

Evaluation, prediction and optimization the ultrasound-assisted extraction method using response surface methodology: antioxidant and biological properties of *Stachys parviflora* L.

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ABSTRACT

Objective(s): To optimize the extraction method using response surface methodology, extract the phenolic compounds, and identify the antioxidant and biological properties of *Stachys parviflora* L. extracts.

Materials and Methods: Maceration and ultrasound-assisted extraction (UAE) (4, 7, 10 min treatment time, 40, 70, 100 % high-intensity and 60, 80, 100 % (v v-1) methanol purity) were applied to obtain the extracts. SEM was conducted to provide the microstructure of the extracted plant. MICs (colorimetric assay), MFCs (colony diameter), total phenolic content, total flavonoid content, radical scavenging capacity and extraction efficiency were determined. HPLC analysis was applied to measure the existent phenolic compounds.

Results: A quadratic model (4 min treatment time, 74.5 % high-intensity and 74.2 % solvent purity) was suggested as the best (TPC: 20.89 mg GAE g-1 d.m., TFC: 6.22 mg QEs g-1 d.m., DPPH IC50: 21.86 µg ml-1 and EE: 113.65 mg g-1 d.m.) UAE extraction model. The optimized UAE extract was generally more effective against Gram-positive microorganisms (MIC: 10-20; MBC: 10-40 (mg ml-1)) than Gram-negative ones (MIC: 40; MBC: >40 (mg ml-1)). Moreover, it (MGI: 2.32-100 %) revealed more anti-mold activity than maceration (MGI: <28.77 %). Explosive disruption of the cell walls, therefore, enhanced extraction yield by acoustic cavitation, was elucidated using SEM. Caffeic acid, tannic acid, quercetin, trans ferulic acid and rosmarinic acid were determined as the phenolic compounds in the optimized extract.

Conclusion: RSM optimization was successfully applied for UAE from *S. parviflora*. The considerable antioxidant and biological properties were attributed to the phenolic compounds.

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Introduction

Stachys is a large genus of perennial and annual herb that belongs to Lamiaceae (previously called Labiateae). This genus, which comprises about 300 species, is one of the largest genera of Lamiaceae. Thirty-five species and four hybrids of the genus are found in Iran, including *Stachys* species (which is widely distributed throughout Iran, India and Turkey) (1). Phenolic compounds are major plant secondary metabolites with several beneficial properties (2, 3). Thus extraction methods to isolate these components with high output are of industrial

importance. Many beneficial effects have been reported for *Stachys* species, such as anti-nephritic, anti-inflammatory, radical-scavenging and anti-microbial properties (4, 5). *Stachys parviflora* L., commonly known as "Baggibuti", is found in temperate and tropical regions of Pakistan and in Iran, and is common across South Khorasan province, Iran (1).

Different extraction techniques have been employed to identify the phenolic compounds in plant materials. Conventionally, phenolic compounds are extracted by refluxing, boiling, heating and

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Soxhlet techniques. However, these methods have disadvantages such as loss of phenols due to oxidation, hydrolysis and ionization during extraction, as well as long extraction time (6). Furthermore, due to the high sensitivity of these compounds to light, heat and oxygen, efficient extraction methods are necessary to maintain the stability of phenolic compounds (7).

Advanced extraction methods have recently been developed and adopted for extracting bioactive compounds from natural plants. These include microwave-assisted extraction (MAE), ultrasound-assisted extraction (UAE), accelerated solvent extraction (ASE) and supercritical fluid extraction (SFE). In comparison with conventional extraction, UAE is an economical, simple, efficient and environmentally friendly alternative. The mechanism of UAE is attributed to cavitation efficacy, which can result in the disruption of cell walls, reductions of particle size and enhancement of mass transfer across cell membranes (6).

Response surface methodology (RSM) is a combination of statistical and mathematical techniques (8) for identifying the effect of individual process variables and efficiently locating optimum process-variable combinations for a multivariable system. As it requires less experimental data, it offers economy of experimental points (9). This method is widely employed for optimizing the extraction of phenolic compounds from different plant materials (6, 10-13). However, a survey of the literature has shown that extraction of phenolic compounds for *S. parviflora* has not yet been optimized using RSM.

On the other hand, Ahmad *et al* (2008) have isolated the flavonoid from *S. parviflora* (14). However, to our knowledge, no study has been conducted on this species in terms of its antioxidant and antimicrobial properties, whereas, some researchers have examined the antioxidant activity of other species of this plant growing in Iran (15-18). Nevertheless, other aspects such as essential-oil composition of the aerial parts of *S. parviflora* have already been studied (19).

The present study aims to optimize the phenolic compounds' extraction factors (solvent purity, ultrasound intensity and extraction time) for *S. parviflora* using RSM and employing a central composite rotatable design (CCDR) in conjunction with UAE. Scanning electron microscopy (SEM) is employed to examine the results of each extraction method. This study also aims to determine the antioxidant activity, antibacterial activity and antifungal properties of *S. parviflora*.

Materials and Methods

Plant materials

The aerial parts (the flowering stage) of *S. parviflora* were collected from South Khorasan Province (East of Iran) during April 2011. A

voucher specimen (12579) was deposited in the Herbarium of the School of Pharmacy, Mashhad University of Medical Sciences, Mashhad, Iran. The plant was dried while shaded at ambient temperature and subsequently ground using an electrical mill. Particle size (less than 149 μm) distributor (sieve No.:100) were used to classify the powder (20). The powder was stored in a dim, cool and low-humid place until the test day.

Chemicals

Caffeic acid, tannic acid, rosmarinic acid, *trans* ferulic, 2,2-diphenyl-1-picrylhydrazil (DPPH), catechin, quercetin, gentamicin and amikacin were Sigma-Aldrich Co. (St. Louis, MO, USA) products. Folin-Ciocalteu reagent, sodium carbonate, sodium nitrite, aluminium chloride, sodium hydroxide, gallic acid, BHA and methanol from Merck Co (Darmstadt, Germany), tryptone soy agar, tryptone soy broth, Mueller Hinton broth, Sabouraud dextrose agar from HiMedia (Mumbai, India) and gold conjugates from Agar scientific (Essex, UK), were used in the experiments. Redistilled water was used throughout the experiments.

Extraction of plant materials

Both maceration and ultrasound-assisted extraction (UAE) methods were employed to obtain the extracts. Extraction efficiency was determined gravimetrically.

Maceration extraction

A previously described extraction method was applied to extract the plant powder (21).

Ultrasound-assisted extraction (UAE)

UAE was applied (200 W, 26 kHz) at 40, 70 and 100 % high-intensity (UP 200Ht, Dr Hielscher GmbH, Germany). Acoustic power and intensity of ultrasonic vibrations were 0.171402 W and 21.8346 Wcm^{-2} corresponding to 100 % high-intensity. The extraction was performed in accordance with the previously described method (21), at temperature of 35 °C, natural pH (20), using methanol (60, 80 and 100 % (v v⁻¹)) as solvent for 4, 7 and 10 min time intervals. Insoluble part was removed from solvent by a centrifuge (model 2-16 KC, Sigma, Laborzentrifugen GmbH, Germany) then the solvent was evaporated at 35 °C using a rotary evaporator. Nonetheless, the soluble part was dried using a vacuum oven at 35 °C and 60 cm-Hg. The dried samples were stored in desiccator up to the test time.

Total phenolic content (TPC)

TPC was determined according to the previously described Folin-Ciocalteu method (20). Total amount of phenolic compounds were

expressed as mg gallic acid equivalent per g dried extract. All tests were carried out in triplicate.

Total flavonoid content (TFC)

TFC was measured using an aluminium chloride (AlCl₃)-based colorimetric assay which has been described before (20). Total flavonoid contents were expressed as mg quercetin equivalent (QEs) per g dried extract. All tests were carried out in triplicate.

DPPH assay

In order to determine the radical scavenging capacity (RSC) of the extracts, a slightly modified method was employed (14, 20). Briefly, methanol was added to 100-1000 µl of the extracts (0.020 g 100 ml⁻¹) to reach a total volume of 4 ml. Subsequently, DPPH solution (1000 µl, 0.012 g 100 ml⁻¹) was added and stirred for 30 sec. The solution was then kept in dim light for 115 min. Thereafter, the absorbance at λ_{max}=517 nm was checked using a spectrophotometer (Shimadzu UV-VIS 1601, Japan). Pure methanol was used as blank and BHA was considered as positive control. DPPH scavenging activity was calculated according to the following formula:

$$\% \text{DPPH scavenging} = [A_{\text{blank}} - A_{\text{sample}} / A_{\text{blank}}] \times 100 \quad \text{Eq. (A.1)}$$

where, A_{blank} is the absorbance of the control (containing all reagents except the test compound), and A_{sample} is the absorbance of the sample.

Extract concentration providing 50% DPPH inhibition (IC_{50}) was calculated by plotting the inhibition (%) against the extract concentrations (22). All tests were carried out in triplicate.

Analysis of phenolic acids

The HPLC apparatus (Model k-1500, Knauer, Germany) equipped with a HPLC pump system (model k_1001, knauer, Germany), a C18: EC 25014_6 column (250×4.6 mm, 5 µm; Nucleodur; 100_5 C18c) and an UV detector (model k_2600, Knauer, Germany) at the wavelength of 254 and 320 nm was employed. The flow rate and the injection volume were 1.0 ml min⁻¹ and 20 µl respectively; where a gradient solvent system consists of methanol-water was used. Briefly, a gradient elution was started with 100 % water and reached to 100% methanol within 20 min, while it continued for 5 min at 100% methanol. The condition was returned to 100% water during another 5 min. The internal standards include caffeic acid, tannic acid, rosmarinic acid, quercetin and *trans* ferulic were utilized (10, 50, 100 and 200 µg ml⁻¹).

Antimicrobial activities

Antibacterial activity

Four microorganisms including *Escherichia coli* (PTCC 1330, Persian Type Culture Collection or ATCC 8739, American Type Culture Collection) and *Pseudomonas aeruginosa* (PTCC 1074 or ATCC 9027) as Gram-negative bacteria as well as *Staphylococcus aureus* (PTCC 1337, 1112 or ATCC 29737, 6538 respectively), *Staphylococcus epidermidis* (PTCC 1114 or ATCC 12228 and ATCC 700576) as Gram-positive bacteria were employed. *In vitro* antibacterial screening tests were performed using 24 hr growth culture at 37 °C on tryptone soy agar. Briefly, a suspension of each bacterium was prepared in sterile normal saline from its fresh colonies. The turbidity was compared to 0.5 Mcfarland as standard to achieve 10⁸ CFU ml⁻¹. Then the suspension was adjusted to 10⁶ CFU ml⁻¹ by 10-fold serial dilution method. In order to achieve and examine the desired concentrations, 200 mg dried extracts were dissolved in 250 µl methanol as co-solvent. The volume was adjusted to 5 ml by Muller Hinton broth to achieve final concentration (40 mg ml⁻¹). Other concentrations were prepared according to the two-fold serial dilution method. Wells of a cell culture plate (96-well) were filled with both 200 µl of each concentration and 20 µl of 10⁶ CFU ml⁻¹ cell suspension.

Culture media as negative control (to show the sterility of media), gentamicin (5 µg ml⁻¹) and amikacin (4 µg ml⁻¹) as positive control for bacteria also were used. Thereafter, the plates were incubated (24 hr, 37 °C). The microorganisms were assessed by 2,3,5-triphenyltetrazolium chloride (TTC) assay as a colorimetric indicator for bacterial growth. In this method, 20 µl TTC (5 mg ml⁻¹) was added to each well and plates were incubated once again at 37 °C for almost 1 hr. The minimum inhibitory concentration (MIC) (defined as the lowest concentration of the extracts against the test microorganisms) were determined as inhibition the color change to pink-red. This change indicates the bacterial growth (20). Minimum bactericidal concentration (MBC) (defined as the lowest concentration with no bacterial growth) was determined by inoculation of 50 µl of each well on the surface of the plates containing Mueller-Hinton agar. The plates were incubated at 37 °C for 24 hr.

Anti-mold activity

The mycelial growth inhibition (MGI) of the extracts against *Penicillium expansum* (ATCC 7861) was determined according to a previously reported method (23) with some modifications. In short, the requisite quantity of each extract was added to Sabouraud dextrose agar (SDA) medium while sterilized agar medium without the extracts was used as negative control. In the aseptic conditions, mycelia discs (5 mm diameter) were

cut off from the periphery of 7-day-old-cultures and inoculated upside-down on the agar surface of the medium. Inoculated medium without the extracts was also used as positive control. Inoculated Petri dishes were incubated at 25 ± 1 °C and the observations were recorded on the 7th day. Mycelia growth inhibition (MGI) was calculated as follows:

$$\text{MGI} = (d_c - d_t) d_c^{-1} \times 100 \quad \text{Eq. (A.2)}$$

Where, d_c is fungal colony diameter in the positive control, d_t is fungal colony diameter in the plates containing different concentrations of each extract.

The minimum fungicidal concentration (MFC) was established by re-inoculated the inhibited fungal discs at MIC on SDA medium. After 7 days of inoculation, all observations were documented. Fungal growth indicated a fungistatic nature, while its absence connoted fungicidal action of the extract.

Scanning electron microscopy (SEM)

SEM was applied to determine the microstructure for untreated, classical extraction and optimized UAE treated samples. A Philips XL 30 scanning electron microscope (Eindhoven, The Netherlands) under high vacuum condition, an accelerating voltage of 20.0 kV and a working distance of 8-9 mm (24) was used. Each sample was fixed on the specimen holder with aluminum tape and then sputtered with gold in a sputter coater (BAL-TEC SCD 005, Balzers, Switzerland).

Experimental design

The highest yield of bioactive phenolic compounds from *S. parviflora* was obtained using central composite design (CCD) in order to optimize the extraction conditions. Three independent variables (time (X_1 , min), ultrasound intensity (X_2 , %) and purity of the solvent (X_3 , %)), six central points, eight factorial points and another six axis points at a distance of ± 1.682 from the center were considered to generate CCD using Design Expert statistical package (version 9.0.2.0; Stat-Ease Inc., Minneapolis, MN, USA). This experiment, lead to 20 sets of run. To avoid the degradation of temperature-sensitive compounds, the samples were kept at room temperature. Therefore, the effect of temperature was not considered in this study. Dependent responses were TPC, TFC, DPPH IC_{50} and extract efficiency (EE).

The variables were coded according to Eq. (A.3).

$$x_i = (X_i - X_0) \Delta X^{-1} \quad \text{Eq. (A.3)}$$

where, x_i is the coded value of the variable, X_i is the actual value for the independent variable, X_0 is the actual value of X_i at the center point and ΔX_i is the step change value. The independent variables in coded and natural values are tabulated in Table 1.

To predict the optimal point, the following equation as a second-order polynomial model was used.

$$Y = \beta_0 + \sum_{i=1}^k \beta_i X_i + \sum_{i=1}^k \beta_{ii} X_i^2 + \sum_{i=1}^{k-1} \sum_{j=2}^k \beta_{ij} X_i X_j \quad \text{Eq. (A.4)}$$

Where, β_0 , β_i , β_{ii} , and β_{ij} are the regression coefficients for intercept, linear, quadratic, and interaction terms, respectively while X_i and X_j are the independent variables. The larger R^2 (goodness-of-fit) and smaller P -value, the more significant corresponding coefficient (25). Correlation between the response and independent variables could be found out from 3D response surface and 2D contour plots, which are simultaneously representing the interaction of three factors on the responses and finding the location of optimum experimental variables (26). Statistical significance of the coefficients in the regression equation was checked by analysis of variance (ANOVA). The fitness of the polynomial model equation to the responses was evaluated by the coefficient of R^2 as well as the lack of fit, using F-test. For this study, 20 treatments were prepared based on the CCD, with three independent variables at five levels for each variable (Table 2).

Results

Model fitting

The most frequently used model, which was implemented for the optimization in this study, is an empirical second-order polynomial model. The accuracy of the generated model was assessed by analysis of variance (ANOVA), R^2 and lack of fit. The ANOVA results in Table 3 suggest that the model has very high F -values and very low P -value (0.0001) for all four responses.

In addition, high R^2 values and insignificance lack of fit ($P > 0.05$) indicate that the implementation of the quadratic model was highly significant for the obtained data. This suggests that the model can describe the relationship between the extraction conditions and responses during UAE. The fitted quadratic models for TPC, TFC, DPPH IC_{50} and EE all in coded form, are given in Eqs. A.5 to A.8, respectively. The significance of each coefficient was determined using T tests and P -value (Table 3).

Table 1. Independent variables and coded values employed for optimization of the extraction procedure

Real independent variables	Coded independent variables				
	Corner points		Centre points	Star points	
	-1.682	-1	0	+1	+1.682
Time treatment (min), X_1	1.96	4	7	10	12.04
UAE treatment (%), X_2	22.36	38	70	84	99.64
Purity of the solvent (%), X_3	50.14	60	80	89	98.86

Table 2. Coded central composite design (CCD) matrices with observed and predicted values for dependent responses of total phenolic content (TPC), total flavonoid content (TFC), inhibition capacity (IC_{50}) and extraction efficiency (EE)

Test runs	Independent factors levels			Dependent responses			
	Time treatment (min)	UAE treatment (%)	Purity of the solvent (%)	TPC (mg GAE g ⁻¹ dry matter)	TFC (mg QEs g ⁻¹ dry matter)	DPPH IC_{50} (μ g ml ⁻¹)	EE (mg g ⁻¹ dry extract)
13	4.00 (-1)	38.00 (-1)	60.00 (-1)	19.45	6.15	23.28	99.32
6	4.00 (-1)	38.00 (-1)	89.00 (1)	19.94	6.12	24.25	99.48
3	4.00 (-1)	84.00 (1)	60.00 (-1)	20.48	6.23	22.99	112.64
15	4.00 (-1)	84.00 (1)	89.00 (1)	20.92	6.19	23.80	108.73
17	10.00 (1)	38.00 (-1)	60.00 (-1)	17.95	5.93	30.85	93.62
10	10.00 (1)	38.00 (-1)	89.00 (1)	18.44	5.92	31.95	97.56
5	10.00 (1)	84.00 (1)	60.00 (-1)	18.98	6.01	30.31	98.43
1	10.00 (1)	84.00 (1)	89.00 (1)	19.48	5.97	31.37	96.15
12	12.05 (+1.682)	61.00 (0)	74.50 (0)	17.96	5.80	30.78	90.34
7	1.95 (-1.682)	61.00 (0)	74.50 (0)	20.40	6.15	18.20	103.17
4	7.00 (0)	99.68 (+1.682)	74.50 (0)	20.12	6.22	28.18	105.46
9	7.00 (0)	22.32 (-1.682)	74.50 (0)	18.45	6.12	28.87	96.16
11	7.00 (0)	61.00 (0)	98.89 (+1.682)	20.44	6.05	29.60	106.99
2	7.00 (0)	61.00 (0)	50.11 (-1.682)	19.69	6.08	28.01	106.70
8	7.00 (0)	61.00 (0)	74.50 (0)	20.47	6.16	25.96	113.32
14	7.00 (0)	61.00 (0)	74.50 (0)	20.47	6.17	25.94	114.12
16	7.00 (0)	61.00 (0)	74.50 (0)	20.48	6.17	25.90	113.98
18	7.00 (0)	61.00 (0)	74.50 (0)	20.48	6.16	25.92	114.45
19	7.00 (0)	61.00 (0)	74.50 (0)	20.44	6.16	25.90	114.14
20	7.00 (0)	61.00 (0)	74.50 (0)	20.45	6.17	25.90	114.34

All the observed and predicted values were identical. TPC ranged from 17.95 to 20.92 mg GAE g⁻¹ dry matter; TFC from 5.80 to 6.23 mg QEs g⁻¹ dry matter; extraction efficiency from 90.34 to 114.45 mg g⁻¹ dry extract; and radical scavenging capacity from 18.20 to 311.95 μ g ml⁻¹ (Table 2).

The results of the 20 runs related to CCD, which include the coded matrices for design conditions, observed responses and corresponding predicted values, are tabulated in Table 2.

Optimization of process variables and verification of the model

The reliability of the fitted model for all four responses (based on the ANOVA results) was supported as the maximal points of the responses were inside the experimental region. As per the response surface plots, a simultaneous optimization for all responses was carried out. The optimal conditions were treatment time of 4 min, 74.5%

high-intensity and solvent purity of 74.2%, which give estimated maximal values of 20.89 mg GAE g⁻¹ dry matter for TPC, 6.22 mg QEs g⁻¹ dry matter for TFC, 21.86 μ g ml⁻¹ for DPPH IC_{50} and 113.65 mg g⁻¹ dry extract for EE.

SEM

Figure 2 shows the microstructure of the optimized UAE extract (4 min treatment time, 74.5 % high-intensity and 74.2% solvent purity), maceration extract and untreated sample of *S. parviflora*. SEM images clearly show that the untreated sample contained no fractured cells (Figure 2 (a)), whereas both treated samples underwent fractural changes. These changes were more obvious for optimized UAE extraction than the sample obtained using the maceration method (Figure 2 (b, c)): the maceration sample contained shrunken cells, while the cells in the optimized UAE sample were shredded.

$$Y_{\text{TPC}} = 20.46 - 0.74X_1 + 0.5X_2 + 0.23X_3 - 0.45X_1^2 - 0.42X_2^2 - 0.14X_3^2 \quad \text{Eq. (A.5)}$$

$$Y_{\text{TFC}} = 6.16 - 0.11X_1 + 0.033X_2 - 0.012X_3 - 0.067X_1^2 - 0.035X_3^2 \quad \text{Eq. (A.6)}$$

$$Y_{\text{DPPH}} = 25.92 + 3.76X_1 - 0.22X_2 + 0.48X_3 - 0.51X_1^2 + 0.92X_2^2 + 1.02X_3^2 - 0.047X_1.X_2 + 0.048X_1.X_3 \quad \text{Eq. (A.7)}$$

$$Y_{\text{EE}} = 114.06 - 4.1X_1 + 3.05X_2 - 6.12X_1^2 - 4.68X_2^2 - 2.54X_3^2 - 2.4X_1.X_2 + 0.68X_1.X_3 - 1.29X_2.X_3 \quad \text{Eq. (A.8)}$$

Table 3. Analysis of variance of the quadratic regression model fitted for total phenolic content (TPC), total flavonoid content (TFC), inhibition capacity (IC₅₀) and extraction efficiency (EE)

Model	TPC (mg GAE g ⁻¹ dry matter) ^a				TFC (mg QEs g ⁻¹ dry matter) ^b			
	Estimate	Std Err	P-value	F-value	Estimate	Std Err	P-value	F-value
	20.46	0.0084	<0.0001	4336.01	6.16	0.0027	<0.0001	611.30
X ₁	-0.74	0.0055	<0.0001	17327.23	-0.11	0.0018	<0.0001	3392.48
X ₂	0.50	0.0055	<0.0001	8151.07	0.03	0.0018	<0.0001	324.74
X ₃	0.23	0.0055	<0.0001	1738.54	-0.01	0.0018	<0.0001	46.98
X ₁ X ₂	0.01	0.0073	0.3284	1.06	0.00	0.0023	0.3181	1.10
X ₁ X ₃	0.01	0.0073	0.3284	1.06	0.00	0.0023	0.3181	1.10
X ₂ X ₃	-0.01	0.0073	0.5089	0.47	-0.01	0.0023	0.0619	4.42
X ₁ ²	-0.45	0.0054	<0.0001	6953.01	-0.07	0.0017	<0.0001	1428.48
X ₂ ²	-0.42	0.0054	<0.0001	5860.78	0.00	0.0017	0.2924	1.24
X ₃ ²	-0.14	0.0054	<0.0001	667.01	-0.04	0.0017	<0.0001	393.56
	DPPH IC ₅₀ (μg ml ⁻¹) ^c				E.E (mg g ⁻¹ dry extract) ^d			
	P-value lack of fit>0.05				P-value lack of fit>0.05			
Model	Estimate	Std Err	P-value	F-value	Estimate	Std Err	P-value	F-value
	25.92	0.0136	<0.0001	22647.62	114.06	0.23	<0.0001	413.78
X ₁	3.76	0.0090	<0.0001	172175.5	-4.10	0.16	<0.0001	692.37
X ₂	-0.22	0.0090	<0.0001	596.5027	3.05	0.16	<0.0001	382.45
X ₃	0.48	0.0090	<0.0001	2860.166	-0.12	0.16	0.4685	0.57
X ₁ X ₂	-0.05	0.0118	0.0025	16.1167	-2.40	0.20	<0.0001	138.56
X ₁ X ₃	0.05	0.0118	0.0025	16.1167	0.68	0.20	0.0077	11.04
X ₂ X ₃	-0.03	0.0118	0.0607	4.4644	-1.29	0.20	<0.0001	39.92
X ₁ ²	-0.51	0.0088	<0.0001	3314.771	-6.12	0.15	<0.0001	1621.69
X ₂ ²	0.92	0.0088	<0.0001	10884.75	-4.68	0.15	<0.0001	949.46
X ₃ ²	1.02	0.0088	<0.0001	13355.27	-2.54	0.15	<0.0001	280.19

^a The R square obtained in fit statistics for the response model of total phenolic content (TPC) was 0.9997

^b The R square obtained in fit statistics for the response model of total flavonoid content (TFC) was 0.9982

^c The R square obtained in fit statistics for the response model of 2,2-diphenyl-1-picrylhydrazil (DPPH) was 1.000

^d The R square obtained in fit statistics for the response model of extraction efficiency (EE) was 0.9973

HPLC

The only study indicating the HPLC analysis of *S. parviflora* was reported by Acikara *et al* (2013) (27). They revealed that chlorogenic acid was just the phenolic acid in both the aerial part (444.77±2.78 μg mg⁻¹) and root (509.96±6.64 μg mg⁻¹). The HPLC

chromatogram from *S. parviflora* optimized UAE extract indicated that this extract abundant caffeic acid (3.88 mg g⁻¹) followed by tannic acid (1.24 mg g⁻¹), quercetin (0.49 mg g⁻¹), *trans* ferulic acid (0.39 mg g⁻¹) and rosmarinic acid (trace) (Figure 3).

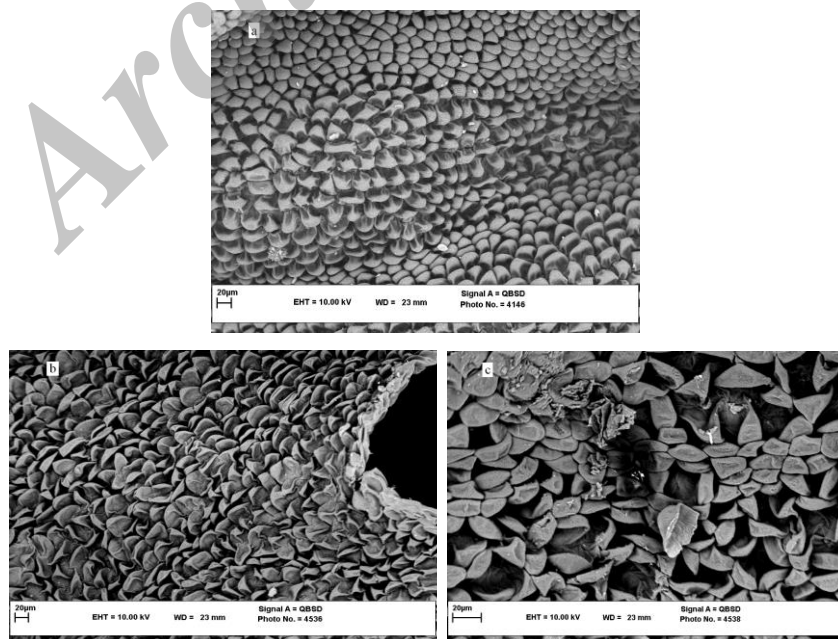


Figure 2. Scanning electron micrographs (SEM) from *Stachys parviflora* surface, (a) untreated sample, (b) maceration, (c) optimized ultrasound-assisted extraction (UAE) (4 min treatment time, 74.5 % high-intensity and 74.2 % solvent purity)

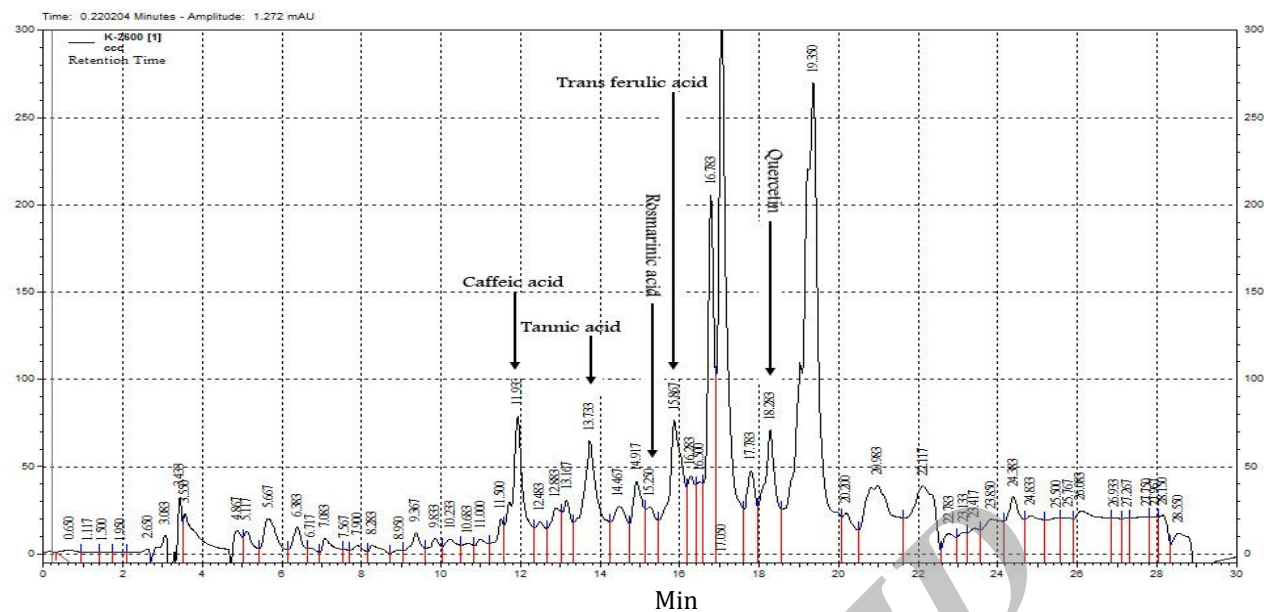


Figure 3. High performance liquid chromatography (HPLC) chromatogram from *Stachys parviflora* optimized UAE extract

Antimicrobial activities

Antibacterial activity

Many studies have demonstrated the antibacterial activity of the *Stachys* genus (28- 30). To our knowledge however, there is no study focusing on the antibacterial activity of *S. parviflora* specifically. Skaltsa *et al* (2003) declared that sesquiterpene hydrocarbons were the main group of constituents of all studied *Stachys* species from Greece responsible for antimicrobial activity. They meanwhile found out a better activity there is in studied species against bacteria than fungi (31). Likewise, the composition and antibacterial activity of *Stachys inflata* Benth. essential oil were determined by Ebrahimabadi *et al* (2010) (30). Linalool, α -terpineol, spathulenol and (2E)-hexenal (in descending order of quantity) were determined as the major components while the two earlier compounds appeared as the most considerable antibacterial

agent. Investigation of the essential oil composition of *S. parviflora* (5) revealed that three oxygenated sesquiterpenes – muurolol $\langle\text{epi-}\alpha\rangle$ (32, 33), caryophyllene $\langle z\rangle$-and caryophyllene oxide (34) – have antimicrobial activities. Phytochemical analyses of *Stachys* species have confirmed the occurrence of diterpenes, phenyl ethanoid glycosides, flavonoids and saponines. Flavonoids are a class of phytochemicals that may be responsible for antimicrobial activity (29, 35, 36).

Table 4 shows the MICs and MBCs of *S. parviflora* on the tested bacterial strains. The MIC values range from 10 to 40 mg ml⁻¹ for both maceration and optimized UAE, while the MBC values range from 20 to >40 mg ml⁻¹ for maceration and 10 to >40 mg ml⁻¹ for optimized UAE. The findings indicate that *S. parviflora* was generally more effective against Gram-positive microorganisms than Gram-negative ones.

Table 4. Antibacterial activity of *Stachys parviflora* extracts

Microorganism	MIC ^a (mg ml ⁻¹) extract		MBC ^b (mg ml ⁻¹) extract		Positive control	MIC (mg ml ⁻¹)
	Maceration	Optimized UAE	Maceration	Optimized UAE		
<i>Escherichia coli</i> (PTCC 1330 or ATCC 8739)	20	40	>40	>40	Gentamicin	0.005
<i>Staphylococcus aureus</i> (PTCC 1112 or ATCC 6538)	20	20	>40	40		
<i>Staphylococcus aureus</i> (PTCC 1337 or ATCC 29737)	10	20	40	20		
<i>Staphylococcus epidermidis</i> (PTCC 1114 or ATCC 12228)	20	10	20	10	Amikacin	0.004
<i>Staphylococcus epidermidis</i> (ATCC 700576)	20	10	40	10		
<i>Pseudomonas aeruginosa</i> (PTCC 1074 or ATCC 9027)	40	40	40	40		

^a MIC: Minimum inhibitory concentration, ^b MBC: Minimum bactericidal concentration

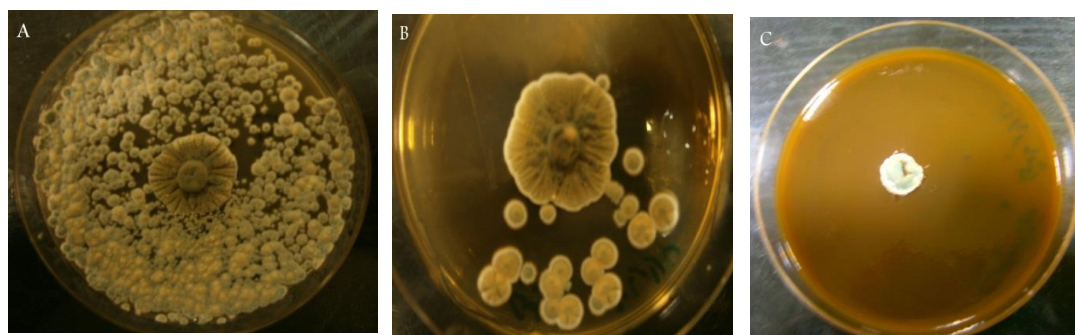


Figure 4. Anti-mold activity of *Stachys parviflora*. blank (A), extract (optimized UAE) concentration less than minimum inhibitory concentration (MIC) (B), extract (optimized UAE) concentration at minimum inhibitory concentration (MIC) (C)

Anti-mold activity

Table 5 summarizes the anti-mold activity of both maceration and optimized UAE extracts. According to the results, the maceration extract at the extract concentration of 0.305 to 5 mg ml⁻¹, did not inhibit the mycelia growth, whereas optimized UAE extract showed an MGI% of 2.32 to 30.78 at the same range of concentration. It also showed that the MGI% of maceration extract was only 28.77%, in contrast to 100% for optimized UAE at the final concentration of 160 mg ml⁻¹. These results indicate that optimized UAE extract of *S. parviflora* possesses the stronger activity against *P. expansum* than maceration extract.

Discussion

The effects of three independent process variables – treatment time (X_1 : 1.95-12.04 min), UAE treatment (X_2 : 22.31-99.68%) and purity of the solvent (X_3 : 50.11-98.88%) – were investigated during ultrasonic extraction of *S. parviflora* aerial parts. A quadratic model has been suggested as the best model for all independent variables (Table 6).

Figure 1 represents the relationship between each pair of process variables in each response. As shown in Table 3, the term of the extraction time (X_1) was the most effective factor on TPC, while the quadratic term of the solvent purity (X_3)

had the smallest effect. The results indicated that none of the interactions of the treatment time and UAE (X_2), the treatment time and the solvent purity and UAE and the solvent purity had a significant effect on TPC ($P>0.05$). The fitness of the predicted TPC model was defined by its R^2 (0.9997) and insignificant lack of fit (0.2093), as shown in Table 3. Figure 1A shows the 3D response surface and the contour plots.

The most effective factor in extracting TFC was extraction time (X_1), followed by its quadratic term, while the interaction terms between all parameters – the extraction time, UAE (X_2) and the solvent purity (X_3) – as well as the quadratic term of UAE had no significant effect on TFC ($P>0.05$). Furthermore, the solvent purity had the lowest effect on the flavonoid compounds extracted from *S. parviflora* (Table 3). The coefficient of R^2 of the predicted TFC model was 0.9982, and P -value for the lack of fit was 0.2294.

Three-dimensional surface and contour plots shown in Figure 1B were generated according to Eq. A.6. Figure 1B (a) shows the effects of for TPC as a function of the process variables. TPC was greatly increased with an increase in the extraction time, reaching to 20.47 mg GAE g⁻¹ dry matter, with the lowest residual value at $t = 7$ min. Thereafter, it started to decrease until the 10th min. Figure 1A (b) confirms that the solvent purity

Table 5. Mycelial growth inhibition (MGI) of different *Stachys parviflora* extracts against *Penicillium expansum*

Extract concentration (mg/ml)	Diameter (mm)				Control zone	Mycelia growth inhibition	
	Maceration		Optimized UAE			MGI (%)	
	Test zone	Result zone	Test zone	Result zone		Maceration	Optimized UAE
0.305	47.37	-	28.69	0.68	29.37	Tr	2.32 ^s
0.612	40.53	-	27.61	1.76		Tr	5.99 ^s
1.25	37.14	-	25.13	4.24		Tr	14.44 ^s
2.5	33.65	-	23.04	6.32		Tr	21.52 ^s
5	31.42	-	20.32	9.04		Tr	30.78 ^s
10	28.33	1.04	12.47	16.89		3.54 ^s	57.51 ^s
20	27.32	2.05	8.23	21.13		6.98 ^s	71.94 ^s
40	25.28	4.09	4.27	25.10		13.93 ^s	85.46 ^s
80	23.96	5.41	1.36	28.01		18.42 ^s	95.37 ^s
160	20.92	8.45	0.00	29.37		28.77 ^s	100.00 ^s

^s Fungistatic; tr: Trace

Table 6. Choosing the best model for total phenolic content (TPC), total flavonoid content (TFC), inhibition capacity (IC₅₀) and extraction efficiency (EE)

Source	TPC (mg GAE g ⁻¹ dry matter)					TFC (mg QEs g ⁻¹ dry matter)			
	DF	P-value	Lack of fit (P-value)	Adjusted R ²	Predicted R ²	P-value	Lack of fit (P-value)	Adjusted R ²	Predicted R ²
Linear	3	0.0002	<0.0001	0.6405	0.5695	0.0003	<0.0001	0.6235	0.5034
2fi	3	1.0000	<0.0001	0.5576	0.2720	0.9970	<0.0001	0.5384	0.3095
Quadratic	3	<0.0001	0.2093	0.9995	0.9985	<0.0001	0.2294	0.9966	0.9896
Cubic	4	0.1120	0.6207	0.9997	0.9989	0.1149	0.7685	0.9981	0.9966

Source	DPPH IC ₅₀ (µg ml ⁻¹)					EE (mg g ⁻¹ dry extract)			
	DF	P-value	Lack of fit (P-value)	Adjusted R ²	Predicted R ²	P-value	Lack of fit (P-value)	Adjusted R ²	Predicted R ²
Linear	3	<0.0001	<0.0001	0.8357	0.7666	0.1330	<0.0001	0.1545	-0.0174
2fi	3	0.9994	<0.0001	0.7980	0.7235	0.8020	<0.0001	0.0336	-0.6234
Quadratic	3	<0.0001	0.1687	0.9999	0.9997	<0.0001	0.1154	0.9949	0.9829
Cubic	4	0.0872	0.5902	1.0000	0.9998	0.0470	0.7983	0.9979	0.9970

had no significant effect on TPC. The linear increase in TPC shown in Figure 1A (c) also demonstrates that UAE had a significant effect on TPC, whereas the effect of the solvent purity was not significant.

The most effective factor in extracting TFC was extraction time (X₁), followed by its quadratic term, while the interaction terms between all parameters – the extraction time, UAE (X₂) and the solvent purity (X₃) – as well as the quadratic term of UAE had no significant effect on TFC ($P > 0.05$). Furthermore, the solvent purity had the lowest effect on the flavonoid compounds extracted from *S. parviflora* (Table 3). The coefficient of R² of the predicted TFC model was 0.9982, and P-value for the lack of fit was 0.2294. Three-dimensional surface and contour plots shown in Figure 1B were generated according to Eq. A.6. Figure 1B (a) shows the effects of the extraction time and UAE on TFC.

Total flavonoid content of the extracts increased linearly with UAE, whereas the extraction time had the opposite effect on TFC; and the highest TFC was obtained at $t = 4$ min. A similar linear decrease in TFC was observed as shown in Figure 1B (b), where TFC decreased with the solvent purity. Figure 1B (c) shows the positive effects of UAE and negative effects of the solvent purity on TFC.

Except the interaction effect of the solvent purity (X₃) and UAE (X₂) ($P > 0.05$), all other parameters effected the free radical scavenging activity of the extracts significantly ($P < 0.05$).

Meanwhile, the interaction effects of the extraction time (X₁) and the solvent purity as well as the extraction time and UAE were in the lowest when compared with other parameters. R² for DPPH scavenging capacity was obtained 1.000, and the lack of fit was not significant (0.1687). As shown in Figs 1C (a) and (b), which give 3D response surface and contour plot for DPPH scavenging capacity, the extraction time had a

linear effect on IC₅₀, and an increase in IC₅₀ corresponded to lower antiradical scavenging capacity. These figures also showed the opposite effect of UAE and linear effect of the solvent purity on DPPH scavenging capacity. Furthermore, as Figure 1C (c) shows, the interaction effects of the solvent purity and UAE were extremely low and insignificant ($P > 0.05$).

All terms except the linear term of the solvent purity (X₃) ($P > 0.05$) had significant effect on EE ($P < 0.05$). Nevertheless, the quadratic term of the extraction time (X₁) was the most significant parameter among all effective parameters. R² was 0.9973 for EE, and the lack of fit was insignificant (0.1154). Figure 1D (a), 3D response surface and contour plot for extraction efficiency, shows the linear effect of the extraction time, which led to a reduction in EE; the highest extraction yield was obtained at $t = 7$ min. In contrast, the linear effect of UAE positively agreed with an increase in EE; the highest extraction yield occurred at 61% high-intensity. However, as Figure 1D (b) shows, the solvent purity had an insignificant effect on EE; the interaction effects of the solvent purity and UAE are shown in Figure 1D (c). To determine the accuracy and reliability of the predicted model, also inspection the deviation between actual and estimated values under the suggested optimal conditions, a verification experiment was carried out. The test was conducted at a treatment time of 4 min, 74% high-intensity and the solvent purity of 74%. The observed response values were 20.888 mg GAE g⁻¹ dry matter for TPC, 6.226 mg QEs g⁻¹ dry matter for TFC, IC₅₀ of 21.849 µg ml⁻¹ for DPPH and 113.667 mg g⁻¹ dry extract for EE. No significant differences were perceived between the observed and estimated values, confirming that the fitted model for each response was valid and reliable for simulating the UAE extraction of phenolic and flavonoid compounds from *S. parviflora*.

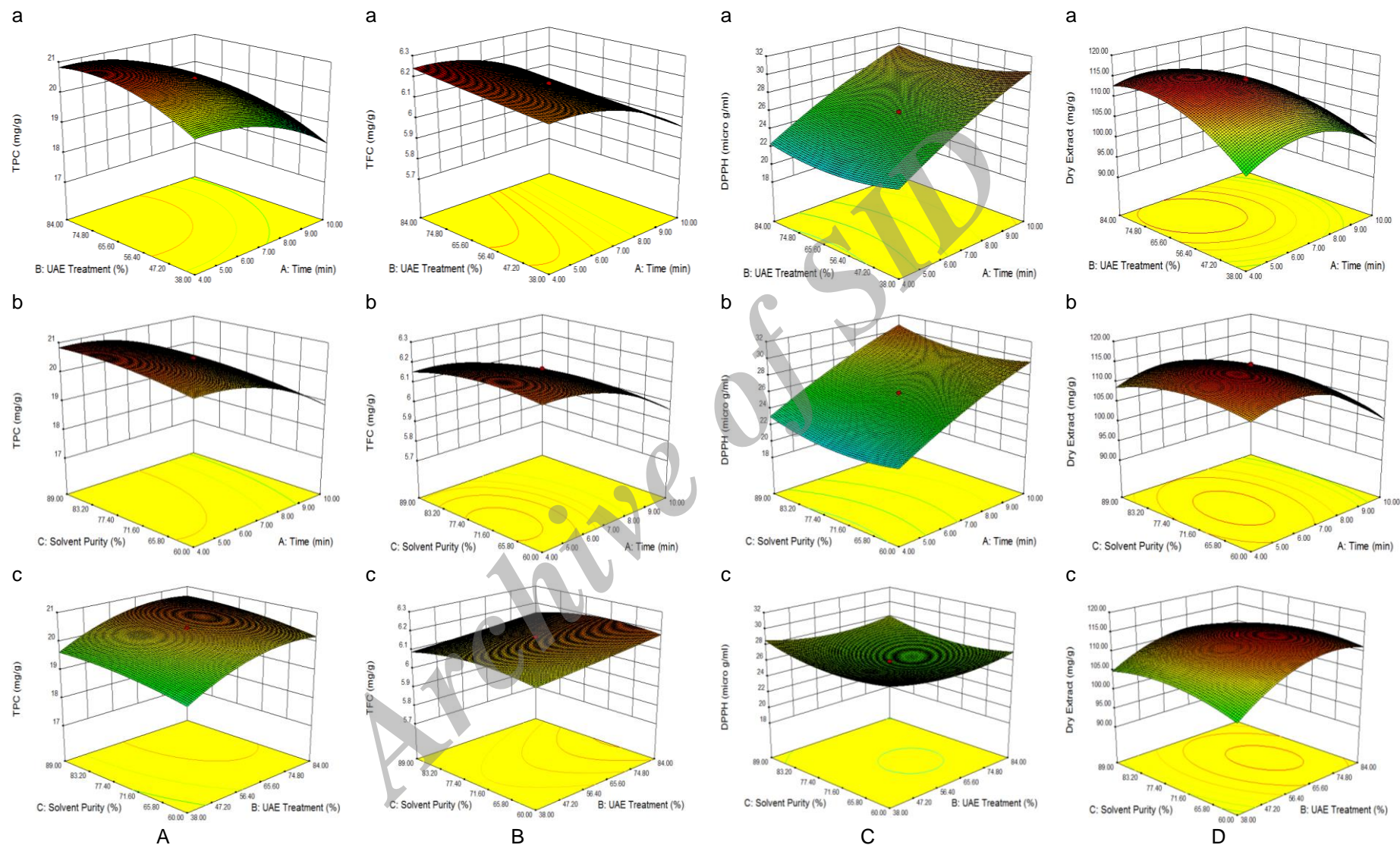


Figure 1. Response surface plots and contour plots show the effect of (a) ultrasound-assisted extraction (UAE) and time, (b) time and the purity of the solvent, and (c) UAE and the purity of the solvent on response of total phenolic content (TPC) (A), total flavonoid content (TFC) (B), radical scavenging capacity (RSC) (C), and extraction efficiency (EE) (D) of *Stachys parviflor* extracts

Based on the SEM images, sonication definitely had an important role in the explosive disruption of the physical structure of vegetal cell walls, and enhanced the extraction yield by the acoustic cavitation (37). Similar results have reported the effect of ultrasonic vibration on the physical structure of *Achillea biebersteinii* Afan. (21) and *Euonymus alatus* (Thunb.) Sieb. (38). The hydration process followed by the swelling process, also mass transfer of the soluble compounds from the plant matrices to the solvent by osmotic and diffusion processes are two stages of the classical extraction process (21). In contrast, the UAE method enhances the swelling and softening process of cell walls via the hydration of pectinous material from the middle lamella. Consequently, it may lead to the breakup of the vegetal tissue during sonication (39) which means, UAE is much faster and more efficient than the classical extraction method. SEM images revealed that the efficiency of UAE for the extraction of phenolic compounds, and consequently the enhancement of their antioxidant and antimicrobial activities could reasonably be considered superior to that of the classical extraction method.

As per the microbial tests, *S. parviflora* extracts was generally more effective against Gram-positive microorganisms than Gram-negative ones. With respect to the antimicrobial properties of phenolics (40, 41) and flavonoids (28), the observations could be partially attributed to the higher phenol and flavonoid content of the obtained extracts. A similar effect of the plant extracts on *S. epidermidis* and *S. aureus*, noted by Salarbashi et al (2014), is also remarkable for *S. parviflora* optimized UAE extract (21).

High anti-mold activity of the optimized UAE extract could be attributed to its higher extraction efficiency when compared with the maceration. A number of studies were conducted to prove the lower MIC of different bacteria is related to the higher extraction of phenolic and flavonoid compounds (20, 21, 26). Therefore, the higher MGI% of UAE can be a consequence of higher concentration of these compounds. Hence, at a concentration of 160 mg ml⁻¹, optimized UAE showed a mold-static effect on *P. expansum*; in contrast, no moldicide effect on *P. expansum* was achieved, based on the growing the mold on all discs after incubation for a week.

Conclusion

RSM was successfully optimized the ultrasound extraction from *S. parviflora*, as a quadratic model was suggested as the best model for the extraction conditions. The obtained extracts were found to be generally more effective against Gram-positive microorganisms than Gram-negative ones; while they were actively

against *P. expansum*. Meanwhile, the optimized UAE appeared more active against the test microorganisms than the classical method. SEM micrographs confirmed the efficiency of UAE in the extraction of active components. Caffeic acid was found abundantly than other phenolic acids in the *S. parviflora* extract and may consider as a responsible antimicrobial agent.

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