%) were of analytical grade (Scharlab, Barcelona, Spain). Ultrapure water was obtained from a Milli-Q water purification system (Millipore Corporation, Bedford, MA, USA).

**Simulated gastrointestinal digestion**

A gastrointestinal in vitro digestion model was performed based on the methodology described by Ortega et al. (34). The discontinuous model describes a three-step procedure to mimic the digestive process in the mouth, stomach (gastric digestion) and small intestine (duodenal digestion) (Fig. 1). In order to evaluate the food matrix effect, two different samples were subjected to the digestion model: 300 mg PC extract (GSPE); 300 mg GSPE + 600 mg carbohydrate-rich food. After the gastric step, the digestion mixture was placed in tubes and centrifuged for 15 min at 12000g at 4°C, yielding the chyme (supernatant or soluble fraction) and the pellet (cloud or non-soluble fraction). Both fractions, the chyme and pellet, were freeze-dried (BMG Labtech,
At the end of the dialysed duodenal digestion step, two fractions were collected and analysed separately (34), the outside dialysis solution (OUT) (Fig. 1), which was considered the dialysable fraction; and the inside dialysis tub content (IN), referring to the non-dialysable fraction. The dialysable fraction (OUT) is considered the fraction that could be available for absorption into the systemic circulation by passive diffusion; while the non-dialysable fraction (IN) is attributed to the digested fraction that would reach the colon fermentation intact. Both fractions were freeze-dried and stored for chromatographic analysis of the PC content.

Chromatographic analysis of procyanidins in digested fractions

The digested freeze-dried fractions (chyme, pellet, IN and OUT) were dissolved in acetone–Milli-Q water–acetic acid (70:29:5·0·5, v/v/v) at a ratio of 1:5. The tubes were vortexed for 5 min and centrifuged at 12 500 rpm. The supernatant containing the solubilised phenols was filtered through 0·22 μm nylon filters before chromatographic analysis by ultradeionisation liquid chromatography–MS/MS using a Waters Acquity ultraperformance liquid chromatography system (Waters, Milford, MA, USA), equipped with a binary pump system (Waters) (32). The ultraperformance liquid chromatography analyses were performed using an Acquity high strength silica T3 column (100 mm × 2·1 mm inner diameter, 1·8 μm particle size; Waters) with a binary mobile phase. Solvent A was water–acetic acid (0·2 %), and the solvent as acetonitrile. Catechin and epicatechin were quantified using the curves of the respective standards. Epigallocatechin, epicatechin gallate and epigallocatechin gallate were quantified using the epicatechin curve. The PC, dimers and trimers were quantified with the calibration curves of the respective standards. The oligomers from tetramer to heptamer were quantified using the calibration curves of the respective standards. The MS spectra gave information about their structure. The structural information was also verified by using product ion scan and neutral loss scan in the MS/MS mode.

All the samples were analysed in triplicate by ultraperformance liquid chromatography–MS/MS according to the method in Serra et al. (33). The quantification of the free forms of the PC and their metabolites in the plasma samples was done using the calibration curves of the respective free forms, and the concentrations were expressed as catechin or epicatechin equivalents.

Statistical analysis

All the data are expressed as the mean of three replicates. In order to simplify the results shown in the tables, we omitted the standard deviation because all these values were lower than 10 %. The data were analysed by a one-way ANOVA test to assess the effect of the carbohydrate-rich food on the in vitro digestibility and in vivo pharmacokinetics of PC from the GSPE extract. A significant difference is considered at a level of P<0·05. All the statistical analysis was carried out using STATGRAPHICS Plus 5·1.

Results

Evaluation of the stability of procyanidins in digestion by the in vitro simulated digestion model

The first part of the study was focused on the evaluation of the stability of PC during digestion using an in vitro model. The composition of the GSPE used as a PC and proanthocyanadin source in the study is summarised in Table 1. Dimers and trimers were the most abundant compounds with 250·18 and 1568·72 μmol/g extract, respectively. The digestion procedure was applied to GSPE dissolved in water, and the same extract was mixed with cereal-based food in order to evaluate the influence of the food matrix with a high-carbohydrate content on the digestion stability of the PC.

The compounds determined initially in GSPE (Table 1) were analysed in all fractions of in vitro digestions to evaluate their digestibility and stability. After the gastric step, the content of PC and proanthocyanidins was measured in the soluble and non-soluble fractions, respectively, called the chyme and pellet (Fig. 1), and the results were expressed as μmol of concentration in the chyme fraction and as μmol per g of

Pharmacokinetic study

Rat plasma samples were obtained from 3-month-old male Wistar rats weighing between 370 and 420 g and were purchased from Charles River Laboratories (Barcelona, Spain). The Animal Ethics Committee of the Rovira i Virgili University approved all the procedures. The rats were housed in temperature-controlled rooms (22°C) and subjected to a standard 12 h light–12 h dark cycle (with a light period of 09.00–21.00 hours). All the animals were fed a standard diet of PanLab A04 (Panlab, Barcelona, Spain) and water.

After the animals had been in fasting conditions for between 16.00 and 17.00 hours with only access to tap water, two experiments were carried out: a dose of the PC-rich extract, 1 g GSPE per kg of body weight dissolved in water; a dose of 2 g carbohydrate-rich food + 1 g GSPE for kg of body weight dissolved in water, which was administered to the rats by intragastric gavage. Three animals per experiment were terminally anaesthetised with ketamine–xylazine and euthanised by exsanguinations at 0 h (fasting conditions), 1, 2 and 4 h (postprandial conditions). Blood samples were collected from the abdominal aorta with heparin-moistened syringes. The plasma samples were obtained by centrifugation (2000 g, 30 min at 4°C) and stored at −80°C until the chromatographic analysis of the PC. The identification of the PC metabolites was performed by analyses in MS (full-scan mode) and MS/MS (based on neutral loss scan and product ion scan mode) (33). First, analyses were carried out in the full-scan mode (from 419 to 1200 m/z) by applying different cone voltages from 20 to 60 V. When low cone voltages were applied, specific fragment ions were generated and the MS spectrum gave information about their structure. The structural information was also verified by using product ion scan and neutral loss scan in the MS/MS mode.

All the samples were analysed in triplicate by ultraperformance liquid chromatography–MS/MS according to the method in Serra et al. (33). The quantification of the free forms of the PC and their metabolites in the plasma samples was done using the calibration curves of the respective free forms, and the concentrations were expressed as catechin or epicatechin equivalents.
pellet (Table 2). After the gastric step, the amount of PC in the chyme was higher than in the pellet fraction independently of the presence of the carbohydrate-rich food in the digestion mixture. The PC oligomers, dimers to heptamers, showed higher concentrations than the monomers (catechin and epicatechin) in the chime and pellet fractions, the trimer form being the most abundant compound in the gastric fractions, reaching a concentration of 50 279·77 μmol in the chyme fraction after the gastric step. Nevertheless, forms with a high degree of polymerisation (pentamer, hexamer and heptamer) were not detected in pellet fraction.

After the duodenal step, an important loss of PC was observed, and most were retained in the non-dialysable fraction (IN). Nevertheless, dimers and trimers were quantified in the dialysable (OUT) fraction (Fig. 1), as well as the monomeric structures. As a consequence of the duodenal digestion conditions, decomposition of oligomeric forms was observed in the respective monomeric forms. Additionally to the depolymerisation of GSPE, a decrease in the ratio catechin/epicatechin was observed in the digestion mixture. Initially, in the GSPE extract, this ratio was approximately 1·1 and after digestion, a change in the ratio of roughly 0·7 was observed in all fractions.

Finally, the comparative analysis between the concentrations of the different PC in the digestion mixtures revealed that the presence of carbohydrate-rich food, as a source of carbohydrates, showed a significant effect on the digestibility of PC in the in vitro digestion model. When PC and procyanidins were submitted to digestion together with carbohydrate-rich food, the amount of catechin and epicatechin present in the digestion mixture after the gastric step was significantly higher (P<0·05) (Table 2). By contrast, the concentration of the other compounds was greater when GSPE was digested without carbohydrate-rich food. Greater differences were observed in the non-soluble fraction (pellet) (Fig. 1), showing a higher concentration of PC (P<0·05) when the extract was digested with carbohydrate-rich food. By contrast, the effect of carbohydrates on the digestibility during duodenal digestion was lower. Thus, the concentration of PC in the dialysed fraction (OUT) was similar independently of the presence of carbohydrates, with the exception of the epigallocatechin gallate that showed a lower concentration (P<0·05) in the presence of carbohydrates. On the other hand, significant differences were observed in the non-dialysable fraction (IN), except the forms with a higher degree of polymerisation (pentamers to heptamers), which showed the same concentration under the two digestion conditions. However, no clear tendency was observed in the results of non-dialysable (IN) fraction in relation to the presence or absence of carbohydrates.

Pharmacokinetic of procyanidins

Figs. 2 and 3 show the results of the pharmacokinetic study of rat plasma after the ingestion of GSPE without and with carbohydrate-rich food. Under basal conditions (time = 0), PC were not detected in rat plasma. Two hours after the GSPE ingestion, several metabolites, such as catechin glucuronide and epicatechin glucuronide, reached peak plasma concentrations with 6·24 and 9·74 nM, respectively, when GSPE was ingested with carbohydrate-rich food, and 6·32 and 8·71 nM when GSPE is ingested without carbohydrate-rich food. In contrast, the concentration of PC, free forms of

Table 1. Phenolic composition of grape seed procyanidin extract (GSPE)*

<table>
<thead>
<tr>
<th>Compound</th>
<th>Concentration (μmol/g)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Catechin</td>
<td>58</td>
</tr>
<tr>
<td>Epicatechin</td>
<td>52</td>
</tr>
<tr>
<td>Epigallocatechin</td>
<td>5·50</td>
</tr>
<tr>
<td>Epicatechin gallate</td>
<td>89</td>
</tr>
<tr>
<td>Epigallocatechin gallate</td>
<td>1·40</td>
</tr>
<tr>
<td>Dimer</td>
<td>250</td>
</tr>
<tr>
<td>Trimer</td>
<td>1568</td>
</tr>
<tr>
<td>Tetramer</td>
<td>8·8</td>
</tr>
<tr>
<td>Pentamer</td>
<td>0·73</td>
</tr>
<tr>
<td>Hexamer</td>
<td>0·38</td>
</tr>
<tr>
<td>Heptamer</td>
<td>0</td>
</tr>
<tr>
<td>Epigallocatechin gallate</td>
<td>1·40</td>
</tr>
<tr>
<td>Epicatechin gallate</td>
<td>89</td>
</tr>
</tbody>
</table>

*The results are expressed as μmol/g of GSPE.

Table 2. Procyanidin and proanthocyanidin contents in the different fractions of the gastric and duodenal steps of the simulated gastrointestinal digestion of grape seed procyanidin extract (GSPE) without and with carbohydrate-rich food

<table>
<thead>
<tr>
<th>Compound</th>
<th>Chyme (μM)</th>
<th>Pellet (μmol/g)</th>
<th>OUT (μM)</th>
<th>IN (μM)</th>
<th>Chyme (μM)</th>
<th>Pellet (μmol/g)</th>
<th>OUT (μM)</th>
<th>IN (μM)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Catechin</td>
<td>1755a</td>
<td>7·4a</td>
<td>95a</td>
<td>451a</td>
<td>1928b</td>
<td>37b</td>
<td>83a</td>
<td>608b</td>
</tr>
<tr>
<td>Epicatechin</td>
<td>2221a</td>
<td>9·6a</td>
<td>126a</td>
<td>613a</td>
<td>2499b</td>
<td>42b</td>
<td>114a</td>
<td>823b</td>
</tr>
<tr>
<td>Epigallocatechin</td>
<td>258a</td>
<td>0·49a</td>
<td>2·6a</td>
<td>37a</td>
<td>186b</td>
<td>1·41b</td>
<td>2·4a</td>
<td>34·5b</td>
</tr>
<tr>
<td>Epicatechin gallate</td>
<td>6305a</td>
<td>29·8a</td>
<td>109a</td>
<td>1772a</td>
<td>5272b</td>
<td>191b</td>
<td>72b</td>
<td>2115b</td>
</tr>
<tr>
<td>Epigallocatechin gallate</td>
<td>136a</td>
<td>0·409a</td>
<td>0·80a</td>
<td>287a</td>
<td>91b</td>
<td>1·9b</td>
<td>0·65a</td>
<td>25·2b</td>
</tr>
<tr>
<td>Dimer</td>
<td>12 953a</td>
<td>37·1a</td>
<td>197a</td>
<td>9517a</td>
<td>11 485b</td>
<td>124b</td>
<td>168a</td>
<td>4005b</td>
</tr>
<tr>
<td>Trimer</td>
<td>50 279a</td>
<td>109a</td>
<td>147a</td>
<td>8101a</td>
<td>37 505b</td>
<td>356b</td>
<td>111a</td>
<td>8659a</td>
</tr>
<tr>
<td>Tetramer</td>
<td>10 076a</td>
<td>21·0a</td>
<td>0·25a</td>
<td>1748a</td>
<td>5388b</td>
<td>82b</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Pentamer</td>
<td>16·8a</td>
<td>ND</td>
<td>ND</td>
<td>12·7a</td>
<td>ND</td>
<td>ND</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Hexamer</td>
<td>185a</td>
<td>ND</td>
<td>ND</td>
<td>16·9a</td>
<td>49b</td>
<td>ND</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Heptamer</td>
<td>0·66a</td>
<td>ND</td>
<td>0·73a</td>
<td>ND</td>
<td>2·34b</td>
<td>ND</td>
<td>ND</td>
<td>ND</td>
</tr>
</tbody>
</table>

OUT, fraction found outside dialysis solution; IN, fraction found inside dialysis solution; ND, not detected.

*Mean values within a column with unlike superscript letters were significantly different (P<0·05) between GSPE + carbohydrate-rich food and GSPE in each digestion step (unpaired Student’s t test).
Fig. 2. Pharmacokinetic curves of monomeric forms, (–B–) catechin, (–K–) epicatechin (–O–) and epicatechin gallate (a and b), and polymeric forms, (–X–) dimer and (–W–) trimer (c and d), detected in rat plasma collected between 0 and 4 h after the ingestion of grape seed procyanidin extract with and without carbohydrate-rich food. The results are expressed as nM. * Mean value was significantly different from that without carbohydrate-rich food (P<0.05; unpaired Student’s t test).

Fig. 3. Pharmacokinetic curves of grape seed procyanidin extract (GSPE) metabolites according to the monomer structure, being (–S–) catechin methyl glucuronide, (–þ–) catechin glucuronide and (–) catechin methyl sulphate (a and b), and (–A–) epicatechin methyl glucuronide, (–V–) epicatechin glucuronide and (–) epicatechin methyl sulphate (c and d), detected in rat plasma collected between 0 and 4 h after the ingestion of GSPE with and without carbohydrate-rich food. The results are expressed as nM. * Mean value was significantly different from that without carbohydrate-rich food (P<0.05; unpaired Student’s t test).
dimers and trimers in the plasma reached a maximum 1 h after administration. These free forms appeared in the plasma faster than the glucuronidated forms of the monomers. Dimer concentration was greater after the ingestion of GSPE without the presence of carbohydrate-rich food, reaching a postprandial concentration in plasma of 0·57 nM as against 0·12 nM when the GSPE was ingested with carbohydrate-rich food. Similarly, the presence of the carbohydrate-rich food reduced the maximum concentration of the trimer in the plasma. Non-conjugated forms of monomers (catechin, epicatechin and epicatechin gallate) were present in the plasma at very low concentration, less than 0·03 nM under the two experimental conditions. Glucuronidated forms of catechin and epicatechin were the most abundant, and the second most abundant metabolite was methyl-glucuronidated forms, independently of the presence of the carbohydrate-rich meal.

Comparison between in vitro digestion method and in vivo bioavailability

Fig. 4(a) and (b) show the percentage of different PC present in the dialysed fraction (OUT) of the duodenal digestion mixture corresponding to the in vitro digestion model (Fig. 1); and Fig. 4(c) and (d) show the percentage of free PC and their metabolites in the rat plasma after the ingestion of the GSPE extract (in vivo model). The expression of the quantification of PC as percentages in both the in vitro and in vivo models aims to normalise the results to facilitate their comparative analysis independently of whether the concentrations of the compounds varied widely in both models. In all the plasma samples, epicatechin was the most abundant compound, glucuronidated forms being 88 and 89 % of the total, when GSPE was digested alone and with carbohydrate-rich food, respectively. This was followed by catechin, which was the second most abundant compound. Nevertheless, the dimer structure was the most substantial PC quantified in the dialysed fraction (OUT) after the duodenal step of the in vitro digestion model, reaching a percentage of almost 30 % (Fig. 4(a) and (b)); however, this compound was a very low percentage in the plasma samples (1·69 % when the GSPE was administered alone and 0·35 % when it was administered with carbohydrate-rich food) (Fig. 4(c) and (d)). The quantification of the trimer in the plasma samples showed percentages under 1 %, independently of the presence or absence of carbohydrate-rich food in the intake, whereas its percentage in the dialysed fraction (OUT) of the duodenal digestion was higher, close to 20 %. Epicatechin gallate behaved similarly.

Discussion

We have shown, using in vitro cell lines, that dimer and trimer PC are the most powerful molecules that mimic the in vivo

![Image](https://doi.org/10.1017/S0007114509992741)
effects of GSPE (2). As there is some controversy about the bioavailability of dimeric and trimeric PC, we used a combination of in vitro and in vivo models in order to measure the bioavailability of dimer and trimer PC. The effects of the digestion process on their stability and the food matrix effect on their potential bioaccessibility were also evaluated.

The GSPE used as a source of PC in the present study contained a substantial amount of high molecular weight polymeric PC, dimer and trimer being the most abundant. After the simulated in vitro digestion, these PC oligomers were present in high concentration in all digestion fractions, showing an important stability under gastric and duodenal digestion conditions. The most important aspect is related to the increase in their concentration observed in the digestion mixtures after the two steps. This is probably related to the partial hydrolysis of the highly polymerised molecules or condensed tannins in the GSPE. However, the pharmacokinetic study revealed a limited absorption. The limitation in the diffusion across the epithelial cells could explain the nanomolar levels quantified in the postprandial rat plasma after the intake, at micromolar levels, of the oligomeric PC. The pharmacokinetic results shown in the present study agree with previous research that has shown the presence of some dimers (B2 and B5), detected at nanomolar levels, in the plasma of rats given cocoa extracts (35, 36). By contrast, in other studies, the dimer (B3) and trimer (C2) were not detected in the plasma of rats given purified compounds (37). With regard to the metabolism, the major metabolites observed in rat plasma in the present study were glucuronidated forms of catechin and epicatechin, and mainly the dimer and trimer oligomers. The pharmacokinetic study of the rat plasma revealed that the glucuronidated forms were the most abundant metabolite. These results agree with those in Vaidyanathan & Walle (43) who observed that rat liver microsomes efficiently glucuronidate epicatechin. Glucuronidation of catechin has been described to be more active in rat livers than in the intestine, in both in vitro (44) and in vivo models (45). Traces of glucuronidated and sulphated metabolites from monomers have also been detected in the lumen of gut from rats treated with GSPE (1). One of the most important factors determining the metabolic fate of polyphenols in human subjects is the quantity ingested (46). According to Hacket et al. (47), when food polyphenols are administered at a pharmacological dose, they are found in free forms in the blood. By contrast, when the dose is decreased to only a few milligrams, the monomers were conjugated and no free forms were detected (18). Related to this, in the present study, the free forms of catechin and epicatechin were found at very low concentrations in the plasma samples after the ingestion of GSPE. The present results may be coherent with the dose administered to the rats, because the dose used is an intermediate position between the pharmacological dose and typical dose of polyphenols present in rich foods.

The last objective in the present study was to evaluate the effect of carbohydrate-rich foods on the bioavailability of the GSPE. Relatively, little is known about the effect of food on the bioavailability of PC. Food within the gastrointestinal tract is known to markedly alter the oral absorption of many xenobiotic compounds (45). In the present in vitro study, when the extract was ingested with carbohydrate-rich food, an enhanced uptake of the monomeric PC was observed. This result is in agreement with the results in Schramm et al. (20), who reviewed that the effect observed with carbohydrate-rich foods (bread, sucrose and grapefruit juice) may
have been mediated by a carbohydrate-specific effect on gastrointestinal physiology (e.g. motility and/or secretion) or a carbohydrate-specific enhancement of the activity of a yet unidentified carbohydrate–flavanol transporter.

In conclusion, dimer and trimer PC are absorbable in vivo, reaching maximum concentrations in the plasma as soon as 1 h after PC ingestion. Their low concentrations in the plasma are not the consequence of instability during the digestion process as the in vitro digestion model shows high levels of dimer and trimer after duodenal digestion. Moreover, absorption of dimer and trimer PC is repressed by the simultaneous presence of carbohydrate-rich food.

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