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Circular valorization of coffee silverskin through supercritical CO₂ for the production of functional extracts†

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The development of sustainable procedures for the valorization of industrial biomass waste represents a major challenge for the scientific community. Among the residual biomasses, coffee silverskin could represent an attractive source of bioactive compounds. In spite of its promising applications, silverskin is still underutilized and nowadays is still discarded by roasters as solid waste in landfill. This work presents the application of a green unconventional extraction strategy, supercritical CO₂, with a particular focus on the extractability of polar compounds from green coffee beans and their related wasted biomass, coffee silverskin, by means of the addition of ethanol as a co-solvent, allowing us to shift the window of extractables towards more polar molecules. The results point out the ability of the supercritical fluid to efficiently target chlorogenic acids, minimizing the excessive use of organic solvents and avoiding the presence of inorganic or organic acids. The use of a co-solvent, at the expense of reducing the selectivity towards target species, was demonstrated to efficiently co-extract other classes of compounds directly connected to the relevance of silverskin's phytochemical profile, dealing mainly with its antioxidant and prebiotic properties, pointing at interesting applications in functional food products.

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Sustainability spotlight

In the context of circular economy, the agri-food chain represents the sector in which there is the highest production of wasted biomasses with a direct strong potential to be further investigated. This, joint with the growing interest in the nutraceutical industry that is continuously raising, pushes research to seek techniques to extract bioactive compounds starting from these wastes. Our research merges the valorization of a biomass residual, coffee silverskin, responsible for the production of 2400 tons of waste per year only in Italy, with the development of a green and sustainable technique, supercritical CO₂, to target its polar fraction to be exploited in the nutraceutical field. This approach is perfectly aligned with the following UN sustainable development goals: responsible consumption and production (SDG 12), industry, innovation, and infrastructure (SDG 9) and climate action (SDG 13).

Introduction

Biomass valorization constitutes a key aspect of the circular economy and an opportunity to unlock the full potential of the bioeconomy and enhance the possible synergies between different sectors. Within circular models, agriculture and in particular the agri-food chain represent the sector in which there is the highest production of waste biomass with a direct strong potential to be further investigated.¹ This, together with a growing interest in the nutraceutical industry, has led to research to find new bioactive molecules and techniques to

extract and isolate them from these wastes. Among the food-chain-related wastes, plant residuals could represent an attractive source of more sustainable bioactive compounds for the nutraceutical and/or functional foods sector.²

The exploitation of byproducts towards the generation of added-value compounds should meet and merge the need to develop alternative and innovative extraction techniques that are able to guarantee the sustainability of the entire process, from biomass selection to the technological methods to final added-value generation. In this context, unconventional extraction techniques are promising approaches able to enhance compounds recovery in a greener way compared to conventional solvent-based technologies. Among them, subcritical water and supercritical carbon dioxide extraction are attracting more and more interest. These extraction techniques are efficient, economical, and promising routes for the recovery of bioactives from different matrices. When the target compounds could undergo thermal degradation, supercritical

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CO₂ (sc-CO₂) might be a preferable choice, in order to maintain all the antioxidant profiles in the extracts.^{3,4} The sc-CO₂ technology, employing CO₂ as extraction fluid is indeed characterized by low environmental impacts, the use of organic solvents can be avoided or minimized, ensuring safer and selective processes directly on the biomass without any pretreatment, with the possibility to recycle the employed CO₂ in industrial plants.⁵ Following these global trends, where the interest in “green” products and technologies is growing, this work presents as a core strategy a supercritical CO₂ extraction methodology with a particular focus on the extractability of polar compounds. Since sc-CO₂ alone displays a solvating power similar to *n*-hexane, most of the literature is focused on the primary and direct use of the supercritical fluid to target the lipidic fraction and, more in general, non-polar compounds.⁶ On the other hand, the addition of a co-solvent in a relatively low ratio enables the modification of the supercritical fluid polarity, allowing the shifting of the window of extractables towards more polar molecules.⁷ This approach has already been discussed in the literature; however, it is still underdeveloped, probably due to the loss of certain selectivity attributable to the presence of a co-solvent.^{8,9} Nevertheless, if the loss of selectivity is evaluated together with an investigation of the co-extracted compounds and their potential added value, a sustainable platform might be designed, in a more complete biorefinery concept, where the effect of co-extraction has an interesting economical outcome from the process perspective. Other molecules, apart from the targeted ones, could be beneficial towards specific applications in particular in the nutraceutical field, where they could have great potential as prebiotics or as building blocks for functional materials.^{10–12}

In this wide context, this work presents for the first time the valorization of the polar fraction from coffee silverskin by means of supercritical CO₂ in the presence of a co-solvent. Silverskin, the thin layer directly in contact with the coffee bean removed during the roasting process, representing about 4.2% (w/w) of coffee beans and accounting for a production of 2400 tons of waste produced per year only in Italy, was selected as an interesting case study together with its related original biomass, green coffee beans, as the term of comparison.¹³ In spite of its promising applications, silverskin is still underutilized and nowadays is still discarded by the roasters as solid waste in landfill. To date, the exploitation of silverskin has been focused mainly on its lipidic content, explored also by means of environmentally-friendly techniques such as sc-CO₂, but resulting in the loss of its high-value polar content^{14,15}. This latter connects with the relevance of silverskin's phytochemical profile, dealing mainly with its antioxidant and prebiotic properties, mostly related to the presence of phenolics and, in particular, chlorogenic acids.^{16–18} To the best of our knowledge, no investigations are present in the literature related to the extraction of phenolics from coffee silverskin by means of a green extraction strategy such as supercritical CO₂. In addition, the aim of the present work was to study the effect of co-solvent addition to the supercritical fluid and investigate the presence and nature of the co-extracted compounds.

Materials and methods

Reagents and starting biomasses

All the chemicals were used without further treatment. Acetonitrile and water, when used for liquid chromatography, were purchased from Sigma-Aldrich (Italy) as ultra-performance liquid chromatography grade. Formic acid, trifluoroacetic acid, ethanol, *n*-hexane, caffeine (powder, ReagentPlus®), and 5-cafeoylquinic acid (≥95%) standards were purchased from Sigma-Aldrich (Italy). A carbon dioxide tube (CO₂ purity 4.5) was purchased from Sapio (Italy). Chemicals, *p*-coumaric, caffeic, and ferulic acid standards were purchased from Sigma-Aldrich (SE).

Green coffee beans (GCB) of Robusta species from Vietnam and micronized coffee silverskin (CS) were kindly donated by an Italian roasting company (Ideal Caffè snc, Verderio) and then pulverized using a knife blender at its maximum velocity for 3 min.

Delipidization of GCB and CS

The extraction yield of lipids was calculated as:

$$\frac{\text{Mass}_{\text{extract}}}{\text{Mass}_{\text{starting biomass}}} \times 100.$$

Three replicates were performed for each experiment.

Supercritical CO₂ extraction. Supercritical CO₂ extractions (sc-CO₂) were performed using a pilot unit SFT110XW system (Supercritical Fluid Technologies, Inc., USA). It consists of an SFT-Nex10 SCF CO₂ pump (constant pressure-piston pump) with a Peltier cooler, a 100 cm³ stainless steel extractor vessel inserted in an oven, and a collection vial. 30.15 g of CS or 47.30 g of GCB were loaded in the vessel during their respective delipidization experiments.

Following the results from our previous work on coffee silverskin,¹⁹ the operative pressure was set at 300 bar in all experiments, while the temperature of the vessel and the temperature of the restrictor block were maintained at 60 °C and 75 °C, respectively, throughout the extraction period. For each extraction experiment, many cycles, each one comprising 30 min of maceration time in static conditions and 10 min of dynamic conditions, were performed until no evident extracted mass was further gained. Under dynamic conditions, the valves were opened and the extract was collected in a vial, keeping a CO₂ gas constant flow rate of 8 SCFH (standard cubic feet per hour). In the case of the lab-scale supercritical fluid extractor, the duration of static and dynamic cycles (and hence the final number of cycles) was optimized on the basis of the need for a specific soaking time under static conditions before opening the valves to run the dynamic process to collect the extract. The extracts were then stored at –20 °C for subsequent analysis, whilst the defatted biomasses were collected from the vessel and subjected to further experiments.

Solvent-based extraction. In a reference conventional method, *n*-hexane was employed as the extraction solvent. 10 g of the starting biomasses (GCB or CS) were suspended in



100 mL of *n*-hexane and: (i) magnetically stirred for 6 h at room temperature, or (ii) extracted by means of Soxhlet apparatus. The residual solid was dried under static air at 40 °C. The solvent was evaporated from the filtered solution using a rotary evaporator (Buchi) and then under a vacuum pump.

Chlorogenic acid extraction

Supercritical CO₂ extraction + co-solvent. The same equipment employed for biomass supercritical delipidization was then connected to a co-solvent pump (515 HPLC pump, Waters). 29.5 g of defatted CS or 46.6 g of defatted GCB were loaded during their respective extraction experiments. Before setting the supercritical conditions inside the vessel, 10% v/w (with respect to the loaded biomass) of water : ethanol (1 : 1) mixture was injected inside the vessel, as a result of preliminary trials aimed to assess the best co-solvent mixture. The operative supercritical pressure was set at 300 bar, while the temperature of the vessel and the temperature of the restrictor block were maintained at 60 °C and 75 °C, respectively, throughout the extraction period. A static maceration 30 min period was maintained at the beginning, followed by a dynamic period of 1 hour maintaining an ethanol flow rate of 0.5 mL min⁻¹ (corresponding to 0.25% v/v of the CO₂ amount) with a CO₂ gas flow rate constant at 8 SCFH (standard cubic feet per hour). After this first cycle, many cycles, comprising 15 min of maceration time in static conditions and 1 hour of dynamic conditions with co-solvent addition, were performed. The extracted solutions were collected and ethanol was evaporated using a rotary evaporator (Buchi) and then under a vacuum pump.

Solvent-based extraction. A conventional solvent-based extraction method was employed as a reference in order to quantify the total chlorogenic acids and caffeine content of the biomasses and identify the co-extracted species. Some reference literature works were considered to select the most appropriate and effective strategy.^{20–22} In detail, 1 gram of biomass powder of CS or GCB was mixed with 10 mL of an acidic hydroalcoholic solution of water : ethanol = 7 : 3 (+1% formic acid). The suspension was magnetically stirred at room temperature for 4 hours, centrifuged at 6000 rpm for 5 min, and then filtered on a Buchner funnel. The solvent was evaporated using a rotary evaporator (Buchi) and then under a vacuum pump.

Characterization of the extracts

Chlorogenic acids and caffeine profiles. The composition of the extracts in terms of chlorogenic acids and caffeine was analyzed using a Waters ACQUITY UPLC (Ultra-high Performance Liquid Chromatography) system (Waters corp., United States) equipped with a quaternary solvent manager system, autosampler, thermostated column compartment and dual-wavelength UV-Vis detector. Water solutions of the samples were filtered (0.2 μm nylon filters) before injection (5 μL) in the system. The analytical separation was performed using an ACQUITY UPLC[®] BEH C18 column (1.7 μm, 2.1 × 50 mm). The mobile phase was composed of water containing 0.1% of formic acid (v/v) (A) and acetonitrile containing 0.1% of formic acid (v/v) (B). The flow rate was set at 0.25 mL min⁻¹ and the linear

gradient elution was: 0 min, 95% A; 3 min, 93% A; 7 min, 90% A; 15 min, 80% A; 18 min, 95% with a re-equilibration time of 3 min before the next injection. The column temperature was maintained at 34 °C and the wavelengths were set at 275 nm and 324 nm, corresponding to the maximum absorption of caffeine and chlorogenic acids, respectively. Data were processed using Empower 3 workstations.

Before sample injections, six dilutions of 5-caffeoylquinic acid (5-CQA) aqueous solution (515 μg mL⁻¹) and six dilutions of caffeine aqueous solution (220 μg mL⁻¹) were prepared in the range 5.15–515 μg mL⁻¹ for 5-CQA and in the range 2.2–220 μg mL⁻¹ for caffeine. Standard solutions were filtered (0.2 μm polypropylene filters) and injected three times in the UPLC system. 5-CQA and caffeine were eluted as sharp peaks. In the operative concentration range, the trend was linear, with no saturation effects that could bend the linearity.

The area under each peak was quantified using the instrumental software and plotted *versus* the concentration. The best fit of experimental data in the plot “peak area at 324 nm *vs.* [standard concentration]” was then used for 5-CQA and caffeine quantification in each extract.

The best fit of experimental data in the plot of “A_{324nm} *vs.* [5-CQA]” was a straight-line represented by the following mathematical equation: $y = 6.1 \times 10^7 x - 3 \times 10^5$, resulting in an R^2 equal to 0.996. The slope m and intercept q of the regression line with their respective standard deviations were $(6.1 \times 10^7 \pm 2 \times 10^6)$ and $(-3 \times 10^5 \pm 4 \times 10^5)$, respectively. The calculated equation of the regression line was then employed to determine the 5-CQA concentration in each extract.

The best fit of the experimental data in the plot of “A_{324nm} *vs.* [caffeine]” was a straight-line, represented by the following mathematical equation: $y = 4.91 \times 10^7 x + 7 \times 10^4$, resulting in an R^2 equal to 0.999. The slope m and intercept q of the regression line with their respective standard deviations were $(4.91 \times 10^7 \pm 8 \times 10^5)$ and $(7 \times 10^4 \pm 9 \times 10^4)$, respectively. The calculated equation of the regression line was then employed to determine the caffeine concentration in each extract.

When necessary, chromatographic separation was followed by mass spectrometry (LCQ Fleet Thermofisher) analysis. A negative electrospray mode was used for the ionization of molecules with a capillary voltage of -42 V, at a capillary temperature of 275 °C. The heater temperature was set at 150 °C, the gas flow rate was 20 (arb) and the spray voltage was 5 kV. The monitored mass range was from m/z 50 to 1000. Before the sample injection, mass spectrometry parameters were optimized using a commercial standard of 5-caffeoylquinic acid.

Monosaccharides profile. The carbohydrates profile of the soluble extracts after the supercritical CO₂ extraction (samples obtained during the 6th cycles in sc-CO₂) and solvent extraction were analyzed by direct solubilization in water or hydrolysed by trifluoroacetic acid (TFA), followed by the analysis of the released monosaccharides by high-performance anion-exchange chromatography-pulsed amperometric detection analysis (HPAEC-PAD). In summary, samples were weighed at 1 mg in glass tubes (triplicate), and 1 mL of MiliQ water or 2 M TFA was added. In the case of TFA, the hydrolysis was performed at 120 °C for 3 h. Hydrolyzed samples were then dried under a stream of



air and then re-suspended in water. Both sets of samples were analyzed using the ICS6000 fitted with a Dionex CarboPac PA1 IC column (Thermo Scientific, Stockholm, Sweden), as previously described.²³ Analysis was performed in triplicate.

Hydroxycinnamic acid profiles. Phenolic acids were released by saponification followed by analysis using high-performance liquid chromatography (HPLC) as described in the literature.^{24,25} 10 mg of each sample (solvent extracts and samples obtained during the 6th cycles in sc-CO₂) were saponified with 500 μ L of 2 M NaOH at either 30 °C (for soluble extracts) or 80 °C (for insoluble ones) for 17 h, with stirring in duplicate. The saponified samples were then acidified to pH 3.0 using 12 M HCl, partitioned with ethyl acetate, and dried under nitrogen. The dried phenolic acid fractions were resuspended in methanol: 2% (v/v) acetic acid (1 : 1, v/v), filtered through Chromacol 0.2 μ m filters (Scantec Nordic, Sweden), and analyzed on the HPLC using the ZORBAX StableBond C18 column (Agilent Technologies, USA) fitted with photodiode array detection at 200–400 nm (Waters 2695, MA, USA). Samples that were not saponified were diluted with methanol 2% (v/v) acetic acid (1 : 1, v/v), filtered through Chromacol 0.2 μ m filters (Scantec Nordic, Sweden), and subjected to the HPLC for analysis. The phenolic acids were separated in accordance with a previous method using *p*-coumaric, caffeic, and ferulic acid as standards.²⁶

Conjugate profiles present in samples. The presence of single and conjugated phenolics of the extract (solvent extracts and samples obtained during the 6th cycles in sc-CO₂) was checked using liquid chromatography electrospray ionization tandem mass spectrometry (LC-ESI-MS), using a Synapt G2 mass spectrometer (Waters Corporation, Milford, MA, USA). Saponified and non-saponified samples were diluted in acetonitrile 50% (v/v) with 0.1% (v/v) formic acid to 0.1 mg mL⁻¹ and filtered through Chromacol 0.2 μ m filters (Scantec Nordic, Sweden). Samples were then briefly passed through an Eclipse Plus C18 column 1.8 μ m (2.1 \times 150 mm) (Agilent Technologies, Santa Clara, CA) for automation and analyzed using positive-ion mode in the ESI-MS. The capillary and cone voltages were set to 3 kV and 25 kV, respectively. The phenolic dimers and conjugates of carbohydrate-phenolics were detected as [M + H]⁺ or [M + Na]⁺ adducts.

Size exclusion chromatography. The molar mass distribution of the extracts (solvent extracts and samples obtained during the 6th cycles in sc-CO₂) was determined using the SECurity 1260 size exclusion chromatography (Polymer Standard Services, Mainz, Germany) coupled to a refractive index, as formerly described.²⁷ The samples were solubilized in water containing 100 mM NaNO₃ and 5 mM NaN₃.

Results and discussion

Biomasses milling

One of the most critical operating parameters affecting the extraction yield is particle size. The solid matrix must be blended to increase the mass transfer area. Enhancing the exposed surface area to the solvent makes the extractions more effective in terms of yield, facilitating the mass transfers. Solutes can be in fact located on the surface of the cells of the

vegetable matrix or in intracellular spaces. As the particle size of the sample decreases, the extracted solute would travel shorter diffusional paths along the solid substrate, and mass transfer barriers, such as cell walls, would be destroyed. However, it was also reported by Coats *et al.* that some adverse hydrodynamic effects may occur when excessively small particles are treated in a packed bed.^{28,29} Particle size is hence considered a critical parameter, influencing the yield of extraction processes.³⁰

Coffee silverskin (CS) was provided as already micronized in the form of fine powder. Optical microscopy assessed an average particle size of about 100 μ m. Green coffee beans (GCB) were subjected to blending before extraction. Due to the high content of lipids, the GCB fatty biomass could not be transformed into a fine powder as silverskin, not even by embrittlement by means of liquid nitrogen before blending. In this case, the minimum achievable particle size was in the range of 500–1000 μ m, as detected by an optical microscope. GCB particle size was in any case in the range of recommendations found in the literature.³⁰ Pictures of the pristine and blended biomasses are shown in the ESI in Fig. S1.†

Delipidization extraction yields and kinetics

The lipid content of CS was already reported in a recent publication.¹⁹ For the purpose of this work, a preliminary delipidization of both biomasses was performed in order to ease the successive extraction of the target polar compounds. The presence of excessive lipids would hinder the accessibility of the supercritical solvent (and the co-solvent) towards the generation of the necessary interface between the fluid and the solutes. The presence of lipids in the biomasses could be responsible for a certain hindrance in the targeted polar compounds extraction, rendering the biomass less prone to be penetrated by the extraction fluid. Moreover, lipids could contaminate the extracts later obtained in sc-CO₂ in the presence of the co-solvent.

The lipid content was determined by conventional solvent-based procedures by triplicate trials, corresponding to 5.0 \pm 0.5% and 1.4 \pm 0.1% (w/w on dry biomass) for GCB and CS samples, respectively.

When the extraction was performed in supercritical CO₂, the optimization of process parameters followed the results from the previous work by Nasti *et al.*, set at 60 °C and 300 bar. Cycle by cycle, with an alternation of static and dynamic periods, an incremental mass of extract was obtained from both GCB and CS, and the extraction process was maintained active until no evident mass gain was observed, thus corresponding to the maximum extractable yield. These values corresponded to 6.2 \pm 0.1% and 1.4 \pm 0.2% for GCB and CS, respectively. It was expectable that these yield values are comparable to the corresponding numbers obtained by conventional extraction. A slight difference was observed for GCB. The lipids yield obtained by means of sc-CO₂ was 20% higher than the conventional extract. This can be attributed to the impossibility of finely micronizing GCB, hence making the biomass-hexane solid-liquid interface contact less effective, due to a lower exposed surface area. The penetration inside the porosity of the



supercritical fluid, partially behaving like a gas in terms of permeability, was, on the other hand more effective, reaching pores inaccessible to hexane, thus enhancing the mass transfer and hence the yield.³¹

sc-CO₂ delipidization extraction kinetics are displayed in Fig. 1. Both extractions follow the model explained by da Silva *et al.*, primarily based on two mass transfer mechanisms: solubility and diffusion.³² The model assumes two distinct modes of extraction: initial constant rate extraction that is controlled by solubility, and falling rate extraction that is controlled by diffusivity. An initial period of extraction (the steepest, represented by the first four points in each dataset) corresponds to the depletion of lipids found on the particle surface, while the final period represents the extraction from inside the porosity, where the mass transfer is more hindered and the extraction rate slows down until it zeroes.^{32,33} For the purpose of this work, during the delipidization process, the extraction was interrupted after 15/16 cycles when a plateau was reached, meaning that no more lipids could be obtained in the process.

All the extracts obtained in the delipidization process were subjected to UPLC analysis, and none of them displayed evidence of chlorogenic acid presence. The details on CS lipidic extract composition have been reported in a recent paper by Nasti *et al.*¹⁹

Chlorogenic acid extraction

Chlorogenic acids, due to their well-known antioxidant and interesting biological activities, were considered the main target compounds adopting supercritical fluid in the presence of a co-solvent.^{20,34} For comparison purposes, a solvent-based extraction was carried out employing a literature method with the aim of effectively extracting all the chlorogenic acid content from the delipidized biomasses. In the conventional extraction, a 1 : 10 solid-to-liquid ratio was employed since it represents a good compromise between not exceeding the amount of solvent and guaranteeing an exhaustive extraction of targeted compounds.

The first three sc-CO₂ extracts obtained in the presence of co-solvent from GCB delipidized biomass and the first five extracts obtained in the presence of co-solvent from delipidized CS biomass were all similar to samples obtained during

delipidization and almost completely insoluble in water. After the mentioned cycles, successive extracts displayed an almost complete water-solubility.

Fig. 2 displays the chromatograms recorded at 324 nm (maximum absorbance for chlorogenic acids) for the solvent-based extracts and a selection of representative samples from sc-CO₂ extraction cycles.

The UPLC data show an almost complete overlap of the chromatograms at 324 nm, corresponding to the maximum absorption of chlorogenic acids, relative to all four samples, irrespective of their extraction method. This leads to the conclusion that sc-CO₂ in the presence of the co-solvent was effective in the extraction of chlorogenic acids to the same extent as the solvent-based method, from a qualitative point of view. Some slight differences are detected in the last part of the chromatogram, after 10 min as the retention time, where some small peaks are present in all chromatograms, differing from sample to sample. Due to the low area of the corresponding peaks, these differences were not further investigated, but tentative attributions to ubiquitous flavonoids such as quercetin and kaempferol could be carried out on the basis of the work by Nzekoue *et al.*³⁵

The UPLC-UV chromatogram shows a main peak characterized by a retention time of 4.1 min (peak 2 in Fig. 2) attributable to 5-CQA, the most abundant chlorogenic acid in coffee, as confirmed by co-injection of the commercial standard and further mass spectrometry results.

The assignments of the other peaks in the chromatograms were confirmed by mass spectrometry. Peaks 1, 2, and 3 were attributed to 3-caffeoyquinic acid (3-CQA), 5-caffeoyquinic acid (5-CQA) and 4-caffeoyquinic acid (4-CQA), respectively, and all were characterized by the presence of the molecular ion signal at $m/z = 353.5$ corresponding to $[M-H]^-$. Isomers were

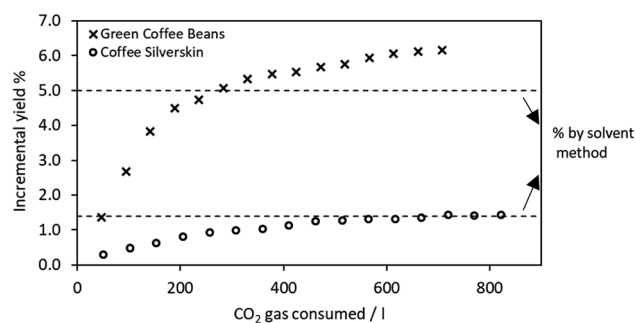


Fig. 1 Extraction kinetics (incremental yield vs. CO₂ utilization) during supercritical delipidization of GCB and CS. Dashed lines correspond to extraction yields obtained by the solvent method.

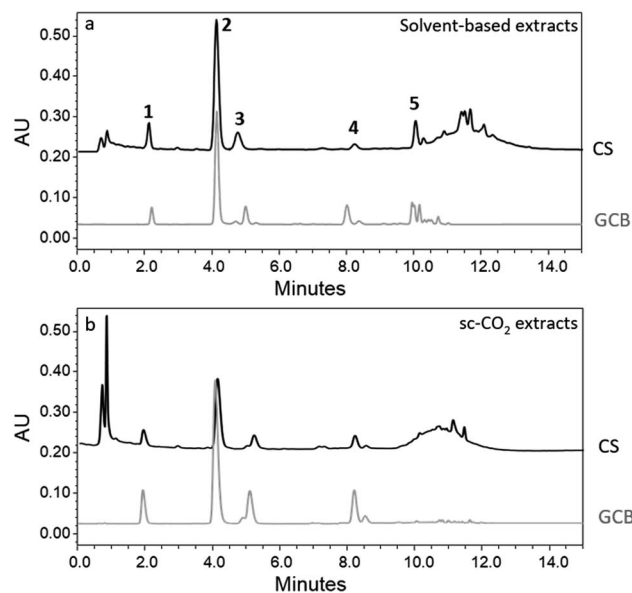


Fig. 2 UPLC-UV chromatographic profile monitored at 324 nm of: (a) conventional solvent-based extracts, (b) sc-CO₂ extracts. Peaks assignments: (1) 3-CQA, (2) 5-CQA, (3) 4-CQA, (4) FQA, (5) di-CQA.



recognized on the basis of the literature data.³⁶ Peak 4 was attributed to feruloylquinic acid (FQA), characterized by the presence of the molecular ion signal at $m/z = 367.7$ corresponding to $[M-H]^-$. Peak 5 was assigned to dicaffeoylquinic acids (di-CQA), due to the presence of the molecular ion signal at $m/z = 515.4$, corresponding to $[M-H]^-$, with no further investigation on the identity of the specific isomers.

Extraction results are consistent with the literature, where a previous work on the coffee silverskin extract obtained by acidified water (1% aqueous formic acid) at 70 °C for 1 h reported the presence of 3-CQA, 4-CQA, 5-CQA, and feruloylquinic acids.³⁷

Supercritical CO₂ has already been employed in the literature targeting phenolics from green coffee beans. De Azevedo employed ethanol and isopropyl alcohol as co-solvents in temperature ranges similar to this work, concluding that chlorogenic acids were extracted only from coffee beans when isopropyl alcohol was used. However, they evaluated chlorogenic acid extraction efficiency as poor, explained by the competition between these molecules and more polar components present in the coffee beans (e.g., sugars) and by the inability of the solvents to break down the caffeine–chlorogenic acid interactions.³⁸ In this work, the presence of an enhanced possibility of hydrogen bonding due to the presence of an ethanol/water co-solvent mixture, could have resulted in an increased competition and hence in breaking the hydrogen bonds between caffeine and chlorogenic acid.

Some other coffee by-products were also investigated targeting the chlorogenic acid by sc-CO₂, such as spent coffee grounds and coffee husks for valorization purposes. 15% of ethanol as co-solvent was the best choice, providing higher recovery of chlorogenic acids in supercritical conditions.³⁹

Concerning silverskin, to the best of our knowledge, no investigations have been reported in the literature related to chlorogenic acid extraction from this biomass by means of supercritical CO₂.

After the identification of the main phenolics, the most abundant chlorogenic acid in the samples, 5-CQA, was quantified.

As reported in Table 1, 5-CQA was quantified as 4.4% of green coffee beans and 0.29% of silverskin, when the extraction was run by means of a conventional solvent-based strategy. Results for green coffee beans are consistent with the literature, depending on the plant species, harvest time, and geographical origin of the plant. 2.7–6.5% of 5-CQA was reported by Ky *et al.*

employing GCB of species *C. liberica* var. *dewevrei* from the Ivory Coast.⁴⁰ Jeszka-Skowron *et al.* found chlorogenic acid content varying from 3.41% per dry mass in Arabica type from Laos or Rwanda to 8.16% in Robusta coffee from Indonesia.⁴¹

The content of chlorogenic acid quantified in coffee silverskin is also consistent with the literature. In particular, 5-CQA was extracted at 0.1–0.2% (w/w dry matter) from CS in the temperature range of 25–180 °C by Narita *et al.*⁴² Bresciani *et al.* reported similar content of chlorogenic acid, with an abundance of *circa*. 0.2% of 5-CQA in the dry biomass.⁴³ Subcritical water was also employed by Ginting *et al.*, targeting chlorogenic acids, reaching a percentage of 0.27% (w/w dry CS) at 147.8 °C.⁴⁴

The solvent-based extraction was supposed to be exhaustive in chlorogenic acids' complete extraction from both the delipidized biomasses. After assessing for consistency the effectiveness of the extraction of chlorogenic acids from GCB and CS with solvents with the literature, data were assumed to represent the reference when studying the recovery of the same phenolics by means of supercritical fluids.

5-CQA was then quantified in all extracts deriving from supercritical extraction and results were compared to the solvent-based extraction. When the water solubility of the sc-CO₂-samples was low (first extracts obtained during the initial cycles in sc-CO₂), they were first dissolved in dichloromethane and then counter-extracted in water. Only the water phase was injected in UPLC.

In terms of 5-CQA recovery, the results are displayed in Fig. 3.

Fig. 3a and b show the % of the cumulative yield of 5-CQA (on dry biomass) extracted over time, for each cycle of sc-CO₂ in the presence of the co-solvent. The dashed lines represent the % of 5-CQA (on dry biomass) extracted by means of the solvent method. The extractions of 5-CQA from both GCB and CS follow similar kinetics, with an initial absence of chlorogenic acids in the samples, then following the model again by da Silva *et al.* with a steep extraction rate followed by a lower extraction rate, where the mass transfer occurs mainly by diffusion inside the solid particles.³² The initial period where no 5-CQA is detected in the samples can be explained by the wetting of the biomass by the co-solvent being too low, rendering it unable to drag out (even with the aid of the gas flow) affine compounds, such as chlorogenic acids, in terms of polarity.

For GCB (Fig. 3a), the first three cycles were ineffective in 5-CQA extraction, whilst, from the 4th cycle onward, 5-CQA was positively found in the extracts. After the 16th cycle, 3.5% of 5-CQA (on dry biomass) was reached. This value is slightly lower,

Table 1 Quantification of 5-CQA in each sample

Sample name	Starting biomass	Extraction method	5-CQA (on dry biomass) (%)	5-CQA (in extract) (%)
GCB-solv	GCB	Solvent	4.4	11.7
CS-solv	CS	Solvent	0.29	2.1
GCB-scCO ₂	GCB	sc-CO ₂ + co-solvent	3.5	16.8 ^a
CS-scCO ₂	CS	sc-CO ₂ + co-solvent	0.37	33.9 ^a

^a The concentration of 5-CQA changes in each extract from the different successive sc-CO₂ cycles. In the table are reported the highest values, obtained during the 11th sc-CO₂ cycle for GCB and during the 6th sc-CO₂ cycle for CS.



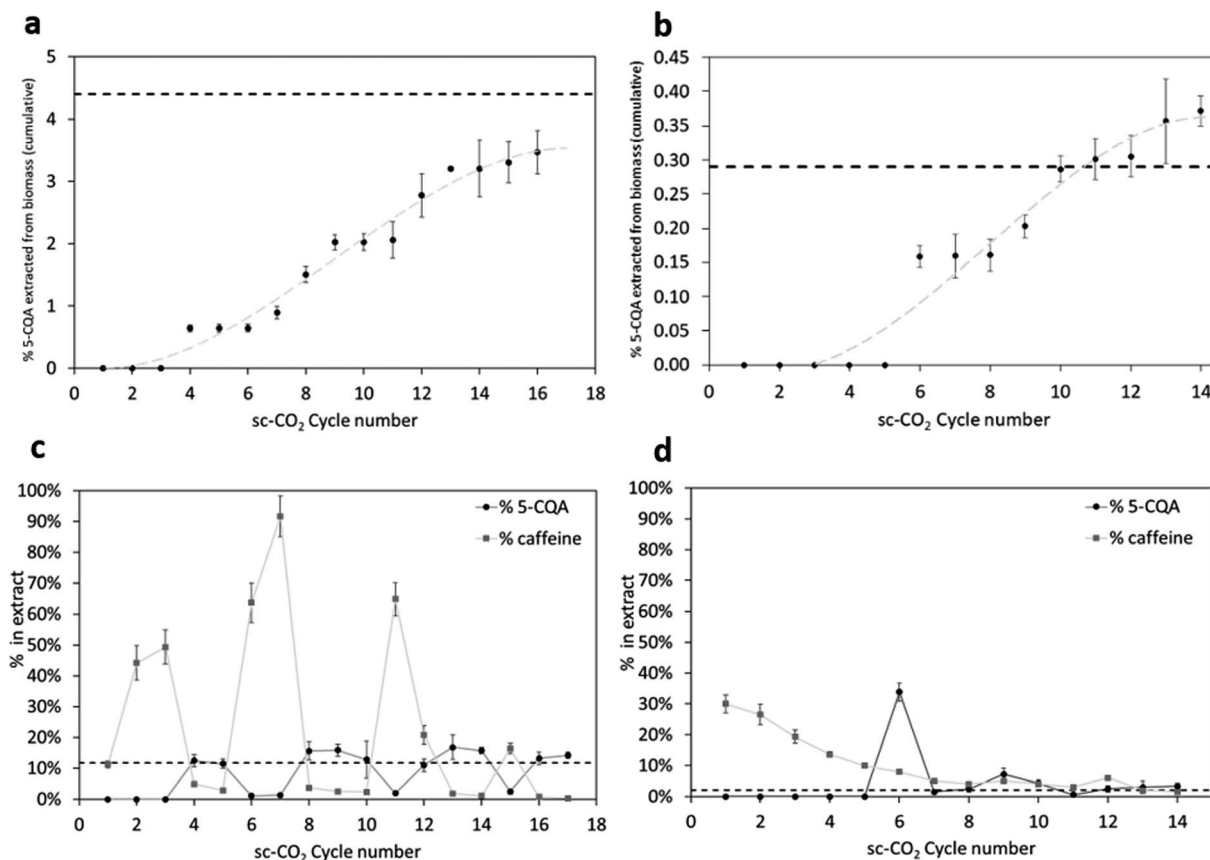


Fig. 3 Extraction kinetics of 5-CQA (incremental yield on dry biomass vs. progressive extraction cycle) from (a) delipidized GCB and (b) delipidized CS. % of 5-CQA and % of caffeine in each extract from (c) delipidized GCB and (d) delipidized CS. Dashed lines represent the % of 5-CQA obtained by solvent-method.

but comparable, to the % of 5-CQA obtained by solvent-based extraction, equal to 4.4%. The extraction was not continued further, due to the amount of co-solvent already used. Further use of the organic solvent, higher than 10 times the mass of the solid to be extracted, would not be reasonable when applying a “green technique” such as supercritical fluid extraction. Regarding chlorogenic acid recovery from CS extraction, as shown in Fig. 3b, the first five cycles provided extracts in which 5-CQA was absent. Its extraction began with the 6th cycle. From the 11th cycle onward, the amount of extracted 5-CQA was even higher than that obtained by means of a solvent strategy, reaching a maximum extraction yield of 0.37% (the content of 5-CQA in CS when the extraction was run in conventional conditions was 0.29%). The micronization of the sample, together with the enhanced penetrability of the fluid mixture in supercritical conditions, might have facilitated the mass transfer process of chlorogenic acids. For the same reason as above (not to exceed the logical amount of co-solvent employed), the extraction was stopped on the 14th cycle.

Besides the 5-CQA recovery, it was also interesting to quantify, cycle by cycle during the sc-CO₂ extraction process, the 5-CQA content of each extract, as a necessary dataset to study how the selectivity towards chlorogenic acids varies over the time of sc-CO₂ extraction. Explicative data are displayed in Table 1 in

the last column, where it appears evident that supercritical extracts are enriched in chlorogenic acids more than the solvent-based samples, confirming the increased selectivity of the sc-CO₂ technique. GCB solvent-based extract contains only 11.7% of 5-CQA, while one of the extracts obtained in sc-CO₂ reaches 16.8%. Results are even more rewarding when considering coffee silverskin: solvent-based extract contains only 2.1% of 5-CQA, while an sc-CO₂ extract reaches 33.9%.

Apart from these data, relative to the most rewarding sc-CO₂ extracts (in terms of 5-CQA content), the % of 5-CQA in each extract changes during the extraction kinetics under supercritical conditions. Fig. 3c and d are explicative. The dashed lines represent the 5-CQA enrichment in the conventional extracts, while each point of the plot represents the sc-CO₂ samples' enrichment in terms of 5-CQA content. For GCB, the trend is oscillating. This might be explained by the non-micronized biomass particles hindering the mass transfer processes and hence releasing phenolics in a non-linear way, being trapped inside different channels' sizes and morphologies. Anyhow, most of the extracted samples display enhanced 5-CQA content compared to the solvent-based extract, with most of the points above the dashed line. Interestingly, the caffeine content follows a complementary behavior: samples characterized by



a high content of caffeine are composed of low content of 5-CQA and *vice versa*, as clearly visible in Fig. 3c and d.

Caffeine is characterized by solubilities in organic solvents that are completely different from the solubilities of chlorogenic acids, almost insoluble in most organic solvents.^{45,46}

When the extraction was performed starting from GCB, caffeine was immediately extracted during the first extraction cycles (caffeine has a higher solubility in pure sc-CO₂ than chlorogenic acids), which were those characterized by a lower wetting of the matrix by the co-solvent. Over time (and over cycles) increasing the wetting of the matrix corresponded to an easier extractability of polar compounds, and in fact, chlorogenic acids were extracted to a higher extent than caffeine, flushed out from the vessel together with the co-solvent. At the end of this process, almost complete co-solvent removal corresponded to a decrease in the wetting of the matrix, thus making caffeine more easily extractable in the subsequent cycles. This behavior can explain the alteration in the presence of more caffeine (or more chlorogenic acids) in the extracts, hence related to a higher/lower presence of the co-solvent in the vessel. In the case of CS, due to the lower percentages of caffeine and chlorogenic acids present in the biomasses, this phenomenon might have been hidden.

CS extracts behave in a completely different way. The first extract containing 5-CQA (6th cycle) is also the highest one in terms of 5-CQA content (33.9%), exceeding more than 15 times the content of the solvent-based extract (2.1%). About 50% by weight of the total 5-CQA present in the dry biomass was extracted during the 6th cycle, likely facilitated by the micronized particles that favor the contact between the fluid and the biomass, and hence the mass transfers. These results point out the possibility to interrupt the extraction at this cycle, to collect an extract that is selectively enriched in 5-CQA. On the other hand, caffeine follows a decreasing trend, being mostly extracted at the beginning of cycles, then decreasing over time and over extraction cycles.

Overall, aiming at extracting polar compounds such as phenolics in sc-CO₂, the presence of a co-solvent (such as ethanol) as a polarity modifier is demonstrated to be indispensable, justified by the increase in the solubility of polar compounds in the mixture ethanol/sc-CO₂, compared to the solubility in pure sc-CO₂.^{41,47}

Carbohydrates, phenolics, and conjugate profiles

As discussed in the previous section, not only does the solubility of the target compound increase with the use of co-solvent, but also the diversity of other compounds solubilized by the same solvent, which increases the yield as mentioned above, at the expense of reducing the process selectivity.

In a biorefinery concept, the effect of co-extraction has an interesting economical outcome from a process perspective. The use of conventional solvent extraction methods could solubilize other molecules apart from the targeted phenolics due to them having similar physicochemical properties or because of molecular interactions of the phenolics with other macromolecules in the biomass such as carbohydrates or

lignin. The effect of sc-CO₂ extraction has been proven for the disruption of the plant cell wall as a pretreatment but also for the solubilization of carbohydrates.^{48–50} Based on that, considering the potential co-extraction of different biomass fractions from the same treatment, the characterization of co-extracted species was performed on both conventional extracts and on the 6th cycles of sc-CO₂ extracts, to understand the profile of the obtained samples. In the case of conventional solvent extraction, the method is based on the capacity of the specific solvent to solubilize polar compounds from a matrix. However, the sc-CO₂ also promotes a physical extraction process by pressurized conditions on the flow rate in the reactor, implying that other polar compounds from the biomass can be co-extracted. Since carbohydrates are the main components of GCB and CS biomasses, they have been analyzed to assess and quantify their co-extraction together with the polar phenolics. For this determination, the initial extracts were characterized by high-performance anion-exchange chromatography-pulsed amperometric detection analysis (HPAEC-PAD) in order to study the presence of free sugars. In addition, acidic hydrolysis of the extracts was performed with trifluoroacetic acid (TFA), aiming to check the presence of the complex carbohydrate molecules in the extracts derived from the biomass used in the study. The comparison allows an understanding of the complexity of the carbohydrates in monomeric, oligomeric, and polymeric forms co-extracted during the process. In the case of sc-CO₂ conditions, the carbohydrate co-extraction is more important in the case of GCB, as displayed in Fig. 4a. After TFA hydrolysis, a clear increase in the complexity of monosaccharide profiles of the different samples was observed, as displayed in Fig. 4b. For instance, the presence of other sugars such as fucose, arabinose, rhamnose, xylose, and mannose was detected by HPAEC-PAD. The sugars confirmed the most probable presence of galactoglucomannans, glucans, arabinogalactans, and, in the case of CS, pectins were also deduced by the presence of rhamnose. This is in agreement with previous studies on CS for the extraction of pectins.^{51,52} The occurrence of those sugars was also followed by an increase of the solubilized glucose by the hydrolytic effect of the polymeric carbohydrate species. However, the galactose content seems to be affected by the TFA hydrolysis, more clearly in the GCB samples; in the case of GCB-sc-CO₂, the galactose after TFA hydrolysis was 20% of the initial free amount (Fig. 4a). The partial degradation of free sugars has been presented previously and the susceptibility of degradation in the case of arabinose and galactose more than in the case of free glucose has been discussed.^{53,54} From the SEC analysis, it was highlighted that most of the extracts were of a lower molecular size until 200 Da, clearly represented in Fig. 4c, with most of the signal in the small range of the molecular size. The greater increase of monosaccharides after TFA hydrolysis from the CS samples is directly correlated with the presence of longer polymeric fractions, close to 100 000 Da presented in the magnified image in Fig. 4d. The molecular distribution of the CS extract showed the presence of longer polymeric fractions that were co-extracted by both conventional solvent and sc-CO₂. This also correlates with the lower amounts of free sugars shown in Fig. 4a. In the case of GCB, the molecular distribution



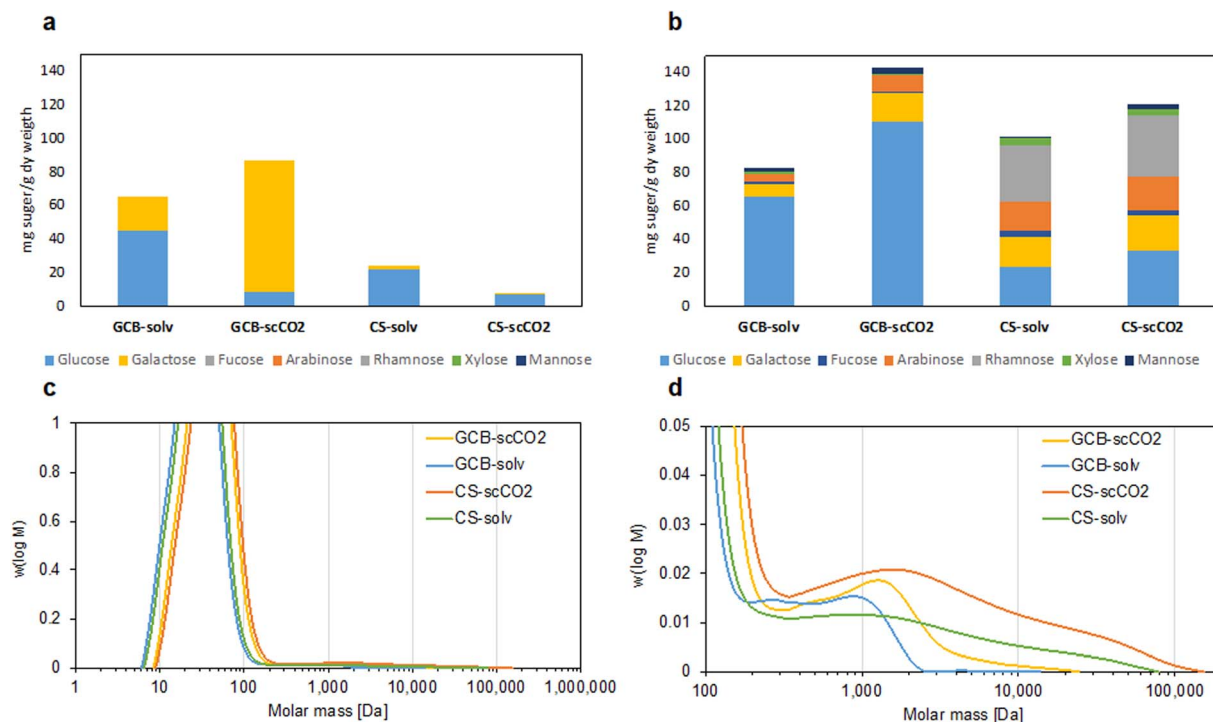


Fig. 4 Monosaccharide profile and molecular weight distribution (Da) of the extracts from conventional solvent extraction and supercritical CO₂ + EtOH. (a) Carbohydrate composition after direct solubilization; (b) carbohydrate composition after TFA hydrolysis; (c) complete molecular weight distribution; (d) oligomeric and polymeric molecular weight distribution of the extracts after direct solubilization (zooming of (c)).

of the extracts is in the range of oligomers and short polymers, with populations around 2000 Da for sc-CO₂ extraction and 10 000 Da for the conventional solvent extraction. The use of LC-ESI-MS before and after saponification of the samples allowed the identification of certain alternative phenolic molecules in monomeric forms such as ferulic acid at $m/z = 195 [M + H]^+$ and $m/z = 177$, corresponding to its dehydrated adduct $[M + H - H_2O]^+$, caffeic acid at $m/z = 181 [M + H]^+$ and $m/z = 163 [M + H - H_2O]^+$, as well as certain dimeric forms such as di-ferulic acid at $m/z = 369 [M + H - H_2O]^+$ and $m/z = 351 [M + H - 2H_2O]^+$ as reported in Table S1.† These fractions, commonly present in other lignocellulosic materials such as cereal brans,⁵⁵ were also identified by comparing them with available standards (reported in Fig. S2†). Moreover, it was possible to identify the presence of carbohydrate-phenolic conjugates at $m/z = 509$ (feruloyl hexo-pentose), $m/z = 345$ (caffeoyl hexose), and $m/z = 329$ (feruloyl pentose), which were distinctly present in the CS samples and were structures that had been presented previously in other types of biomasses such as tomatoes or cereals^{56,57}. In the case of GCB, the possible presence of conjugates is reported only after sc-CO₂ treatment, at $m/z = 499$ (caffeoyl hexo-pentose) and $m/z = 367$ (caffeoyl hexose), demonstrating the increased potential of co-extraction of this technique (Fig. S3†). The different elution times could correspond with different linkages in the sugar-phenolic conjugates. These results correlate with the previously mentioned co-extraction of phenolics with carbohydrates reported in Fig. 4. In the case of sc-CO₂, it is the first time, to our knowledge, that these conjugates have been recognized after extraction. Other studies using subcritical

water extraction have proven the protection of the link between carbohydrates and phenolic moieties present in natural cell wall extracts.²⁷ These conjugates could have great potential as prebiotics or as building blocks for functional materials since the phenolics are able to provide their characteristic antioxidant properties to the carbohydrates fraction.^{58,59}

Conclusions

The results highlighted the ability of the supercritical fluid to efficiently target polar compounds, such as chlorogenic acids, in the presence of other beneficial co-extracted compounds, minimizing the excessive use of organic solvents and avoiding the presence of inorganic or organic acids conventionally employed when solvent-based methods are used. To the best of our knowledge, no other works are present in the literature targeting chlorogenic acids from coffee silverskin employing the sc-CO₂ strategy. From the view of the application, the feasibility of the technique of supercritical CO₂ in the presence of a co-solvent should meet a solvent usage threshold in order to be environmentally and economically convenient in comparison to conventional solvent-based methods. For example, less use of an organic co-solvent is advisable and should be the object of future studies and technological improvements, in order to further reduce the use of ethanol/water and to decrease the time and energy requirements for solvent removal.

The scalability of sc-CO₂ technology is not well-documented in the literature, and future studies are needed to better



understand the practical application of this technology and related implications when co-solvents are employed.

The advantage of sc-CO₂ connects also with the possibility to recover extracts in different batches over time since it requires alternation of static and dynamic periods. Extract separation over time allows a selection and isolation of the most promising as the ones characterized by high polyphenol content and a negligible amount of caffeine. This profile is desirable since caffeine itself has some biological properties that are not always compatible with the nutraceutical purposes of extracts containing chlorogenic acids.

Overall, the results further confirm that CS can be considered a good raw material for the production of chlorogenic acid-enriched extracts, with the advantage of being cheaper and more sustainable than green coffee beans. A very interesting application could be related to prebiotics in food products such as yogurt and derivatives since coffee chlorogenic acids have already been demonstrated to increase the growth of total bacteria to a similar magnitude to fructooligosaccharides, which are the positive prebiotic control.⁶⁰ This assertion could also be supported by the fact that a considerable proportion of ingested chlorogenic acids reaches the large intestine, giving them the possibility of exerting beneficial effects in the large gut.⁶¹ Upcoming studies are desirable to assess the nutraceutical and antioxidant profiles of these extracts in order to confirm the potential commercial interest in this ambitious and virtuous “waste-to-products” valorization chain.

Author contributions

Data curation: S. M., A. M., R. N. and A. J. Q.; funding acquisition: L. V., S. M. and A. J. Q.; investigation: A. M., R. N., S. M. and A. J. Q.; project administration: L. V., S. M. and A. J. Q.; supervision: L. V., S. M. and A. J. Q.; conceptualization: L. V., S. M., R. N. and A. J. Q.; validation: A. M., R. N., S. M. and A. J. Q.; visualization: A. M., R. N., S. M. and A. J. Q.; writing – original draft: S. M. and A. J. Q.; writing – review and editing: A. M., R. N., L. V., S. M. and A. J. Q.

Conflicts of interest

There are no conflicts to declare.

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References

- I. Vural Gursel, B. Elbersen, K. P. H. Meesters and M. van Leeuwen, *Sustain*, 2022, **14**, 1–21.
- C. Zhou and Y. Wang, *Sci. Technol. Adv. Mater.*, 2020, **21**, 787–804.
- S. O. Essien, B. Young and S. Baroutian, *Trends Food Sci. Technol.*, 2020, **97**, 156–169.
- A. Vandeponseele, M. Draye, C. Piot and G. Chatel, *Green Chem.*, 2020, **22**, 8544–8571.
- E. Ibáñez, J. A. Mendiola and M. Castro-Puyana, *Encyclopedia of Food and Health*, 2015, pp. 227–233.
- R. M. Couto, J. Fernandes, M. D. R. G. da Silva and P. C. Simões, *J. Supercrit. Fluids*, 2009, **51**, 159–166.
- M. D. Luque de Castro and M. M. Jiménez-Carmona, *TrAC, Trends Anal. Chem.*, 2000, **19**, 223–228.
- M. D. Luque de Castro and M. T. Tena, *TrAC, Trends Anal. Chem.*, 1996, **15**, 32–37.
- M. W. Tham and K. C. Liew, *Eur. J. Wood Wood Prod.*, 2014, **72**, 67–72.
- S. R. Falsafi, S. P. Bangar, V. Chaudhary, E. Hosseini, Z. Mokhtari, A. C. Karaca, M. K. Samota, D. Goswami, V. Krishnan, G. Askari and H. Rostamabadi, *Carbohydr. Polym.*, 2022, **298**, 120074.
- S. Yilmaz-Turan, P. Lopez-Sanchez, A. Jiménez-Quero, T. S. Plivelic and F. Vilaplana, *Food Hydrocoll.*, 2022, **28**, 107575.
- V. Manasa, A. Padmanabhan and K. A. Anu Appaiah, *Waste Manag.*, 2021, **120**, 762–771.
- N. Kumar, R. Weldon and J. G. Lynam, *Biocatal. Agric. Biotechnol.*, 2021, **36**, 102145.
- A. Procentese, F. Raganati, G. Olivieri, M. E. Russo and A. Marzocchella, *Appl. Microbiol. Biotechnol.*, 2019, **103**, 1021.
- R. Nasti, A. Galeazzi, S. Marzorati, F. Zaccheria, N. Ravasio, G. L. Bozzano, F. Manenti and L. Verotta, *Waste and Biomass Valorization*, 2021, **12**, 6021–6033.
- V. S. Ribeiro, A. E. Leitão, J. C. Ramalho and F. C. Lidon, *Food Res. Int.*, 2014, **61**, 39–47.
- F. Rodrigues, A. Palmeira-de-Oliveira, J. Das Neves, B. Sarmento, M. H. Amaral and M. B. P. P. Oliveira, *Pharm. Biol.*, 2015, **53**, 386–394.
- Y. Narita and K. Inouye, *Food Res. Int.*, 2014, **61**, 16–22.
- R. Nasti, A. Galeazzi, S. Marzorati, F. Zaccheria, N. Ravasio, L. B. Giulia, F. Manenti and L. Verotta, *Waste and Biomass Valorization*, 2021, **12**, 6021.
- G. Navarra, M. Moschetti, V. Guarrasi, M. R. Mangione, V. Militello and M. Leone, *J. Chem.*, 2017, **2017**, 6435086.
- M. L. Suárez-Quiroz, A. Alonso Campos, G. Valerio Alfaro, O. González-Ríos, P. Villeneuve and M. C. Figueroa-Espinoza, *J. Food Compos. Anal.*, 2014, **33**, 55–58.
- A. Rai, R. Shukla, S. Sawant, R. Shetye, D. Boppte and H. Gavandi, *NCIFEH Conference Proceeding*, 2018, p. 269.
- L. S. McKee, H. Sunner, G. E. Anasontzis, G. Toriz, P. Gatenholm, V. Bulone, F. Vilaplana and L. Olsson, *Biotechnol. Biofuels*, 2016, **9**, 1–13.
- P. Comino, H. Collins, J. Lahnstein, C. Beahan and M. J. Gidley, *Food Hydrocoll.*, 2014, **41**, 219–226.
- C. Antoine, S. Peyron, V. Lullien-Pellerin, J. Abecassis and X. Rouau, *J. Cereal. Sci.*, 2004, **39**, 387–393.
- C. Menzel, C. González-Martínez, A. Chiralt and F. Vilaplana, *Carbohydr. Polym.*, 2019, **214**, 142–151.



- 27 A. C. Ruthes, A. Martínez-Abad, H. T. Tan, V. Bulone and F. Vilaplana, *Green Chem.*, 2017, **19**, 1919–1931.
- 28 J. M. Del Valle, M. Jiménez and J. C. De la Fuente, *J. Supercrit. Fluids*, 2003, **25**, 33–44.
- 29 H. B. Coats and M. R. Wingard, *J. Am. Oil Chem. Soc.*, 1950, **27**, 93–96.
- 30 C. G. Pereira and M. A. A. Meireles, *Food Bioprocess Technol.*, 2010, **3**, 340–372.
- 31 P. A. Uwineza and A. Waśkiewicz, *Molecules*, 2020, **25**, 3847.
- 32 R. P. F. F. da Silva, T. A. P. Rocha-Santos and A. C. Duarte, *TrAC, Trends Anal. Chem.*, 2016, **76**, 40–51.
- 33 R. N. Patel, S. Bandyopadhyay and A. Ganesh, *Energy Convers. Manag.*, 2011, **52**, 652–657.
- 34 S. I. M. L. F. Ballesteros and J. A. Teixeira, *Food Bioprocess Technol.*, 2014, **7**, 3493.
- 35 F. K. Nzekoue, S. Angeloni, L. Navarini, C. Angeloni, M. Freschi, S. Hrelia, L. A. Vitali, G. Sagratini, S. Vittori and G. Caprioli, *Food Res. Int.*, 2020, **133**, 109128.
- 36 L. Regazzoni, F. Saligari, C. Marinello, G. Rossoni, G. Aldini, M. Carini and M. Orioli, *J. Funct. Foods*, 2016, **20**, 472.
- 37 L. Bresciani, L. Calani, R. Bruni, F. Brighenti and D. Del Rio, *Food Res. Int.*, 2014, **61**, 196–201.
- 38 A. B. A. De Azevedo, P. Mazzafera, R. S. Mohamed, S. A. B. Vieira De Melo and T. G. Kieckbusch, *Braz. J. Chem. Eng.*, 2008, **25**, 543–552.
- 39 K. S. Andrade, R. T. Gonálvez, M. Maraschin, R. M. Ribeiro-Do-Valle, J. Martínez and S. R. S. Ferreira, *Talanta*, 2012, **88**, 544–552.
- 40 C. L. Ky, M. Noirot and S. Hamon, *J. Agric. Food Chem.*, 1997, **45**, 786–790.
- 41 A. Daraee, S. M. Ghoreishi and A. Hedayati, *J. Supercrit. Fluids*, 2019, **144**, 19–27.
- 42 Y. Narita and K. Inouye, *Food Chem.*, 2012, **135**, 943–949.
- 43 L. Bresciani, L. Calani, R. Bruni, F. Brighenti and D. Del Rio, *Food Res. Int.*, 2014, **61**, 196–201.
- 44 A. R. Ginting, T. Kit, W. Mingvanish and S. P. Thanasupsin, *Sustainability*, 2022, **14**, 8435.
- 45 Q. V. Vuong and P. D. Roach, *Separ. Purif. Rev.*, 2014, **43**, 155–174.
- 46 K. Dibert, E. Cros and J. Andrieu, *J. Food Eng.*, 1989, **10**, 1–11.
- 47 Y. Wu, B. Liu, Y. Chang and Q. Wang, *J. Liq. Chromatogr. Relat. Technol.*, 2015, **38**, 443–450.
- 48 L. Vaahtera, J. Schulz and T. Hamann, *Nat. Plants*, 2019, **5**, 924–932.
- 49 Y. Jiang, Y. Feng, B. Lei and H. Zhong, *Int. J. Biol. Macromol.*, 2020, **161**, 1506–1515.
- 50 T. Gu, M. A. Held and A. Faik, *Environ. Technol.*, 2013, **34**, 1735–1749.
- 51 L. Wen, Z. Zhang, M. Zhao, R. SenthamaraiKannan, R. B. Padamati, D. W. Sun and B. K. Tiwari, *Int. J. Food Sci. Technol.*, 2020, **55**, 2242–2250.
- 52 V. Gottstein, M. Bernhardt, E. Dilger, J. Keller, C. M. Breitling-Utzmann, S. Schwarz, T. Kuballa, D. W. Lachenmeier and M. Bunzel, *Foods*, 2021, **10**, 1–18.
- 53 S. I. Mussatto, L. M. Carneiro, J. P. A. Silva, I. C. Roberto and J. A. Teixeira, *Carbohydr. Polym.*, 2011, **83**, 368–374.
- 54 Y. Mao, R. Lei, J. Ryan, F. Arrutia Rodriguez, B. Rastall, A. Chatzifragkou, C. Winkworth-Smith, S. E. Harding, R. Ibbett and E. Binner, *Food Chem.*, 2019, **2**, 100026.
- 55 R. C. Rudjito, A. C. Ruthes, A. Jiménez-Quero and F. Vilaplana, *ACS Sustain. Chem. Eng.*, 2019, **7**, 13167–13177.
- 56 R. C. Rudjito, A. Jiménez-Quero, M. Hamzaoui, S. Kohnen and F. Vilaplana, *Green Chem.*, 2020, **22**, 8337–8352.
- 57 A. Carrillo-López and E. Yahia, *J. Food Sci.*, 2013, **78**, 1839–1844.
- 58 J. Ou and Z. Sun, *J. Funct. Foods*, 2014, **7**, 90–100.
- 59 S. Yilmaz-Turan, A. Jiménez-Quero, C. Menzel, D. M. de Carvalho, M. E. Lindström, O. Sevastyanova, R. Moriana and F. Vilaplana, *Carbohydr. Polym.*, 2020, **250**, 116916.
- 60 L. Wang, X. Pan, L. Jiang, Y. Chu, S. Gao, X. Jiang, Y. Zhang, Y. Chen, S. Luo and C. Peng, *Front. Nutr.*, 2022, **9**, 1–22.
- 61 F. Tomas-Barberan, R. García-Villalba, A. Quartieri, S. Raimondi, A. Amaretti, A. Leonardi and M. Rossi, *Mol. Nutr. Food Res.*, 2014, **58**, 1122–1131.

