

An Insight on Salting-out Assisted Liquid–Liquid Extraction for Phytoanalysis

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Abstract

Introduction

Salting-out assisted liquid–liquid extraction (SALLE) is a technique that, although simple and not requiring any complex equipment, is very powerful and versatile. It has obtained growing interest in bioanalysis particularly when combined with chromatographic techniques.

Objectives

Herein, fennel seeds (*Foeniculum vulgare* Mill.) were used as a case-study to show the application of SALLE in phytochemical analysis.

Material and Methods

SALLE combined with HPLC-UV-MS/MS and GC–MS.

Results

By HPLC-UV-MS/MS analysis of the organic extract it was possible to identify various phenolic compounds, including quercetin derivatives, caffeic acid, p-coumaric acid and chlorogenic acid. The main compounds identified by GC–MS were estragole, fenchone, anisaldehyde, anethole, benzaldehyde, camphor and apiole.

Conclusion

HPLC and GC analysis of the extracts showed that it is possible to isolate, in only one step, a wide range of compounds with distinct properties, allowing a detailed phytochemical analysis.

Keywords

chromatography; *Foeniculum vulgare*; phytochemical characterisation; polyphenols; sample preparation

Introduction

The extraction and characterisation of active compounds from plants have resulted in the discovery of several compounds with therapeutic value or even new sources for already known compounds (Huie, 2002). For the identification of new compounds, sample preparation is a crucial first step (Huie, 2002). Therefore, considering the economic importance of plants and the increasing relevance of medicinal plants, it is not surprising that a lot of research in this field has employed techniques like solid-phase micro-extraction, supercritical-fluid extraction, pressurised-liquid extraction, microwave-assisted extraction, solid-phase extraction, and surfactant-mediated extraction (Huie, 2002). Another option, somewhat simpler and cheaper yet very powerful, is salting-out assisted liquid-liquid extraction (SALLE) (Valente et al., 2013a). The salting-out effect has an interesting application for the creation of liquid biphasic systems in mixtures composed of water and water-miscible organic solvents. Adding salts, two distinguishable liquid phases are formed in which the upper phase is mainly composed of the organic solvent. Although its initial applications can be traced to the 1970s (Matkovich and Christian 1973), new analytical methodologies using the salting-out effect continue to be developed, as is the case in the field of biotechnology where the precipitation of proteins in biological samples is aimed (Ahmed and Mahmoud, 2015; Yang et al., 2015). Moreover, it is not in any way misplaced to state that the QuEChERS (portmanteau word formed from “Quick, Easy, Cheap, Effective, Rugged and Safe”) method embody the SALLE concept (Silva et al., 2012; Valente et al., 2013b). Of course, this type of sample preparation, where analytes are concentrated and incorporated in cleaner matrices that are more column-friendly, is ideal for good chromatographic separations targeted for chemical analysis.

Fennel (*Foeniculum vulgare* Mill.) is a popular plant in Mediterranean cuisine; it is crunchy with a minor sweet flavour, and owns a pleasant odour (Steuer and Schulz, 2003; Díaz-Maroto et al., 2006; Miguel et al., 2010; Azevedo et al., 2012). Most of the plant (bulb, leaves, seeds and stalk) is edible, with a considerable content of vitamin C and fibre. Fennel belongs to the Apiaceae family (also known as Umbelliferae), the same that includes plants like parsley, parsnip, cumin, carrot and many other widely known plants, not all of them edible as is clearly the case of hemlock. Many health applications are attributed to fennel, from antispasmodic to diuretic or anti-inflammatory properties (Gori et al., 2012); however most of these claims have not been subject to serious studies. Nevertheless some trials can be found in the literature advocating, for example, minor therapeutic effects for dysmenorrhoea (Jahromi et al., 2003) or hirsutism (Akha et al., 2014).

Herein, SALLE was applied to the phytochemical study of fennel seeds as a proof-of-concept of the capabilities of this technique applied to plants' chemical analysis. Extracts were then subsequently analysed by liquid and gas chromatography (LC and GC), both coupled to mass spectrometry (MS) detection.

Material and methods

Chemicals and samples

High-purity water (resistivity not lower than 18.2 M Ω cm), from a Direct-Q 3 UV water purification system (Millipore Iberia, Madrid, Spain), was used for the preparation of solutions, chromatographic eluents and glassware washing. HPLC gradient methanol (Fisher, Waltham, MA) was used for standard solutions preparation and as chromatographic mobile phase. HPLC gradient grade acetonitrile (Fisher) was used for the extractions. Formic acid (99%, VWR, Radnor, PA) was used to acidify the aqueous mobile phase, prepared with high-purity water. All eluents were filtered through a Nylon filter (0.45 μ m pore size, Whatman, Pittsburgh, PA) and degassed before use.

Caffeic acid (98%), chlorogenic acid (95%), p-coumaric acid (98%), quercetin dihydrate (98%), trans-anethole (99%) were purchased from Sigma-Aldrich (Steinheim, Germany). Stock standard solutions of these compounds were prepared in methanol. Ammonium acetate (Merck, Darmstadt, Germany), sodium carbonate (Panreac, Barcelona, Spain), potassium carbonate (Merck), dihydrogen phosphate (Pronalab, Tlalnepantla, Mexico) and ammonium sulphate ((NH₄)₂SO₄) (BDH Prolabo, Radnor, PA) were of analytical grade and were used without further purification. Xanthohumol (98%, Hopsteiner, Mainburg, Germany) was prepared in ethanol (95.5%, Merck) for use as internal standard. Terpene standards (CAN-TERP-MIX1 and CAN-TERP-MIX2) used for GC analysis were obtained from SPEX Europe (Stanmore, UK).

Fennel (*Foeniculum vulgare* Mill.) dried seed samples were purchased in local markets.

Extraction procedure

The experimental procedure used for the development of this work was based on previous procedures using the SALLE technique, developed by our research group (Valente et al., 2013a) (Figure 1). In 50 mL plastic tubes, fennel seeds (0.5 g) were extracted with 10 mL of solvent (50% of acetonitrile in water) and 1 g of (NH₄)₂SO₄ by decoction using mechanical stirring for 37 min, at room temperature. After the extraction, the tubes were centrifuged for 2 min at 7750 relative centrifugal force (RCF) for phase separation and an aliquot of the upper (organic) phase was collected for HPLC analysis. For GC analysis, an extra step was included to remove any water residue and possible interfering compounds (sugars, pigments, etc.); 1 mL of organic extract was added to a dispersive solid-phase extraction (d-SPE) tube containing 150 mg magnesium sulphate (MgSO₄), 50 mg of primary and secondary amine (PSA) and 50 mg C18. All analyses were carried out in triplicate.

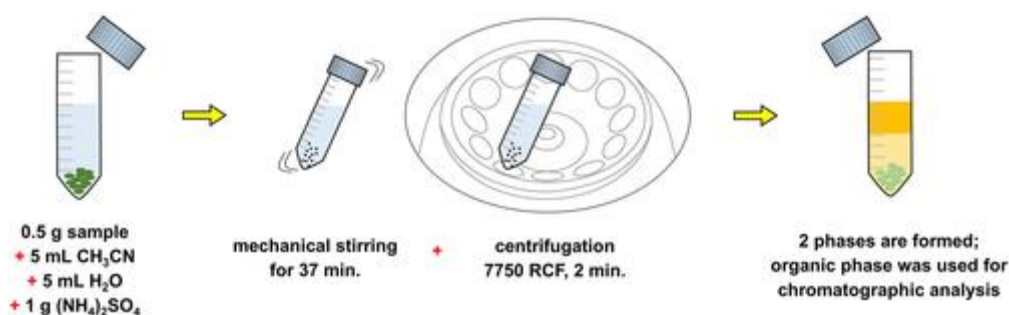


figure 1- Schematics of the salting-out assisted liquid-liquid extraction (SALLE) experimental procedure

HPLC-UV analysis

Fennel extracts were analysed by HPLC-UV using a Jasco (Jasco Corporation, Tokyo, Japan) chromatographic system consisting of a low-pressure quaternary gradient unit (model PU-2089 Plus) with an on-line degasser (model DG-1580-54), an autosampler (model AS-950) and a photodiode array detector (model MD-1510). The system control and data analysis was done with Jasco ChromPass Chromatography Data software (version 1.7.403.1). Separations were achieved at room temperature on a Gemini C18 column (250 mm × 4.6 mm; 5 µm particle size) and a guard column (4 mm × 3.0 mm) from Phenomenex (Torrance, CA). The mobile phase, under gradient conditions at 0.8 mL/min, was composed of (A) methanol and (B) 0.1% formic acid in water. Gradient programme was set as follows: 0 to 40 min, linear increase from 10% to 30% of A; 40 to 60 min, increase to 45% of A; 60 to 90 min, linear increase to 100% A and conditions maintained for 5 min; return to initial conditions in 15 min and conditions maintained for 10 min before the next injection. Sample injection volume was 20 µL. The photodiode array detection was conducted by scanning between 190 and 600 nm. Analytes in each sample were identified by comparing their retention times and UV-vis spectra with those of standard compounds, and by HPLC-MS/MS analysis. A chromatogram obtained for the HPLC-UV analysis (at 280 nm) of an extract is shown in Figure 2.

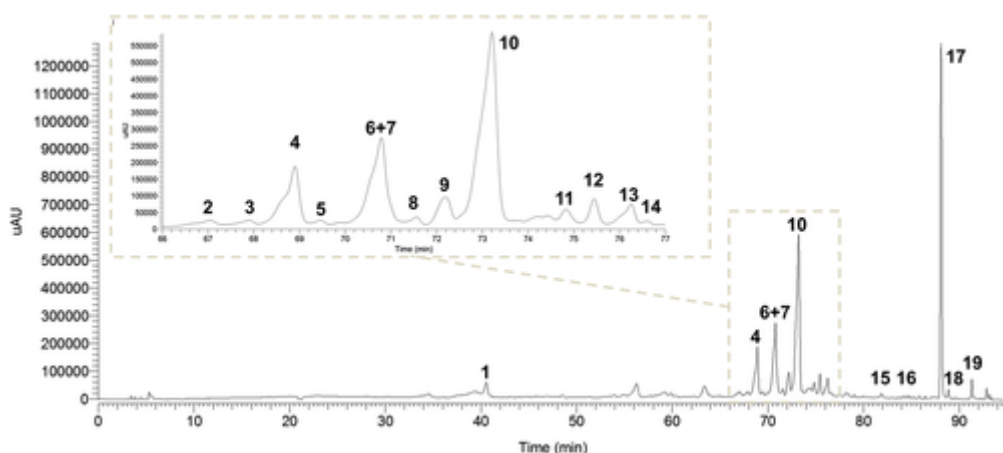


figure 2 -

HPLC-UV chromatogram (280 nm) of fennel seeds extract. Numbered peaks correspond to tentatively identified compounds that can be ascribed to Table 2

HPLC-DAD-MS/MS analysis

The qualitative study of the extracts' composition was performed by HPLC coupled on-line with electrospray ionisation (ESI) MS. The HPLC system (Thermo Electron Corporation, Waltham, MA) was composed of a low-pressure quaternary pump with autosampler (200-vial capacity sample) and a diode array detector (DAD) (Finnigan Surveyor Plus, Thermo Fisher Scientific). A Gemini C18 column (150 mm × 4.6 mm; 3 µm particle size) and a guard column (4 mm × 3.0 mm) from Phenomenex were used at room temperature. Separations were achieved in the same conditions as for HPLC-UV analysis but at a flow rate of 0.4 mL/min with injection of 25 µL of sample. A quadrupole ion-trap mass spectrometer (Finnigan LCQ Deca XP Plus) equipped with an ESI source was used in the following conditions: capillary temperature, 325°C; source voltage, 5.0 kV; capillary voltage, 4.0 V; sheath gas (N₂) flow at 40 arbitrary units and auxiliary gas (N₂) flow at 15 arbitrary units. The mass detection was performed in the range 160–1000 m/z. XCalibur software Version 2.2 (Thermo Electron Corporation) was used for data acquisition and processing.

GC–MS analysis

Fennel extracts were characterised by GC–MS analysis using a Thermo Scientific Trace 1300, ISQ Single Quadrupole MS and a TraceGOLD TG-5MS (60 m × 0.25 mm; 0.25 µm) from Thermo Scientific. The injector and detector temperatures were maintained at 250°C. The oven temperature was programmed to 40°C for 2 min, raised to 250°C at a rate of 6°C/min, and kept steady at 250°C for additional 5 min. Helium was the carrier gas at a flow rate of 1 mL/min. A sample of 1 µL of extract was injected in splitless mode.

The MS detector was operated in electron-impact ionisation (EI) mode with a mass scan range from m/z 50 to 350 at 70 eV. The compounds were identified by comparing their GC retention indices and mass spectra with published data and National Institute of Standards and Technology (NIST) mass spectra library data provided by the software of GC–MS system, and with standards. Compounds are reported as a relative percentage of the total extract by peak area.

Experimental design

A total of 19 runs were performed for optimising the three individual experimental parameters in the current model. Eight points of a factorial design with three factors at two levels ($\alpha = \pm 1.00$), six axial star points at a distance $\alpha = \pm 2.00$ from the centre point. The centre point had five replicates. The experiments were performed in a randomised order to minimise bias effect (Mendes et al., 2016). The optimisation was carried out using the response surface methodology.

All statistical analyses were made using the software Statistica version 6.0 (StatSoft, Tulsa, OK), namely, multifactor analysis of variance (ANOVA) and response surface three-dimensional (3D) plots.

Results and discussion

The optimisation of SALLE for phytochemical analysis was evaluated by the analysis of the non-volatile phenolic compounds extracted from fennel seeds. For this purpose, the relative extraction efficiency was evaluated using the total peak area obtained by the analysis of the extracts by HPLC-UV analysis.

Preliminary studies

Before using the experimental design, preliminary experiments were carried out in order to select the relevant variables for the extraction of phenolic compounds from the fennel seeds as well as the experimental range for the factors.

The first studied experimental parameter was the sample mass/solvent volume ratio. It was observed that higher ratios resulted in higher chromatographic signals. A sample mass of 0.5 g and a solvent volume of 10 mL were considered to result in a good extraction efficiency and was the best experimental conditions to achieve a good phase separation.

One of the most important parameters in SALLE is the type of salt used as salting-out agent (Valente et al., 2013a). The salts used for these studies were selected according to our previous results (Valente et al., 2013a). $\text{CH}_3\text{COONH}_4$, Na_2CO_3 , K_2CO_3 , NaCl , NaH_2PO_4 , Na_2SO_4 and $(\text{NH}_4)_2\text{SO}_4$ were tested at a concentration of 0.5 mol/L in a mixture of acetonitrile/water (1:1, v/v). The extraction of phenolic compounds was possible with all the salts tested (Figure 3). However, it was verified that the extraction efficiency of phenolics using carbonate salts was the least efficient. From the salts tested, the best result was observed for $(\text{NH}_4)_2\text{SO}_4$ being the salt chosen to the subsequent experiments. Besides, this salt has the advantage of being compatible with MS analysis.

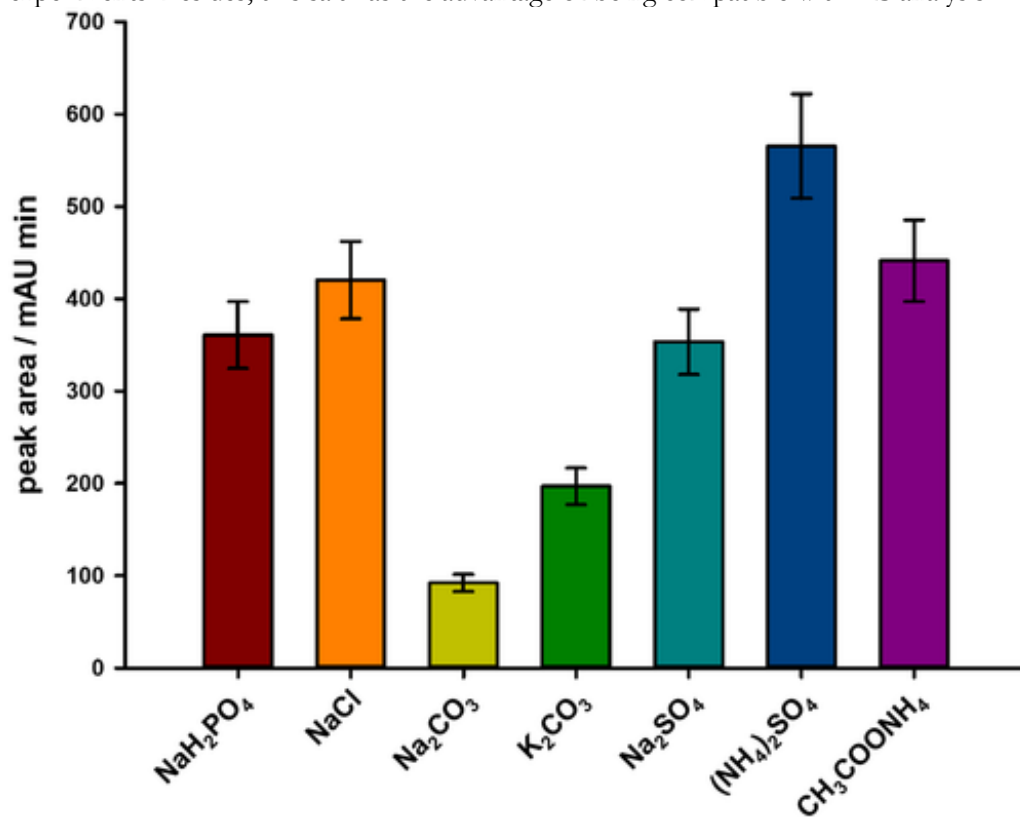


figure 3 -Effect of different salts used for SALLE in the extraction efficiency of non-volatile phenolic compounds.

Another important parameter is the solvent composition; the acetonitrile/water ratio not only affects the formation of the biphasic system, but also affects the extraction efficiency (Valente et al., 2013a). Mixtures composed of acetonitrile and water in the range 40–60% were shown to be the best conditions for phase separation.

Surface response methodology analysis

In the preliminary tests, it was possible to establish the sample's quantity, the solvent's volume and the salt nature; the other important parameters (salt concentration, solvent composition and extraction time) were optimised using a central composite design. The response values, expressed as total peak area, at different experimental combination for coded variables are listed in Table S1 in the Supporting Information. The results in the initial set of experiments (runs 1–13 in Table S1) were fitted to a first-order model, additional runs were performed to improve model adjustment (Moreira et al., 2012). The final optimised conditions are shown in Table 1.

Variable	Studied range	Optimum value
Organic solvent percentage (%)	33–67	50
Extraction time (min)	3–37	37
Salt concentration (mol/L)	0.66–2.30	0.7

Table 1. Final optimised conditions as well as tested intervals for percentage of organic solvent, extraction temperature and salt concentration.

Phenolic profile of the fennel extracts

The compounds extracted from fennel seeds by the developed methodology were identified by HPLC-MS/MS analysis by comparing retention time, UV and MS data with those of reference standards. Otherwise, the extracted phenolics were tentatively identified based on literature data. The list of identified compounds is presented in Table 2. Identification of the compounds was mainly based on the results from the negative ion mode analysis; in some cases the positive ion mode allowed the confirmation of the peaks identification.

Table 2. Identified compounds by HPLC-MS/MS in the fennel seeds extracts obtained using the developed methodology

Table 2. Identified compounds by HPLC-MS/MS in the fennel seeds extracts obtained using the developed methodology

Peak	t _R (min)	UV bands (nm)	[M-H] ⁻ m/z (% base peak)	MS/MS m/z (% base peak)	[M + H] ⁺ m/z (% base peak)	MS/MS m/z (% base peak)	Assigned identity
1	40.5	246, 297sh, 327	353(100), 421(93), 399(91), 489(40)	MS ² [353]: 191 (100)	355(100), 372(97), 731(28)	MS ² [355]: 163 (100)	5-caffeoylquinic acid (chlorogenic acid)
2	67.0	252, 300sh, 324	515(100), 583(63)	MS ² [515]: 353(100) MS ³ [515 → 353]: 191 (100)	—	—	1,5-dicaffeoylquinic acid
3	67.9	252, 300sh, 327	601(100)	MS ² [601]: 395(100), 515(45), 557(42), 439(17), 233(6) MS ³ [601 → 395]: 179(100), 233(71), 215(65), 335(49)	—	—	dicaffeoylquinic acid derivative
4	68.9	246, 300sh, 330	515(100), 583(44)	MS ² [515]: 353(100) MS ³ [515 → 353]: 191 (100), 179 (11)	539(100), 517(46), 534(34), 355(17)	MS ² [539]: 377(100), 359(48), 331(12) MS ³ [539 → 377]: 218(100), 215(66), 163(39), 185(40), 197(37), 359(35), 179(24)	3,5-dicaffeoylquinic acid
5	69.5	290sh, 320	475(100)	—	—	—	unknown
6	70.8	249, 300sh, 330	601(100), 623(23), 463(12)	MS ² [601]: 395(100), 515(85), 557(82), 439(58), 377(20), 421(12) MS ³ [601 → 395]: 233(100), 335(16)	625(100), 620(85), 465(29)	MS ² [625]: 463(100), 581(41), 419(23), 563(23), 401(17), 445(12) MS ³ [625 → 463]: 401(100), 419(77), 377(67), 257(39), 221(25), 319(21)	malonyl-1,4-O-dicaffeoylquinic acid
7	71.0	255, 305sh, 342	463(100), 601(88), 577(43)	MS ² [463]: 301(100) MS ³ [463 → 301]: 179(100), 271(51), 255(30), 151(26)	—	—	quercetin-3-O-glucoside (iso quercitrin)
8	71.6	252, 300sh, 327	515(100), 583(72), 561(61)	MS ² [515]: 353(100) MS ³ [515 → 353]: 191 (100)	—	—	1,3-dicaffeoylquinic acid
9	72.2	252, 300sh, 327	601(100), 623(41), 512(12), 644(12)	MS ² [601]: 395(100), 557(59), 439(53), 515(43), 233(22) MS ³ [601 → 395]: 233 (100)	—	—	malonyl-4,5-O-dicaffeoylquinic acid

Table 2. (Continued)

Peak	t _R (min)	UV bands (nm)	[M-H] ⁻ m/z (% base peak)	MS/MS m/z (% base peak)	[M + H] ⁺ m/z (% base peak)	MS/MS m/z (% base peak)	Assigned identity
10	73.2	255, 300sh, 354	477(100), 955(34)	MS ² [477]: 301(100)MS ³ [477 → 301]: 179(100), 151(45), 229(14), 273(11)	479(100), 501(22), 978(18)	MS ² [479]: 303(100)MS ³ [479 → 303]: 257(100), 229(73), 285(49), 247(30), 165(32)	quercetin-3-O-glucuronide (miquelianin)
11	74.4	252, 300sh, 330	477(100), 447(42), 593(49), 615(45), 545(34)	MS ² [477]: 314(100), 285(19)MS ³ [477 → 314]: 285(100), 271(46), 243(17), 300(21)	—	—	isorhamnetin-3-O-glucoside
12	75.4	264, 300, 324	—	—	163(100)	MS ² [163]: 145(100), 107(14), 135(14), 107(12)	unknown
13	76.3	264, 300sh, 345	461(100), 529(82), 507(81), 483(25)	MS ² [461]: 285(100)MS ³ [461 → 285]: 257(100), 229(75), 213(59), 199(28), 165(16)	463(100), 485(57), 947(22)	MS ² [463]: 287 (100)MS ³ [463 → 287]: 241(100), 287(66), 199(58), 135(45), 165(37), 213(40)	kaempferol-3-O-glucuronide
14	76.6	255, 309	491(100)	MS ² [491]: 315(100)MS ³ [491 → 315]: 300(100), 255(0), 273(0), 287(0)	—	—	isorhamnetin-3-O-glucuronide
15	81.87	261, 303, 363	285(100)	—	—	—	kaempferol
16	84.21	—	329(100)	MS ² [329]: 229(100), 211(55), 311(43)MS ³ [329 → 229]: 125 (100)	—	—	unknown
17	88.1	276	—	—	—	—	unknown
18	88.9	258, 300	—	—	—	—	unknown
19	91.3	243, 276, 369	353(100)	MS ² [353]: 233(100), 247(7)MS ³ [353 → 233]: 218(100), 189(49), 159(45), 165(29), 189(29)	647(100)	—	unknown

Several caffeoylquinic and dicaffeoylquinic acids were identified in the fennel seed extracts through the presence of the characteristic fragments with m/z 515 and 353 in the negative ion mode (Clifford et al., 2003) and a characteristic UV spectrum with an absorption maximum at 324 nm.

Peak at 40.54 min was assigned as 5-caffeoylquinic acid (chlorogenic acid) showing an intense molecular ion $[M-H]^-$ with m/z 353 and a MS2 fragment at m/z 191. Three isomers of dicaffeoylquinic acid (m/z 515 in negative ion mode) were identified in the extracts at 67.04, 68.90 and 71.56 min (Parejo et al., 2004). Their identification and the discrimination of the isomers was based on the literature data (Clifford et al., 2005).

Two dicaffeoylquinic acid derivatives containing a malonyl group in their structure were present in the extract. Peaks at 70.79 and 72.19 min displayed a $[M-H]^-$ ion at m/z 601 and gave characteristic fragment ions of malonyl at m/z 557 ($[M-H-44]^-$) and m/z 515 ($[M-H-86]^-$). The differences between the fragment ions intensities enabled the identification of compounds based on the literature data (Gouveia and Castilho, 2011). Hence, peaks at 70.79 min and 72.19 min were attributed to malonyl-1,4-O-dicaffeoylquinic acid (due to the presence of an intense ion at m/z 515) and malonyl-4,5-dicaffeoylquinic acid, respectively.

Several flavonoids derivatives were also identified in the obtained extracts, quercetin, isorhamnetin and kaempferol glycosides.

Peak at 71.00 min displayed a $[M-H]^-$ ion at m/z 463. The MS2 spectrum had a base peak at m/z 301, corresponding to the loss of a hexoside residue (162 Da). Fragmentation of base peak led to quercetin characteristic ions at m/z 179, 271 ($[M-H-CH_2O]^-$), 255 ($[M-H-H_2OCO]^-$) and 151. This compound was identified as quercetin-3-O-glucoside (isoquercitrin) by comparison of MSn with the literature data (Gouveia and Castilho, 2011).

A parent ion at m/z 477 was found at $t_r = 73.21$ min, attributed to quercetin-3-O-glucuronide (miquelianin), due to the presence of the characteristic MSn fragment ions of quercetin described earlier.

An m/z of 477 was also found at 74.43 min although producing a different fragmentation pattern. The MS2 fragmentation of this ion produced an intense ion peak of m/z 314 ($[M-H-163]^-$) and a MS3 resulted in characteristic isorhamnetin ions of m/z 285, 271 and 300. This peak was attributed to isorhamnetin-3-O-glucoside (Parejo et al., 2004). Another isorhamnetin derivative (isorhamnetin-3-O-glucuronide) was found at 76.62 min with a parent ion of m/z 491 which produced a MS2 intense fragment of m/z 315 and MS3 fragment ion m/z 300.

Kaempferol-3-O-glucuronide was identified at 76.25 min due to the presence of a MS2 fragment ion at m/z 285 corresponding to kaempferol, which resulted in the fragmentation of the intense parent ion m/z 461 by the loss of 176 Da.

Volatile composition of the extracts

The volatile composition of the fennel seed extracts was characterised by GC–MS analysis. The identified compounds, their retention time and relative peak area are listed in Table 3. A total of 26 compounds were identified in the extracts. The main component was estragole (58.47%) followed by limonene (17.70%), fenchone (5.15%), α -pinene (2.39%) and trans-anethole (1.61%). Other compounds were found to be minor components (< 1% of the total peak area). The profile obtained

for the SALLE extracts of fennel seeds was similar to that reported by other authors (Yu et al., 2013). Shahat et al. (2011) identified 18 compounds in the essential oil of dried seeds of *Foeniculum vulgare* var. *vulgare* and also found that estragole (57.94%), limonene (20.64%), fenchone (7.22%), α -pinene (3.61%) and trans-anethole (4.99%) were the main components of the seeds oil. The results obtained with the developed SALLE methodology were also in agreement with those obtained by other authors (Miguel et al., 2010) that verified that estragole was the main volatile compound present in dried parts of the fennel plant.

Table 3. Identified compounds by GC–MS in the fennel seeds extracts obtained using the developed methodology

Peak	t_R (min)	Assigned identity	Peak area (%) ^a
1	13.23	α -pinene	2.65
2	13.69	camphene	0.10
3	14.03	benzaldehyde	Trace ^b
4	14.39	sabinene	0.76
5	14.53	β -pinene	0.15
6	14.75	β -myrcene	0.96
7	15.24	3-carene	0.36
8	15.60	<i>o</i> -cymene	0.25
9	15.91	limonene	19.56
10	16.04	1,8-cineole (eucalyptol)	0.58
11	16.31	<i>cis</i> - β -ocimene	0.04
12	16.71	γ -terpinene	1.07
13	17.6	fenchone	5.70
14	18.47	1,5,5,6-tetramethyl-1,3- cyclohexadiene (α -pyronene)	0.47
15	19.14	camphor	0.08
16	20.4	estragole	64.61
17	20.88	cumin aldehyde	0.13
18	21.29	fenchyl acetate	0.22
19	21.54	carvone	0.09
20	21.68	<i>cis</i> -anethole	0.04
21	21.77	<i>p</i> -anisaldehyde	0.13
22	22.49	<i>trans</i> -anethole	1.77
23	27.1	germacrene D	0.09
24	27.35	2,4-bis(1,1-dimethylethyl)- phenol	0.07
25	29.79	apiole (dill.)	0.10
26	34.89	methyl palmitate	0.03

^aRelative percentage of the total peak area of the identified compounds.
^bTrace: $\leq 0.01\%$.

Final remarks

This work was used as a proof-of-concept for the SALLE aiming plant characterisation. Authors advocate its merits namely its simplicity, low-cost and versatility. SALLE was applied in the LC–MS and GC–MS analysis. Optimised conditions, obtained by crafted design of experiments, were the following: 1:1 organic/aqueous phases, 37 min time of extraction and 0.7 mol/L as the salt

((NH₄)₂SO₄) concentration. Several relevant volatile and non-volatile compounds were found ranging from phenolic compounds to quercetin derivatives and various terpenic compounds.

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Supporting information

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