



Article

Phenolic Compounds from Irradiated Olive Wastes: Optimization of the Heat-Assisted Extraction Using Response Surface Methodology

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Abstract: Olive pomace, an environmentally detrimental residue generated during olive oil extraction, contains bioactive compounds in demand by the food industry. To valorize this waste product a suitable yield for the extraction process is required. Heat-assisted extraction of bioactive compounds from olive pomace was optimized by a circumscribed central composite design and response surface methodology. Our previous studies indicated that irradiation could improve 2.4-fold the extractability of the main phenolic compounds from olive pomace. The effect of extraction time, temperature and solvent concentration on the yield of polyphenols from irradiated olive pomace at 5 kGy was tested. Hydroxytyrosol-1- β -glucoside, hydroxytyrosol, tyrosol and caffeic acid were quantified by High Performance Liquid Chromatography to calculate the total polyphenol content. The optimal general conditions by RSM modeling were extraction time of 120 min, temperature of 85 °C, and 76% of ethanol in water. Using these selected conditions, 19.04 \pm 1.50 mg/g dry weight, 148.88 \pm 8.73 mg/g extract of total polyphenols were obtained, representing a yield of 13.7%, which was consistent with the value predicted by the model. This work demonstrated the potential of residues from the olive oil industry as a suitable alternative to obtain compounds that could be used as ingredients for the food industry.

Keywords: olive pomace; bioactive compounds; ionizing radiation; heat-assisted extraction; extraction optimization

1. Introduction

The olive industry is one of the most important activities in the Mediterranean region countries, which produce 95% of the world's olive oil. This industry generates large amounts of wastes, where many potentially interesting compounds remain. Olive pomace is rich in bioactive compounds that can be divided in several classes: simple phenols (e.g., tyrosol and hydroxytyrosol) and their derivatives, benzoic and cinnamic acid derivatives (e.g., gallic acid, syringic acid, caffeic acid, p-coumaric acid, verbascoside), flavonoids (e.g., apigenin, luteolin, rutin) and secoiridoids (e.g., oleuropein and oleuropein aglycone isomers) [1–3]. Although olive oil wastes can have a negative impact to the environment when discharged without treatment, the recovery of these compounds increases the sustainability of the sector, obtaining high added-value products with low production costs and reducing



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the environmental risk. In fact, society's growing demand for healthier foods challenges the scientific community to search and develop new ingredients [4]. Nowadays, with the COVID-19 pandemic situation, more efforts have to be done to develop sustainable and modern food systems. Although the lack of information in correlating the consumption of bioactive ingredients with the prevention or recovery from COVID-19 disease, it became even more imperative to have a healthier immune system which can be achieved with the supplementation of consumers' diets with vitamins, tannins, polyphenols, flavonoids, bioactive lipids and herbs [5]. In this respect, the compounds obtained from olive pomaces could be both a suitable alternative in the food industry to the use of synthetic antioxidants in order to improve the quality of foods, as well as employed in the formulation of functional foods [6].

Numerous studies have reported the extraction of bioactive compounds from olive pomace [1,3,7–9]. The extraction of these compounds can be performed by using conventional or emerging technologies, which can be advantageous since they take reduced extraction time, accelerate heat and mass transfer, increase the extraction selectivity and purity, and use safer solvents comparing to the conventional ones [10,11]. Alu'datt et al. [12] achieved the highest phenolic content extracted from olive pomace using methanol as solvent and performing the extraction for 12 h at 70 °C. Also, Vitali Čepo et al. [13] tried to optimize the extraction of hydroxytyrosol, tyrosol and oleuropein from olive pomace using simple solvent extraction and different conditions (solvent, pH, temperature and duration of extraction). The results demonstrated that the optimum conditions for phenols were 120 min at 70 °C using 60% ethanol as extraction solvent, at solvent-to-sample ratio 5:1 (v/w). Under these conditions, high recoveries of oleuropein, tyrosol and hydroxytyrosol were obtained averaging 115.14 \pm 0.19, 86.05 \pm 0.34 and 81.80 \pm 0.41 mg/kg of fresh olive pomace, respectively. More recently, also Böhmer-Maas et al. [14] studied the extraction optimization of individual compounds from olive pomace, observing that 80% methanol, 45 °C and 180 min were the optimal conditions to recover 154.90 mg/kg dry weight of hydroxytyrosol, 1115.40 mg/kg dry weight of tyrosol, and 153.20 mg/kg dry weight of syringic acid. Zuorro [15] evaluated the effects of temperature, extraction time, solvent composition and liquid-to-solid ratio on the yield of phenolics extraction from olive pomace, showing that temperature was the most influential factor.

Ionizing radiation, a clean and environmentally friendly technology, has proven to be capable of improving phenolic extraction and antioxidant activity on industrial wastewater [16], fresh fruits such as cherry tomatoes [17], raspberries [18,19] and strawberries [20], and dried medicinal plants [21]. More recently, the potential of gamma radiation as an enhancer for phenolic compounds extraction and antioxidant capacity has been shown, with the application of low doses (5 kGy), enough to increase the extractability of the main phenolic compounds from olive pomace by 2.4-fold compared to non-irradiated samples [1].

The current study aims to explore the extraction of some phenolic compounds from olive pomace by heat-assisted extraction (HAE) using a circumscribed central composite design (CCCD) testing different conditions, namely the percentage of ethanol (0–100%), extraction times (20–120 min) and temperature (25–85 °C). The olive pomace samples used in this work were the irradiated ones for which the best results in a previous work of the authors were obtained [1]. The characterization of the individual phenolic compounds was performed, summarized and modeled by Response Surface Methodology (RSM) in order to understand the combined effects of operating variables and to maximize the responses analyzed. RSM is an efficient statistical method for optimizing processes and was originally described by Box & Wilson [22] as a statistical and mathematical tool. RSM allows a more efficient and easier presentation and interpretation of experiments compared to other methodologies. To the best of our knowledge, the optimization of tyrosol, hydroxytyrosol-1- β -glucoside and caffeic acid recovery by RSM has never been reported.

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2. Materials and Methods

2.1. Standards and Reagents

HPLC-grade acetonitrile was obtained from Fisher Scientific (Lisbon, Portugal). Ethanol and formic acid were acquired from Sigma-Aldrich (St. Louis, MO, USA) and Honeywell (Charlotte, NC, USA), respectively. Caffeic acid (\geq 99%) was purchased from Extrasynthese (Genay, France), whereas hydroxytyrosol (\geq 99%) and tyrosol (\geq 98%) were obtained from Applichem (Darmstadt, Germany) and TCI (Tokyo, Japan), respectively. Water was treated in a Milli-Q water purification system (Merck Millipore, Burlington, MA, USA).

2.2. Olive Pomace Samples

Samples used in this work were olive pomaces collected in 2018 from UCASUL (União de Cooperativas Agrícolas do Sul, located in the Alentejo region, Alvito, Portugal) and further submitted to an irradiation-assisted extraction [1].

2.3. Irradiation Experiments

Irradiation was carried out in a Co-60 semi-industrial unit (with an activity of 187 kCi in April 2018) located at Technological Unit of Radiosterilization (UTR-IST), University of Lisbon (Portugal). Sealed bags (10×7 cm) containing 30 g of extracted olive pomace were irradiated at room temperature at 4.9 kGy using a dose rate of 13 kGy/h. The absorbed doses were measured by Amber Perspex routine dosimeters [23] (dose uniformity DUR = 1.2). The irradiations were performed in triplicate.

2.4. Heat-Assisted Extraction (HAE)

All irradiated olive pomace samples were immediately lyophilized (Heto CD8, Allerod, Denmark) and stored until used. Heat assisted extraction (HAE) was performed according to a methodology previously described by Pinela et al. [24], using 0.6 g of the olive pomace with 20 mL of solvent with different conditions previously defined by the RSM design (Table 1): time (t, 20 to 120 min), temperature (T, 25–85 °C) and ethanol proportion (S, 0–100%). The solid/liquid ratio (S/L) was kept constant at 30 g/L. After the extraction in a thermostatic water bath under continuous electromagnetic stirring, the samples were centrifuged (6000 rpm for 10 min at room temperature) and filtered (paper filter Whatman n° 4) and the supernatant was collected and evaporated at 35 °C under reduced pressure (rotary evaporator Büchi R-210, Flawil, Switzerland) to remove the solvent. The obtained residue was redissolved in ethanol: H_2O (20:80, v/v) and the solution was divided into two portions for HPLC-DAD and extraction yield analysis [25].

2.5. Analysed Responses

2.5.1. Extraction Yield

The residue resulting from each extraction was determined gravimetrically using crucibles. A portion (5 mL) of the redissolved extraction liquid was dried in an oven (Memmert, Schwabach, Germany) at 60 °C to evaporate the ethanol, and then at 105 °C to evaporate the water. Afterwards, the dried sample was cooled down and the residue was calculated by difference. The results were expressed in percentage (%, w/w).

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Table 1. Circumscribed central composite design and experimental data for 5-level-3-factor response surface.

		Factors						Responses											
Run	X ₁	X ₂	X ₃	X ₁ :t	X ₂ : T [◦] C	X ₃ : S %	Y ₁	Y ₂	Y ₃	Y ₄	Y ₅	Y ₆	Y ₇	Y ₈	Y ₉	Y ₁₀	Y ₁₁		
1	-1	-1	-1	40.3	37.2	20.3	4.28	10.23	2.08	0.21	16.79	27.01	63.36	12.92	1.28	104.57	16.12		
2	-1	-1	1	40.3	37.2	79.7	2.19	7.25	1.72	0.16	11.31	20.08	75.23	15.81	1.57	112.70	10.90		
3	-1	1	-1	40.3	72.8	20.3	5.26	12.93	2.55	0.24	20.98	28.05	69.41	14.33	1.40	113.18	16.98		
4	-1	1	1	40.3	72.8	79.7	3.23	11.35	2.37	0.23	17.18	23.84	88.49	18.41	1.71	132.44	13.23		
5	1	-1	-1	99.7	37.2	20.3	3.98	9.61	2.09	0.19	15.87	28.34	69.60	15.13	1.40	114.47	13.81		
6	1	-1	1	99.7	37.2	79.7	2.21	8.35	1.84	0.16	12.55	20.66	78.26	17.24	1.65	117.81	10.67		
7	1	1	-1	99.7	72.8	20.3	5.19	12.50	2.62	0.26	20.57	27.91	66.37	13.56	1.37	109.20	18.83		
8	1	1	1	99.7	72.8	79.7	3.55	12.22	2.59	0.24	18.61	25.30	91.16	19.30	1.78	137.55	13.67		
9	1.68	0	0	120	55	50	4.87	12.55	2.64	0.26	20.33	29.51	73.64	14.24	1.48	118.87	17.80		
10	-1.68	0	0	20	55	50	4.23	11.17	2.38	0.25	18.02	28.03	72.34	14.50	1.59	116.46	14.59		
11	0	-1.68	0	70	25	50	3.46	8.81	1.79	0.16	14.22	27.38	68.68	13.51	1.25	110.82	12.63		
12	0	1.68	0	70	85	50	5.04	13.22	2.91	0.25	21.42	27.23	73.01	14.94	1.37	116.55	18.49		
13	0	0	-1.68	70	55	0	4.58	11.38	2.32	0.23	18.51	26.95	67.10	13.74	1.36	109.15	16.99		
14	0	0	1.68	70	55	100	0.66	6.59	1.51	0.13	8.89	11.37	106.03	24.85	2.06	144.31	5.83		
15	-1.68	-1.68	-1.68	20	25	0	2.86	7.13	1.42	0.14	11.54	28.68	69.86	14.28	1.33	114.15	9.96		
16	-1.68	-1.68	1.68	20	25	100	0.26	1.78	0.52	0.04	2.60	9.04	62.59	18.30	1.35	91.28	2.89		
17	-1.68	1.68	-1.68	20	85	0	5.09	12.43	2.46	0.23	20.21	27.84	68.06	13.44	1.26	110.60	18.28		
18	-1.68	1.68	1.68	20	85	100	0.99	9.08	2.00	0.16	12.23	13.93	127.65	28.10	2.29	171.97	7.11		
19	1.68	-1.68	-1.68	120	25	0	2.87	7.99	1.74	0.15	12.75	28.76	80.72	17.06	1.54	128.08	11.01		
20	1.68	-1.68	1.68	120	25	100	0.32	2.57	0.67	0.05	3.61	10.97	88.03	23.02	1.72	123.74	2.93		
21	1.68	1.68	-1.68	120	85	0	4.88	11.91	2.54	0.22	19.55	27.42	65.49	14.95	1.30	109.16	17.22		
22	1.68	1.68	1.68	120	85	100	1.28	10.44	2.35	0.18	14.26	16.21	130.81	28.85	2.26	178.12	8.30		
23	0	0	0	70	55	50	5.02	12.73	2.60	0.25	20.61	26.54	68.09	13.12	1.35	109.09	18.72		
24	0	0	0	70	55	50	4.66	12.03	2.36	0.23	19.28	27.53	70.12	15.04	1.39	114.08	17.17		
25	0	0	0	70	55	50	4.92	12.59	2.65	0.26	20.42	27.79	70.03	14.23	1.39	113.44	17.97		
26	0	0	0	70	55	50	4.94	12.58	2.55	0.25	20.32	24.84	65.56	13.31	1.28	104.99	19.31		
27	0	0	0	70	55	50	5.11	12.95	2.62	0.25	20.92	27.81	70.57	14.98	1.38	114.74	18.35		
28	0	0	0	70	55	50	4.92	12.30	2.44	0.24	19.90	27.51	68.82	14.28	1.34	111.96	17.88		

 X_1 : Time (min), X_2 : Temperature (°C), X_3 : Solvent concentration (% ethanol), and Y_1 : Hydroxytyrosol –1-beta-glucoside (mg/g DW), Y_2 : Hydroxytyrosol (mg/g DW), Y_3 : Tyrosol (mg/g DW), Y_4 : Caffeic acid (mg/g DW), Y_5 : Total phenolic content (mg/g DW), Y_6 : Hydroxytyrosol-1-beta-glucoside (mg/g Ext), Y_7 : Hydroxytyrosol (mg/g Ext), Y_8 : Tyrosol (mg/g Ext), Y_9 : Caffeic acid (mg/g Ext), Y_{11} : Yield (%).

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2.5.2. Phenolic Fingerprinting and Quantification

The phenolic fingerprinting of the extracts was determined by High Performance Liquid Chromatography (HPLC) (Prominence CBM 20-A, Shimadzu, Kyoto, Japan) with UV-DAD detector. The HPLC column was a Kinetex C18 XB-C18 (5 μ m, 250 mm, 4.0 mm, Phenomenex, Torrance, CA, USA) and the detection was made at 280, 330 and 370 nm as preference wavelengths. The phytochemical molecules were analyzed using a previously described methodology [26]. Each re-dissolved solution (3 mL) was filtered through an LC filter disk (nylon filter 0.2 μ m, 25 mm diameter, WhatmanTM, GE Healthcare, Buckinghamshire, UK). Quantitative analysis was performed using 9-level calibration curves (0.78–200 μ g/mL) obtained from commercial standards of hydroxytyrosol (y = 19203x + 8392.3, R² = 0.9999), tyrosol (y = 11762x - 7109.5, R² = 0.9999) and caffeic acid (y = 38473x - 6243.4, R² = 0.9996). The results were expressed in mg per g of dry weight (Y₁₋₅) and extract (Y₆₋₁₀), and the final responses processed for all compounds were summed up to calculate total phenolic content (TPC).

2.6. Extraction Optimization by Response Surface Methodology

2.6.1. Screening Test of Factors and Level Range for Phenolic Compounds Extraction

Primary selection and evaluation of factors and levels were carried out to determine the appropriate experimental domain for the RSM design. Independent variables, which include time (t), temperature (T) and solvent percentage (S) were preliminarily tested.

2.6.2. Experimental Design

A circumscribed central composite design (CCCD) was used, consisting of 28 randomized runs with six replicates at the central point (three replicates per condition). Fixed variables in the designed experiment were defined as X_1 : time (20–120 min), X_2 : temperature (25–85 °C), and X_3 : solvent concentration (0–100% ethanol/water). These variables were selected from a screening analysis, based on experimental data previously obtained by the research group. Each factor was tested at five different levels.

The dependent variable studied were expressed according to eleven responses (Y) format values: the total extraction yield (Y_{11}) and levels of hydroxytyrosol-1- β -glucoside $(Y_1$ and $Y_6)$, hydroxytyrosol $(Y_2$ and $Y_7)$, tyrosol $(Y_3$ and $Y_8)$ and caffeic acid $(Y_4$ and $Y_9)$ as well as the total amount resulting from the sum of the four compounds, TPC, $(Y_5$ and $Y_{10})$. To be clear, the results were expressed in mg of phenolic per g of dry weight (mg/g DW) (Y_{1-5}) and in mg of phenolic per g of extract residue (mg/g ext.) (Y_{6-10}) , both used to evaluate the total phenolic purity in the extract.

The experimental data were fitted to the second-order polynomial model (Equation (1)) to obtain the regression coefficients (b) using Statgraphics Centurion XVI (StatPoint Technologies, Inc., Warrenton, VA, USA) and Design expert 12.0.1. (Stat-Ease, Inc., Minneapolis, MN, USA) software programs. The generalized second-order polynomial model used in the response surface analysis was the following:

$$Y = b_0 + \sum_{i=1}^k b_i X_i + \sum_{i=1}^k b_{ii} X_i^2 + \sum_{i=i+1}^k b_{ij} X_i X_j$$
 (1)

where Y is the dependent variable (response variable) to be modelled, b_0 is a constant coefficient (intercept); b_i , b_{ii} and b_{ij} are the coefficients of the linear, quadratic, and interactive terms, respectively; k is the number of tested variables (k = 3); X_i and X_j are the independent variables.

2.7. Statistical Analysis

The analysis of variance (ANOVA) was carried out to determine individual linear, quadratic and interaction regression coefficient as well as model significance using Stat-graphics Centurion XVI software (StatPoint Technologies, Inc.), and the fitness of the polynomial equation to the responses was estimated using the coefficient of determination (R^2) . The significance of all the terms of the polynomial equation was analysed statisti-

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cally by computing the F value at p < 0.05. Design expert software was used to optimize the conditions of extraction throughout response surface methodology (RSM) with their respective 3D graphs and statistics diagnostics, such as predicted vs actual, residuals vs predicted, and run vs predicted points, which were assessed along the prior information mentioned in order to fit the model.

3. Results and Discussion

A previous study by the authors identified hydroxytyrosol, hydroxytyrosol-1- β -glucoside and tyrosol as the most abundant phenolic compounds present in olive pomace [1]. It was also verified that irradiation at 5 kGy increased the extractability of bioactive compounds from olive pomace by 2.4-fold compared to the non-irradiated ones. Based on this information, the aim of this work was to optimize the extraction of these three compounds for the 5 kGy irradiated samples. Caffeic acid was chosen to be studied in representation of phenolic acids present in olive pomace.

3.1. Single-Factor Effects for Polyphenolic Extractions

The results of 28 runs (all of them analysed in triplicate) using a circumscribed central composite design (CCCD) are given in Table 1, that includes the experimental design and the corresponding response data. The different response criteria used (yield and Y_{1-11}) are interesting for industrial sectors that promote the recovery of high added-value compounds from agro-industrial residues that can be used as food ingredients. The applied response criteria provide important information about the amount of olive wastes needed to achieve a certain quantity of the target compounds and their concentration in the obtained extracts.

The extraction yield ranged from 2.89 to 19.30% (Table 1). The higher value was achieved with the run 26, which combined the following conditions: t = 70 min, T = 55 °C and S = 50%. A lower yield (10.88%) was obtained by Chanioti & Tzia [27] in the ultrasound-assisted extraction of oil from olive pomace at 50 °C.

TPC (Y_5) ranged from 2.60 to 21.42 mg/g DW and the highest content was attained at the run 12. These values are higher than those reported by Böhmer-Maas et al. [14], which obtained up to 1.48 mg/g DW performing an extraction of 180 min with 80% ethanol in water solvent and at 45 °C.

3.1.1. Effect of Extraction Time on Polyphenolic Content and Extractability Yield

In order to understand the effect of time on phenolic compound extraction (Y_{1-10}) and yield (Y_{11}) from olive pomace, the experimental tests were carried out at different extraction times $(X_1 = 20, 40.3, 70, 99.7 \text{ and } 120 \text{ min})$, while the other two factors $(X_2 = \text{Temperature})$ and $X_3 = \text{Solvent}$ concentration) were adjusted according to the defined *CCCD* points presented in Table 1.

Table 2 shows the ANOVA estimated coefficients. The extractability of compounds had a small but positive trend implying that, in the range of the used extraction times, the extraction of compounds was higher when longer times were employed. Variability within the collected dataset had shown significant effect (p < 0.05, highlighted in green) in some of the tested phenolic compounds, but no significant repercussion on the extraction yield (grey coloured). This information is displayed at different points in the responses (Table 1) and is shown in Table 2, at the decoded optimization values section. Numerically, inflection points in the responses are mostly at the long end positive points tested (120 min), but also at some lower ones, e.g., 68 min for Y_2 and Y_5 . The lowest inflection point (53 min) is located at Y_1 , although this response showed no significant effect. Vitali Čepo et al. [13] and Böhmer-Maas et al. [14] also observed higher amounts of polyphenolic content and antioxidant activity in olive pomace with long extraction times, 120 min and 180 min, respectively. Furthermore, Garcia-Castello et al. [28] reported maximum total polyphenols content and antioxidant activity at 413 and 270 min, respectively, in extracts from wastes of grapefruit (*Citrus paradisi* L.)

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Table 2. Statistical analysis (ANOVA) of the *CCCD* design and decoded optimization values, including response terms for building the predictive models and optimal response values for the parametric response criteria.

ANOVA Estimated Coefficients																	
Term	Y ₁	Y ₂	Y ₃	Y_4	Y ₅	Y ₆	Y ₇	Y ₈	Y ₉	Y ₁₀	Y ₁₁						
Intercept	4.75	12.28	2.53	19.81	19.56	27.29	69.50	14.02	1.40	112.22	17.56						
A-time	-0.02	0.08	0.04	0.10	0.09	0.20	1.90	0.57	0.04	2.70	-0.10						
B-Temp	0.47	1.71	0.35	2.57	2.53	0.61	5.21	0.78	0.07	6.67	1.70						
C-Solvent	-0.99	-1.11	-0.18	-2.30	-2.28	-4.22	9.38	2.71	0.17	8.04	-2.63						
AB	0.01	-0.03	0.00	-0.02	-0.02	-0.01	-1.54	-0.26	-0.02	-1.83	0.03						
AC	0.03	0.11	0.01	0.15	0.15	0.19	0.83	0.06	0.00	1.09	0.06						
BC	-0.10	0.27	0.06	0.23	0.23	0.59	5.24	0.80	0.07	6.70	-0.20						
A^2	-0.03	-0.09	-0.01	-0.12	-0.13	0.50	0.70	0.10	0.04	1.35	-0.32		Not sig	gnificant		Significant	
B^2	-0.14	-0.39	-0.06	-0.60	-0.59	-0.02	-0.06	0.05	-0.04	-0.07	-0.54						
C^2	-0.72	-1.11	-0.22	-2.06	-2.04	-2.90	5.51	1.85	0.10	4.56	-2.01						
\mathbb{R}^2	0.98	0.97	0.97	0.98	0.98	0.96	0.96	0.97	0.95	0.94	0.96						
Lack of fit	0.10	0.06	0.38	0.49	0.09	0.25	0.02	0.35	0.08	0.14	0.12						
								Deco	ded Optim	ization							
Factor	Y ₁	Y ₂	Y ₃	Y_4	Y ₅	Y ₆	Y ₇	Y ₈	Y ₉	Y ₁₀	Y ₁₁	Global					
Time (min)	53	68	120	120	68	120	120	120	120	120	66	120					
Temperature (°C)	84.94	84.94	84.94	72.55	84.94	81.19	84.94	84.94	84.87	84.87	84.94	84.94					
Solvent (%)	26	41	45	40	36	35	100	100	100	100	28	76	-1.6		0	1	6
Single Opt Value	5.63	14.16	2.99	0.28	22.86	30.65	127.46	28.23	2.27	174.86	20						
Global Opt Value																	
(mg/g dry	$3.50 \pm$	12.55	$2.75~\pm$	$0.24~\pm$	19.04	25.09	100.77	21.19	$1.83 \pm$	148.88	13.70						
weight or mg/g	0.43	± 0.99	0.21	0.02	± 1.50	\pm 2.45	\pm 7.19	± 1.61	0.12	\pm 8.73	± 1.90						
extract)																	
Desirability	0.64	0.88	0.92	0.89	0.83	0.78	0.57	0.53	0.60	0.66	0.66	0.71					

ANOVA estimated coefficients: all numerical terms to construct second-order equations display color separated (grey: not significant). Decoded Optimization values: separated in single response optimization values (with their respective factorial conditions exhibited numerically and colored using coded values light blue for negative extreme point -1.68 light blue to positive extreme point 1.68 dark blue), global optimization, and desirability values for each of the eleven responses. Y_1 : Hydroxytyrosol-1- β -glucoside (mg/g DW), Y_2 : Hydroxytyrosol (mg/g DW), Y_3 : Tyrosol (mg/g DW), Y_4 : Caffeic acid (mg/g DW), Y_5 : Total phenolic content (mg/g DW), Y_6 : Hydroxytyrosol-1- β -glucoside (mg/g Ext), Y_7 : Hydroxytyrosol (mg/g Ext), Y_8 : Tyrosol (mg/g Ext), Y_9 : Caffeic acid (mg/g Ext), Y_{11} : Yield (%).

The range of temperature tested in this experimental design comprised five different levels ($X_2 = 25$, 37.2, 55, 72.8 and 85 °C). Many phenolic compounds present thermolabile characteristics and it is important to understand which temperatures are within the safe zone before occurring degradation, along with the identification of optimum recovery points.

Positive trend was also found for every analysed response with statistical significance (p < 0.05) (Table 2). Optimum decoded values for temperature were 85 °C in almost all responses, except for Y₄ (73 °C) and Y₆ (81 °C). It worth noting that the studied compounds seemed to be better extracted at maximum tested temperatures (85 °C). In fact, it is known that high temperatures improve the extraction efficiency [29], since it might weaken the cell membranes and hydrolyze the bonds of phenolic compounds (phenol–protein or phenol–polysaccharide) thus increasing the solubility of the compounds in the solvent [30]. The obtained results are consistent with those reported by Vitali Čepo et al. [13] who determined the higher extraction yields from olive pomace at temperatures of 70 °C and above. Also, Böhmer-Maas et al. [14] and Alu'datt et al. [12] demonstrated the higher extraction of phenolic compounds from olive pomace at higher temperatures (70 °C) using methanol as solvent.

3.1.2. Effect of Solvent Concentration on Polyphenolic Content and Extractability Yield

For solvent concentration monitoring, the whole spectrum of mixture of the two solvents were evaluated ($X_3 = 0$, 20.3, 50, 79.7 and 100% of ethanol in water). Besides exhibiting significant statistical differences (p < 0.05) in all the responses (Table 2), the employed broad ranges allowed to distinguish the maximum inflection point in the eleven different responses with strong variability among them. The results described better yield recovery on slightly polar solvents but very variable percentages within the different phenolic compounds. In fact, the optimum decoded value for the yield extraction (Y_{11}) was 28% of ethanol. The highest inflection points (100%) are located at Y_7 – Y_{10} , whereas the lowest ones (26 to 45%) are located at Y_1 – Y_6).

The selection of a suitable extraction solvent is one of the most important steps in assuring a successful extraction and the identification of optimal points is mandatory [31]. Thus, even using the same solvents, different percentages of combination are often reported [13,15,24,28,30], which may be due to the large diversification of polyphenolic compounds present in the different matrices. In previous studies, the ethanol percentage to obtain the optimum recoveries of compounds could also be very variable. A 60% ethanol in water was selected as optimal condition for extracting hydroxytyrosol, tyrosol and oleuropein from olive pomace by Vitali Čepo et al. [13], whereas for *Citrus paradisi* L. wastes the maximum total polyphenol content and total antioxidant activity were achieved using 20% and 50% aqueous ethanol, respectively, as extraction solvent [28]. On the other hand, for *Hibiscus sabdariffa* calyces the optimal condition to extract anthocyanins was using water (0% ethanol) [24] using HAE, probably due to the polarity of anthocyanins. Another study using methanol as solvent verified that the highest individual phenolics extracted from olive pomace was achieved at the concentration of 80% methanol [14].

Moreover, the obtained results demonstrated that the interaction between extraction temperature and ethanol concentration was statistically significant for all the studied phenolic compounds. On the other hand, the interaction between extraction time and ethanol concentration presented no significant effect for all the studied responses.

3.2. Optimization of HAE Process in Olive Pomace

3.2.1. Extracting Modelling and Analysis of Variance

In this study, a one-block experiment circumscribed central composite design (Table 2) based on RSM was used to optimize the HAE of TPC from olive pomace. For this purpose, a complete response analysis was performed with statistical determination of outliers of the raw data, followed by an ANOVA analysis (Table 2) coupled with numerical and graphical statistical diagnostics (Figures 1–5).

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The increasing of the yield during HAE for longer times has been explained by several authors due to the mechanical effect of stirring coupled with the physicochemical parameters employed which causes a disruption of plant cell releasing the inner compounds embedded. Additionally, mass transfer is enhanced with small particles of the matrix analyzed due to the increment of the contact surface area between solvent and plant material [32,33].

According to the performed analysis of variance, model adjustment and the coefficients for a subsequent second order equation are shown in the first part of Table 2. As mentioned above, for the significant values of the eleven responses, the cells were colored in green, while the non-significant values were colored in grey. The coefficients $R^2 > 0.94$ for each response demonstrated the high correlation. The construction of the second order equations followed the typical notation and only significant values (p < 0.05, green colored) were considered. Some examples are displayed below in the Equations (2)–(4).

$$Y_1 = 4.73810 + 0.474459 * Temperature - 0.985634 * Solvent - 0.097691 * Temperature * Solvent - 0.150406 * Temperature^2 - 0.727928 * Solvent^2$$
 (2)

$$Y_{10} = 112.86260 + 2.70371 * time + 6.67001 * Temperature + 8.04413 * Solvent - 1.83357 * time * Temperature + 6.70288 * Temperature * Solvent + 5.33699 * Solvent^2$$
 (3)

$$Y_{11} = 17.46436 + 1.69988 * Temperature - 2.62870 * Solvent - 0.663910 * Temperature^2 - 2.13429 * Solvent^2$$
 (4)

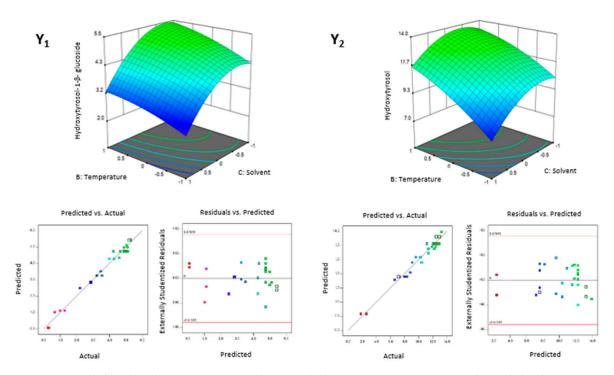


Figure 1. RSM graphs from the eleven responses with statistical diagnostic representations. Each graph displays temperature vs solvent factors, with a time factor adjusted to 0, along with two statistical diagnostics attached. On the top, the graphical representation of the adjustment between predicted vs actual points, and underneath, the graphical representation of residuals vs predicted points.

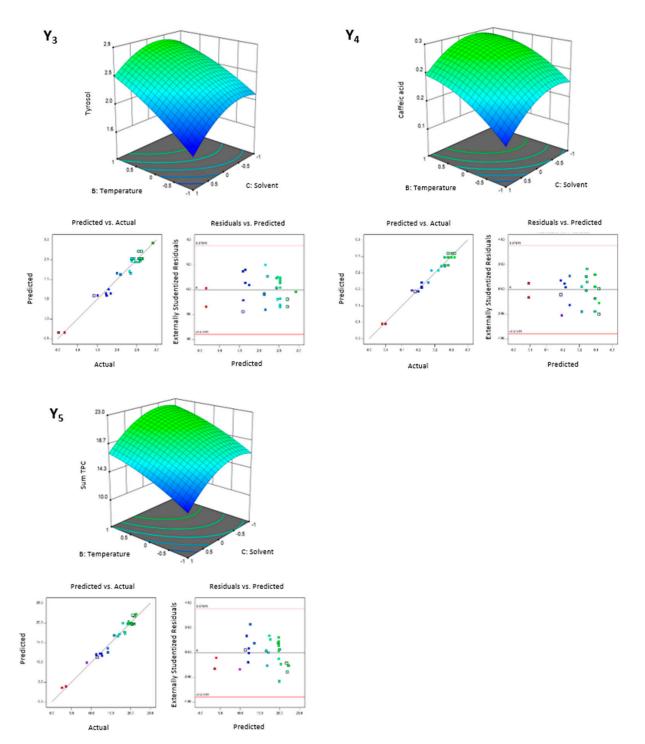


Figure 2. RSM graphs from the responses Y_3 (Tyrosol, mg/g DW), Y_4 (Caffeic acid, mg/g DW) and Y_5 (Total phenolic content, mg/g DW), with statistical diagnostic representations. Each graph displays temperature vs solvent factors, with a time factor adjusted to 0, along with two statistical diagnostics attached. On the left, the graphical representation of the adjustment between predicted vs actual points, and on the right, the graphical representation of residuals vs predicted points.

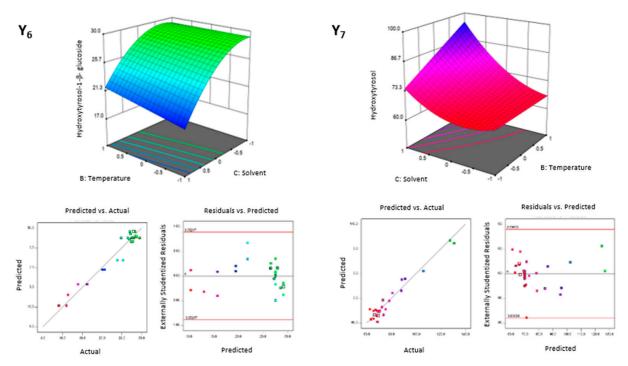


Figure 3. RSM graphs from the responses Y_6 (Hydroxytyrosol-1-beta-glucoside, mg/g Ext) and Y_7 (Hydroxytyrosol, mg/g Ext), with statistical diagnostic representations. Each graph displays temperature vs solvent factors, with a time factor adjusted to 0, along with two statistical diagnostics attached. On the left, the graphical representation of the adjustment between predicted vs actual points, and on the right, the graphical representation of residuals vs predicted points.

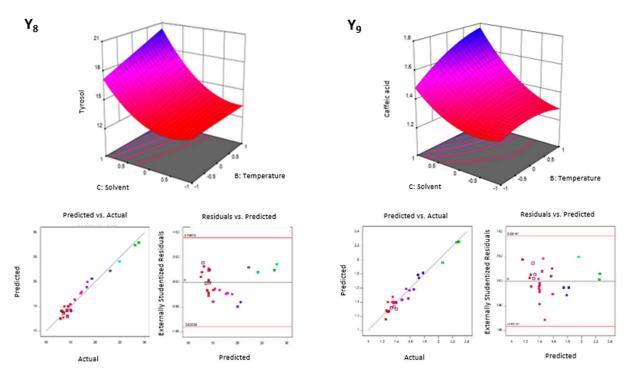


Figure 4. RSM graphs from the responses Y_8 (Tyrosol, mg/g Ext) and Y_9 (Caffeic acid, mg/g Ext), with statistical diagnostic representations. Each graph displays temperature vs solvent factors, with a time factor adjusted to 0, along with two statistical diagnostics attached. On the left, the graphical representation of the adjustment between predicted vs actual points, and on the right, the graphical representation of residuals vs predicted points.

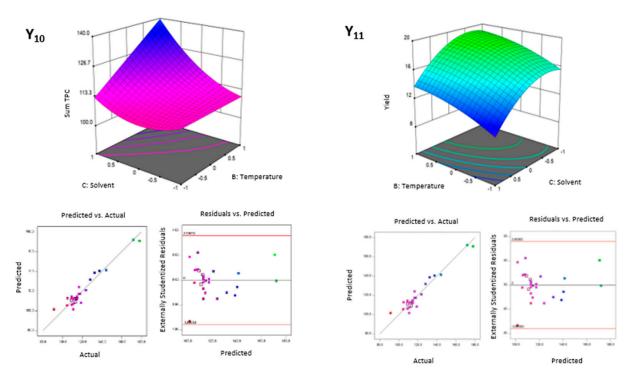


Figure 5. RSM graphs from the responses Y_{10} (Total phenolic acids, mg/g Ext) and Y_{11} (Yield, %), with statistical diagnostic representations. Each graph displays temperature vs solvent factors, with a time factor adjusted to 0, along with two statistical diagnostics attached. On the left, the graphical representation of the adjustment between predicted vs actual points, and on the right, the graphical representation of residuals vs predicted points.

3.2.2. Factorial and General Optimization

From the 28 runs (each one performed in triplicate) data acquired, the most efficient extraction conditions for maximizing each of the eleven responses using HAE were obtained by constructing 3D response surface curves with underlying contour plots (Figures 1–5) and determined by interpolation of experimental values according to the equations mentioned above.

Although ANOVA analysis becomes mandatory in understanding the patterns of the responses, 3D surface plots are the best way to visualize the effects of any independent variable on the extraction of any type of response, helping to comprehend the influence and interaction between the variables. These plots are obtained depicting two variables, temperature vs solvent, within experimental range and keeping the third variable, time, constant at zero level (Figures 1–5).

The information summarized in the eleven RSM graphs is the behavior of the responses through all the tested point, which, along with the other statistics diagnostics and mathematical equation, helps creating a net of untested points in order to give us a visual representation of data. From this representation, it becomes easy to observe that in each one of the responses, while temperature increases, the responses do it in the same manner. At the same time, it is interesting to focus on the solvent axis. Although the increments are obvious, it always presents a point of inflection that exhibits the saddle shape form on the graphs on the phenolic compound from the dry weight (Y_1-Y_5) , yield response (Y_{11}) and hydroxytyrosol-1- β -glucoside from extract (Y_6) . Those combinations of shapes indicate a uniform and increasing value for the factor temperature. Nevertheless, the same cannot be said for solvent factor, since this factor had the optimum point in the middle of the scale, somewhere between the factorial and central points for a big part of responses (Y_1-Y_6) and (Y_1-Y_6) , while the other extract responses (Y_7-Y_{10}) highlighted at the positive factorial points analyzed. This could be explained by the polarity of the compounds as well as by yielding response. On the dry weight, other polar compounds tend to increase the interaction of the

quantified compounds, hence, different profile was found when comparing with the purity of the extract. Lastly, with their own statistics diagnostic attached, it was observed how the predicted point matched well with the actual point showing a smooth straight line. Taking into consideration the lack of fit and R^2 values in Table 2, it was possible to realize how the modeling fitted with high precision (Lack of fit < 0.05 and R^2 > 0.94), except in Y_7 which might be caused by an unexpected outlier in the run 16 obtaining 62.59, instead of 71.17 mg/g of extract. On the other hand, at the bottom, the residual is shown in order to describe the behavior within the runs and check that it does not exist any type of pattern which could influence the modeling.

Furthermore, a compilation of factorial terms is presented in the second part of Table 2, which takes into account the single and interaction effects, the RSM, the statistical diagnostic plots and the quadratic equation. Giving particular interest to a unique response, for example Y_2 or Y_8 (hydroxytyrosol or tyrosol), maximal optimum values of each response are displayed (14.16 mg/g DW and 28.23 mg/g extract, respectively), as well as the specific combination of the three factors time, temperature and solvent that has to be used in order to obtain the predicted values. Besides presenting the numerical values of the used decoded ranges, the cells were colored from light to strong blue to represent the coded factorial points, intending a better observation of the factorial points.

Even if the main objective of this work was to optimize the total phenolic compounds recovery from olive pomace in order to maximize the eleven responses, a combination of the three factors (time, temperature and solvent percentage) had been adjusted and displayed at the column "Global" (Table 2). Additionally, the last row of Table 2 presents the desirability values obtained from the global optimization for each response. The global factor values for maximizing the responses were obtained at 120 min (X_1) , 85 °C (X_2) and 76 % of ethanol (X_3) , with a global desirability of 0.71. In these conditions, it was possible to extract 25.09 \pm 2.45, 100.77 ± 7.19 , 21.19 ± 1.61 and 1.83 ± 0.12 mg/g of extract of hydroxytyrosol-1- β -glucoside, hydroxytyrosol, tyrosol and caffeic acid, respectively. Moreover, extraction yield (Y_{11}) of the global optimization value was 13.70 \pm 1.9 %. The global results are similar to the obtained by Vitali Čepo et al. [13] which reported 120 min of time, 70 °C of temperature and 60% of ethanol as the best conditions to extract hydroxytyrosol, tyrosol and oleuropein from olive pomace. Concerning specifically the hydroxytyrosol extraction, the obtained results are higher than those found by Pavez et al. [34] using Pressurized Liquid Extraction and by Böhmer-Maas et al. [14] using maceration, where 0.258 mg/g DW and 0.154 mg/g DW, respectively, were extracted while 12.55 ± 0.99 mg/g DW were reached in the present work. Thus, olive pomace is a promising source of phenolic compounds namely hydroxytyrosol for potential use as natural preservatives.

Further studies have to be performed in order to valorize olive wastes as a source of potentially valuable food ingredients. For instance, other green technologies can be explored to maximize the extraction of bioactive compounds, such as supercritical extraction, microwave-assisted extraction (MAE), ultrasound-assisted extraction (UAE) [32,35,36], or pressurized-liquid extraction (PLE) [34]. The fortification of food, especially meat [37–39], beverages [40], bakery [41–43] and dairy [44,45] products, with bioactive compounds from olive wastes not only as pure compounds but also in the form of rich extracts, was previous described due to the antioxidant ability of phenols to reduce lipid oxidation during cooking and storage or to their antimicrobial properties.

4. Conclusions

Nowadays, consumers are more interested in what they eat, thus increasing the demand for sustainable and healthier foodstuffs. The food industry tries to address this challenge developing efficient extraction methods for natural added-value ingredients from industrial by-products. In this work, a heat-assisted extraction (HAE) procedure has been optimized combining three independent variables (extraction time, temperature and solvent concentration), in order to maximize the recovery of four relevant bioactive phenolic compounds (i.e., hydroxytyrosol-1- β -glucoside, hydroxytyrosol, tyrosol and

caffeic acid) present in olive pomace, a residue generated during olive oil extraction. The achieved experimental data were fitted to theoretical models to determine the optimal extraction conditions, which were established at: t = 120 min, T = 85 °C and S = 76 % of ethanol, with an extraction yield of 13.70%, allowing to recover a total amount of 148.88 ± 8.73 mg/g extract.

This work evidenced that residues produced during olive oil extraction can be valuable sources of bioactive compounds to produce ingredients that can be used by the food industry. More studies have to be performed to further improve the recovery of these and other valuable compounds to compare with the HAE results obtained in this work, such as the application of green technologies like ultrasound-assisted extraction (UAE) or microwave-assisted extraction (MAE). The isolated pure compounds from olive wastes, as well as the obtained rich extracts can be used as preservatives in foods being suitable alternatives to synthetic food additives and/or as functional ingredients to provide consumer health benefits.

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