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A Quick, Green and Simple Ultrasound-Assisted Extraction for the Valorization of Antioxidant Phenolic Acids from Moroccan Almond (*Prunus dulcis* (Mill.) D.A.Webb) Cold-Pressed Oil Residues

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Featured Application: A quick, green and simple ultrasound-assisted microextraction was here developed and validated for a quick and simple evaluation of total phenolic content from almond oil residues for their valorization as a source of antioxidant compounds.

Abstract: Almond (*Prunus dulcis* (Mill.) D.A.Webb) is one of the most important nut crops both in terms of area and production. Over the last decades, an important part of the beneficial actions for health associated with their consumption was attributed to the phenolic compounds, mainly accumulated in almond skin. Interestingly, after cold-pressed oil extraction, most of these antioxidant phenolic compounds are retained in a skin-enriched by-product, so-called almond cold-pressed oil residue. In Morocco, ranked fifth producer in the world, this production generates an important part of this valuable byproduct. In the present study, using a multivariate Box-Behnken design, an ultrasound-assisted extraction (USAE) method of phenolic compounds from Moroccan almond cold-pressed oil residue was developed and validated. Response surface methodology resulted in the optimal extraction conditions: the use of aqueous EtOH 53.0% (v/v) as green solvent, applying an US frequency of 27.0 kHz for an extraction duration of 29.4 min. The present USAE allowed substantial gains in terms of extraction efficiency compared to conventional heat reflux extraction. Applied to 3 different local *Beldi* genotypes growing at 3 different experimental sites, the optimal USAE conditions led to a total phenolic content of 13.86 mg/g dry weight (DW). HPLC analysis revealed that the main phenolic compounds from this valuable byproduct were:

chlorogenic acid followed by protocatechuic acid, *p*-hydrobenzoic acid and *p*-coumaric acid. The accumulation of these phenolic compounds appeared to be more dependent on the genetic background than on the environmental impact here represented by the 3 experimental culture sites. Both *in vitro* cell free and cellular antioxidant assays were performed, and revealed the great potential of these extracts. In particular, correlation analysis evidenced the prominent roles of chlorogenic acid, protocatechuic acid and *p*-hydrobenzoic acid. To summarize, the USAE method presented here is a quick, green, simple and efficient validated USAE for the possible valorization of antioxidant phenolic compounds from Moroccan almond cold-Pressed oil residues, making it possible to generate extracts with attractive antioxidant activities for future nutraceutical and/or cosmetic applications.

Keywords: Almond; Antioxidant; Byproducts; Chlorogenic Acid; Design of Experiment; Phenolic Acids; Ultrasound-Assisted Extraction

1. Introduction

Consumption of fruits, vegetables, nuts and seeds has been associated with lower risks of chronic and degenerative diseases [1–3]. Particularly, given their many beneficial effects on human health, in recent decades there has been growing interest in the consumption of nuts as a nutrient-rich food [3]. Produced and consumed worldwide, almond (*Prunus dulcis* (Mill.) D.A. Webb) is one of the most popular nuts. It can be consumed in the form of whole nuts, flour, beverages proposed in the food industry. A large part of almond health benefits has been ascribed to their lipid profile [3]. Almond oil is also a sought-after and attractive component for many cosmetic formulations. Over the last few decades, the part of the beneficial actions for health, but also of the growing interest for industrial applications, ascribed to almond phenolics have become increasing [3,4].

The high antioxidant capacity of almond phenolics make it an attractive alternative to synthetic antioxidants. Synthetic antioxidants were largely used to maintain the oxidative stability of emulsions and commonly used in food products and pharmaceutical and cosmetic preparations. However, synthetic antioxidants such as butylated hydroxyanisole (BHA) or butylated hydroxytoluene (BHT) have adverse health effects, including carcinogenesis [5–7]. Therefore, the use of some of these synthetic antioxidants is now prohibited for food applications in Japan, Canada and Europe and they have been removed from what is generally recognized as safe (GRAS list). The replacement of these widely criticized synthetic molecules with natural molecules would meet the expectations of manufacturers and consumers. Therefore, it is now important to identify the natural antioxidants with a pronounced and safer radical scavenging capacity for consumers. Interestingly, after cold-pressed oil extraction, most of the antioxidant phenolic compounds accumulated in the almond skin are retained in a skin-enriched by-product [3,4,8], making this almond cold-pressed oil residue (AOR) an attractive raw material for extraction and the valorization of these natural antioxidant phenolics. In Morocco, ranked fifth producer in the world, almond is the most important nut crop both in terms of area and production value. The almond plantations cover a total area of 151,000 ha for an average annual production estimated at 99,000 tons of shelled products, of which 9% of this area which provides up to 14% of the Moroccan almond production is located in Eastern Morocco (Figure 1a) [9]. This production generates an important part of byproducts, in particular of cold pressed almond oil residues. For this purpose, several local genotypes, called *Beldi*, which means “from here” as opposed to acclimatized genotypes called *Romi* (*i.e.*, from elsewhere) [10], are of special interest [9].

However, for optimal valorization of these natural co-products, the development of effective extraction methods is necessary. By the past, there were many methods developed for the extraction of natural antioxidants from various natural matrices based on conventional methods such as maceration or Soxhlet extraction. More recently, green extraction methods including microwave-

assisted extraction or ultrasonic-assisted extraction (USAE) have been found to be particularly effective [11–15]. These green extraction technologies have also aroused great interest for industrial applications, and USAE is now considered as one of the most efficient energy saving processes in terms of duration, selectivity and reproducibility, operating under mid-extraction conditions [11]. It is accepted that the improvement in extraction efficiency obtained using the USAE is based on both acoustic cavitation and mechanical effects [11]. Indeed, ultrasound (US) produces an acoustic cavitation effect facilitating the penetration of the extraction solvent. Therefore, easier release of the intracellular contents of the plant material is observed through greater agitation of the solvent resulting in increased surface contact between the solvent and the target compound as well as increased solubility of the target compound in the solvent of extraction [11].

Here, we report on the development and validation of a USAE method for the extraction of antioxidant phenolic acids from an enriched skin fraction made up of cold pressed almond oil residues (AOR) from *Beldi* Moroccan genotypes produced in Eastern Morocco (Figure 1).

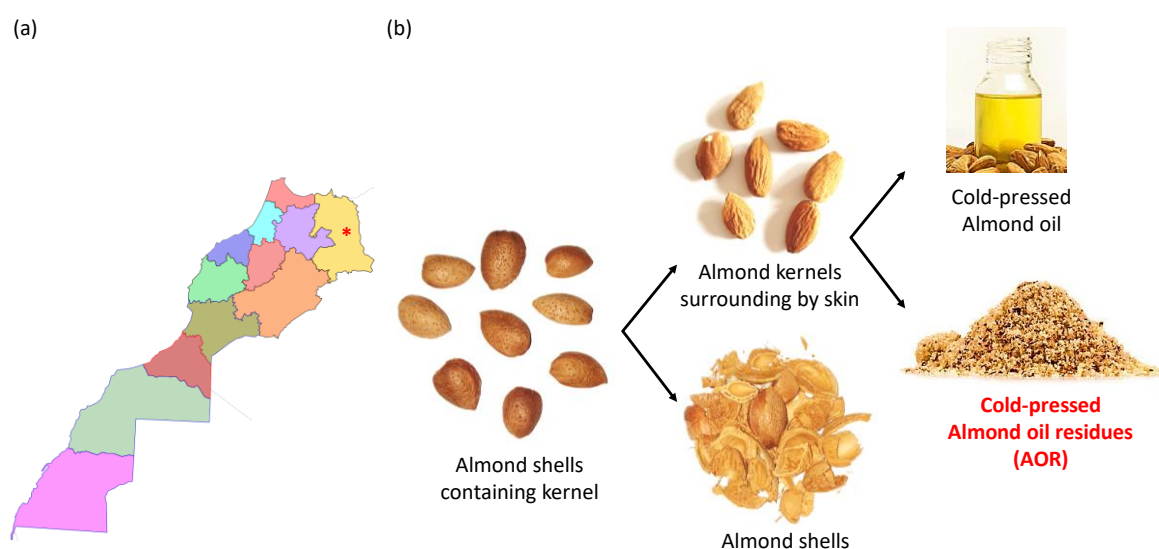


Figure 1. (a) Map of Morocco showing the region Eastern Morocco indicating by “*”. (b) Parts of almond fruits leading to cold-pressed almond oil and its residue used as byproduct in the present study to extract phenolic compounds.

Recently, Prgomet *et al.* [8] have also developed a method for comparing the polyphenol fractions from different almond byproducts including the skin using almond varieties from Portugal, but using a conventional heat reflux method. An USAE methods were developed by Kahlaoui *et al.* [16] for the extraction of polyphenols from another almond byproduct: the hulls (the part surrounding the shell itself surrounded by the thin skin; Figure 1b) from Italian and Tunisian varieties. It is thus of special interest to compare our method optimized using a different genotype, but more importantly either a green extraction method or a different (by)product. The optimal extraction conditions of this USAE using ethanol as solvent were obtained through a multivariate technique (Behnken Box design) coupled with response surface methodology (RSM) and then validated according to international standards of the association of analytical communities (AOAC). This USAE was applied to investigate the influence of the genetic and environment on the phenolic contents by considering 3 different local *Beldi* genotypes growing at 3 different experimental sites. Both *in vitro* cell free and cellular antioxidant assays were performed to evaluate the evolution of antioxidant activity of the corresponding extracts. Finally, correlations linking phytochemical profile and antioxidant activities of the extracts are presented.

2. Materials and Methods

2.1. Chemicals and Reagents

Extraction solvents (ethanol and water) used in the present study were of analytical grade (Thermo Scientific, Illkirch, France). Reagents for antioxidant assays as well as standards (chlorogenic acid, *p*-coumaric acid, protocatechuic acid and *p*-hydroxybenzoic acid) were purchased from Sigma-Aldrich (Saint-Quentin Fallavier, France).

2.2. Plant Materials and Culture Conditions

Almond oil residues were obtained from Moroccan almond (local ecotypes *Beldi*) grown in 3 different pilot locations in the Eastern Morocco (Sidi Bouhria (SID; 34°44'13.6" N, 002°20'15.0" W); Ain Sfa (AIN; 34°46'42.4" N, 002°09'28.9" W); Rislane (RIS; 34°44'59,8" N, 002°26'44.7" W)) using growing conditions as previously described by Melhaoui *et al.* [9]. Almonds were then triturated using an oil screw press (KOMET DD85G, IBG Monforts Oekotec GmbH & Co. KG, Monchengladbach, Germany) and the residues was ground to *ca* 100-150 μ m particles using a blender equipped with rotating blades (Grindomix GM 200 blender, Retsch France, Eragny, France) used as raw materials for ultrasound-assisted extraction optimization.

2.3. Ultrasound-Assisted Extraction Method Development

Ultrasound-assisted extraction (USAE) was completed with an ultrasonic bath (USC1200TH, Prolabo, Sion, Switzerland) composed of a 300 x 240 x 200 mm (inner dimension) tank, with electric power of 400W corresponding to an acoustic power of 1W/cm² and maximal heating power of 400W. The variable frequencies of this device can be selected thanks to a frequency controller, and it also has a temperature regulator as well as an automatic digital timer. Each sample was placed in 50-mL quartz tubes equipped with a vapor condenser, and was suspended in 10 ml extraction solvent. A liquid to solid ratio of 10:1 mL/g DW (dry weight) was used and extraction was performed at 45°C.

For Extraction optimization a Box-Behnken design was used and the resulting response surface plots drawn with the help of XLSTAT2019 software (Addinsoft, Paris, France). For this purpose, 3 variables (aqueous Ethanol (aqEtOH) concentration (X_1), US frequency (X_2) and extraction duration (X_3)) were studied and coded at three levels (+1, 0 and -1) as described in Table 1:

Table 1. Identity, code unit, coded levels and actual experimental values of each variable used for USAE of TPC from almond oil residues

Variable	Code unit	Coded variable levels		
		-1	0	+1
Ethanol concentration (% v/v) ¹	X_1	0	50	100
US frequency (kHz)	X_2	0	22.5	45
Extraction duration (min)	X_3	20	30	40

¹ % of ethanol (analytical grade) concentration in mixture with ultrapure water (HPLC grade).

The different batches were obtained by using the DOE (design of experiment) function of XLSTAT 2019 (Addinsoft, Paris, France), which take values of selective variables at different levels (Table 2). The experiments were carried out in triplicate. Equation of the model for the extraction of total phenolics from almond oil residues was calculated using the XLSTAT 2019 DOE analysis tool (Addinsoft, Paris, France). The corresponding response surface plots were obtained with 3D option of XLSTAT 2019 (Addinsoft, Paris, France).

2.4. Determination of Total Phenolic Content

After extraction, each extract was centrifuged for 15 min at 3,000 rpm and the resulting supernatant filtered using a syringe filter (0.45 μm , Millipore, Molsheim, France) prior to analysis. The total phenolic content (TPC) was determined spectrophotometrically using the Folin-Ciocalteu reagent (Sigma Aldrich, Saint-Quentin Fallavier, France) and according to the protocol adapted for microplate reader described by Abbasi *et al.* [17]. Briefly, 10 μL of extract were homogenized with 180 μL of a mixture composed of 4% Na_2CO_3 (prepared in NaOH 0.1 M), 0.02% potassium sodium tartrate tetrahydrate and 0.02% CuSO_4 . Following a 10-min of incubation at 25°C, 10 μL of the Folin-Ciocalteu reagent were added, and the homogenized mixture was incubated for 30 min at 25°C. Absorbance was measured at 650 nm with a spectrophotometer (BioTek ELX800 Absorbance Microplate Reader, BioTek Instruments, Colmar, France). A standard curve (0–40 $\mu\text{g}/\text{mL}$; $R^2 = 0.998$) of gallic acid (Sigma Aldrich, Saint-Quentin Fallavier, France) was used to express the TPC in mg of gallic acid equivalents per g DW (mg GAE /g DW).

2.5. Validation Parameters

Method validation was carried out using the recommendations of the association of analytical communities (AOAC) in terms of precision, repeatability and recovery as described in details in Corbin *et al.* [13].

For HPLC, 6-point calibration curves were obtained by means of diluted solutions of each authentic commercial standard (Sigma Aldrich, Saint-Quentin Fallavier, France). Each sample was injected three times, and arithmetic means was calculated to generate linear regression equations plotting was done by the peak areas (y) against the injected quantities (x) of each standard. Coefficients of determination (R^2) were used for linearity verification. The limits of detection (LOD) and of quantification (LOQ) was calculated using signal-to-noise ratios of 3:1 and 10:1, respectively.

2.6. HPLC Analysis

After extraction, each extract was centrifuged for 15 min at 3,000 rpm and the resulting supernatant filtered using a syringe filter (0.45 μm , Millipore, Molsheim, France) prior to analysis. Separation and identification of the main extract constituents was done by HPLC (High-Performance Liquid Chromatography) with a Varian system (Varian, Les Ulis, France) composed of: Prostar 230 pump, Metachem Degasit, Prostar 410 autosampler, Prostar 335 Photodiode Array Detector (PAD) and driven by Galaxie version 1.9.3.2 software. A Purospher RP-18 column (250 x 4.0 mm internal diameter; 5 μm) (Merck Chemicals, Molsheim, France) was used for the separation performed at a temperature set at 35 °C. The mobile phase was a mixture of: i) A, which was acidified HPLC grade water with acetic acid (0.2% (v/v)), and ii) B, which was HPLC grade methanol. During the separation run, the mobile phase composition varied according to a nonlinear gradient as follow: 8% B (0 min), 12% B (11 min), 30% B (17 min), 33% B (28 min), 100% B (30–35 min), 8% B (36 min) at a flow rate of 1 ml/min. Between each injection, a 10-min re-equilibration time was applied. The detection of compounds was set at 295 and 325 nm (corresponding to the λ_{max} of the main compounds). Quantification was done based on assessment of retention times of commercial standards (Saint-Quentin Fallavier, France).

2.7. In Vitro Cell Free DPPH Free Radical Scavenging Assay

The *in vitro* cell free DPPH (2,2-diphenyl-1-picrylhydrazyl) assay was used to evaluate the free radical scavenging activity of the samples as described by microplate protocol of Shah *et al.* [18]. In details, 20 μL of extract was mixed with 180 μL of DPPH reagent solution in a microplate well, and incubated in the dark at 25°C for 60 min. DPPH in mixture with 20 μL of the corresponding extraction solvent were used as negative control. Microplate reader (BioTek ELX800 Absorbance Microplate Reader, BioTek Instruments, Colmar, France) was used to record the solution absorbance at 517 nm.

For each sample, antioxidant capacity was denoted as TEAC (Trolox C equivalent antioxidant capacity) using a standard curve (0-500 μM ; $R^2=0.996$).

2.8. *In Vitro* Cell Free ABTS Antioxidant *In Vitro* Cell Free Assay

ABTS (2,2-azinobis(3-ethylbenzthiazoline-6-sulphonic acid)) *in vitro* cell free antioxidant activity of each extract was determined as described by Ullah *et al.* [19]. Briefly, after its preparation, the absorbance at 734 nm ABTS solution (ABTS salt (7 mM) and potassium persulphate (2.45 mM), incubation in the dark for at least 16 h) was adjusted to 0.7. The extract (10 μL) was mixed to 190 μL of this prepared ABTS solution, and incubated in the dark at 25°C for 15 min. After incubation, absorbance at 734 nm was determined using a micro-plate reader (Synergy II reader, BioTek Instruments, Colmar, France). For each sample, antioxidant capacity was denoted as TEAC (Trolox C equivalent antioxidant capacity) using a standard curve (0-500 μM ; $R^2=0.998$).

2.9. *Cupric Ion Reducing Antioxidant Capacity (CUPRAC) In Vitro* Cell Free Assay

CUPRAC assay was performed in microplate as described by Drouet *et al.* [7]. In short, 10 μL of extract were used, homogenized in 190 μL of the CUPRAC solution (mixture of 10 mM Cu (II), 7.5 mM neocuproine prepared in EtOH and 1 M acetate buffer pH 7 prepared in a 1:1:1 ratio (v/v/v)). After 15 min incubation at 25 °C in the dark, absorbance value at 450 nm of the reaction mixture was measured using a microplate reader (BioTek ELX800; BioTek Instruments, Colmar, France). For each sample, antioxidant capacity was denoted as TEAC (Trolox C equivalent antioxidant capacity) using a standard curve (0-500 μM ; $R^2=0.999$).

2.10. *Determination of Membrane Lipid Peroxidation Using Thiobarbituric Acid-Reactive Substances (TBARS) Assay*

An *in cellulo* antioxidant assay, using yeast cells, based on the measurement of membrane lipid peroxide was carried out with the thiobarbituric acid (TBA; Sigma Aldrich, Saint-Quentin Fallavier, France) method as described by Garros *et al.* [20]. In details, yeast cells (*Saccharomyces cerevisiae*) strain MAV203 (Invitrogen, Thermo Fisher Scientific, Villebon-sur-Yvette, France) were grown aerobically at 30 °C in an orbital shaker (150 rpm) in complete 2.0% (w/v) glucose YPD medium (Sigma Aldrich, Saint-Quentin Fallavier, France). After their evaporation under nitrogen flow, and dissolution in DMSO at 50 $\mu\text{g}/\text{mL}$, each extract was added to the cells 6 h before oxidative stress induction at a final concentration of 1 mg/mL. Here, the final concentration of DMSO applied on the cell was 1 % (v/v). Control sample was obtained by DMSO addition to 0.1% of the final volume. Cells were irradiated with 106.5 J/m² UV-C (254 nm) under a Vilber VL-6.C filtered lamp (Thermo Fisher Scientific, Villebon-sur-Yvette, France), and then incubated overnight at 30 °C prior to the evaluation of membrane lipid peroxidation. For this purpose, ca. 10⁷ cells were ground in liquid nitrogen using a mortar and pestle in distilled water, and centrifuged at 10,000xg for 10 min. Supernatant fractions (75 μL) were mixed with 25 μL of 3% (w/v) SDS, 50 μL of 3% TBA (w/v) in 50 mM NaOH, and 50 μL of 23% (v/v) of HCl throughout mixing between each addition. The mixture was then heated to 80 °C for 20 min, and after cooling on ice, the absorbance at 532 nm was measured, and non-specific absorbance at 600 nm was subtracted using Cary50 UV-Vis spectrophotometer (Varian, Les Ulis, France).

2.11. *Statistical Analysis*

Means and standard deviations of three to five independent replicates were used to present the data. Model analysis (ANOVA) and 3D plots resulting from the combination of variables were performed using XLSTAT 2019 and R analysis following the manufacturer instructions (Addinsoft, Paris, France). Student's t-test was performed for comparative statistical analysis of the impact of the different cultivation sites (XLSTAT 2019, Addinsoft, Paris, France). Correlation analysis was

performed with Past 3.0 (Øyvind Hammer, Natural History Museum, University of Oslo, Oslo, Norway) using the Pearson parametric correlation test and visualized using Heatmapper [21]. Principal Component Analysis (PCA) was performed with Past 3.0 (Øyvind Hammer, Natural History Museum, University of Oslo, Oslo, Norway). Significant thresholds at $p < 0.05$ or $p < 0.05$, < 0.01 and < 0.001 were used for all statistical tests and represented by different letters or by *, ** and ***, respectively.

3. Results and Discussions

3.1. Development of the Ultrasound-Assisted Extraction using Box-Behnken Design

Multivariate techniques are used very effectively to optimize the extraction method from complex plant matrices such as food products and by-products [22]. Among the different multivariate techniques, when 3 factors are considered, the Behnken Box design is one of the most effective technique [22,23]. The Behnken Box matrix is a spherical and rotating design, which, view on a cube, consists of the central point and the middle of the edges [22,23]. Many parameters can influence the extraction of phenolic compounds from plant matrices [24], but 3 parameters are very widely distinguished when developing an ultrasound-assisted extraction method: the type of solvent used, the frequency of ultrasound applied and the extraction time [13–15].

The choice of solvent is a crucial parameter to define when developing an extraction method. Various solvents, including methanol, ethanol (EtOH) or acetone, are regularly used for the extraction of plant polyphenols [11,25]. Here, given our objective of developing an extraction method in accordance with green chemistry principles for future nutraceutical and/or cosmeceutical applications of the resulting extract, EtOH was considered as extraction solvent. First, EtOH is one of the less toxic solvent for humans and more respectful of the environment than other organic solvents such as methanol for example [24,26]. In addition, its extraction capacity can easily be modulated by the addition of water, making it an ideal solvent for the extraction of a wide range of polyphenols of low to high polarity. Last, these two universal solvents (namely EtOH and water) have been commonly used for various food and / or cosmetic applications [11,14,15,24,26].

US frequency is a crucial parameter to consider because of its significant impact on the extraction efficiency. Indeed, this parameter modulates the cavitation effect as well as the diffusion coefficient of the target compound in the extraction solvent. Thus, it improves the solubilization of the compound in the considered extraction solvent and increases extraction efficiency [11]. In addition, increasing US frequency can also lead to a drastic reduction in extraction time, thereby reducing energy consumption, which is in accordance with the green chemistry principles [27]. However, depending on the compound and the plant matrix subjected to the extraction, application of high US frequency can alter or even destroy the native structure of the compound, which not only decreases the extraction yield, but also considerably reduces its biological activity, thus negating any valuation interest [14]. Therefore, during the development of an USAE method, US frequency must be optimized very carefully depending on the compound, and the plant matrix subjected to the extraction.

Finally, regarding the extraction time, it is important to consider that its increase does not necessarily lead to a gain in terms of extraction yield, since, on the contrary, a prolonged exposure to US can lead to the increased degradation of the compound [14]. In addition, in order to reduce the impact of energy consumption in the green chemistry context, optimizing the extraction time also appears to be essential [27].

Having these considerations in mind, in order to develop a rapid, green and efficient ultrasound-assisted extraction (USAE) of phenolic compounds (TPC) for the valorization of almond oil residues (AOR), we therefore considered a Behnken Box matrix with the following 3 parameters: aqueous

ethanol (aqEtOH) concentration (X_1), ultrasound (US) frequency (X_2) and extraction duration (X_3) as described in Table 1.

Table 2 presents the experimental and predicted total phenolic contents (TPC) obtained from almond oil residues for the 18 different observations (run ID) corresponding the different USAE conditions of the Behnken Box matrix having been determined randomly (run order) after an *in silico*-assisted procedure generated by the XL-Stat2019.4.1 software.

Table 2. Results of Box-Behnken experimental design of USAE of TPC from AOR.

Run ID	Run order	X_1	X_2	X_3	Experimental	Predicted
					TPC (mg/g DW)	TPC (mg/g DW)
Obs1	10	0	+1	-1	7.93 ± 0.14	7.93
Obs2	6	+1	0	-1	6.16 ± 0.22	6.31
Obs3	1	-1	-1	0	5.03 ± 0.11	5.18
Obs4	17	0	0	0	11.33 ± 0.10	11.37
Obs5	15	0	0	0	11.37 ± 0.08	11.37
Obs6	7	-1	0	+1	7.51 ± 0.07	7.35
Obs7	12	0	+1	+1	9.06 ± 0.13	9.23
Obs8	4	+1	+1	0	6.01 ± 0.11	5.86
Obs9	18	0	0	0	11.41 ± 0.12	11.37
Obs10	5	-1	0	-1	5.52 ± 0.05	5.36
Obs11	13	0	0	0	11.32 ± 0.17	11.37
Obs12	2	+1	-1	0	5.54 ± 0.18	5.56
Obs13	8	+1	0	+1	5.18 ± 0.14	5.16
Obs14	9	0	-1	-1	7.69 ± 0.19	7.52
Obs15	11	0	-1	+1	6.88 ± 0.13	6.88
Obs16	16	0	0	0	11.44 ± 0.17	11.37
Obs17	3	-1	+1	0	7.66 ± 0.16	7.64
Obs18	14	0	0	0	11.35 ± 0.15	11.37

Experimental values are means ± RSD of 3 independent replicates

Here, the TPC extracted from AOR ranged from: 5.03 mg/g DW (Obs3; obtained after 30 min extraction in water bath (no US application) using pure water as extraction solvent) to 11.44 mg/g DW (Obs16; obtained after 30 min at an ultrasonic bath running at a US frequency of 22.5 kHz using 50% (v/v) aqueous EtOH (aqEtOH) as extraction solvent) (Table 2). These results provide a first indication on the interest of using ultrasound and on the choice of extraction solvent. We noted a good repeatability of the central point (*i.e.*, Obs4, 5, 9, 11, 16 and 18), with a mean TPC of 11.37 ± 0.05 mg/g DW corresponding to a relative standard deviation (RDS) of 0.47%, thus highlighting the high reliability of these results. Given the nature of the starting material used in the present study, this range of TPC is in fairly good agreement with the data in the literature obtained with almond and/or almond by-products from California, Portugal, Italia and Tunisia [3,4,16,28].

A multiple regression analysis was applied to model of the TPC as a function of the 3 different extraction variables. Under the described conditions, the TPC (Y_{TPC} , in mg/g DW) as a

function of the 3 different extraction variables (*i.e.*, X_1 : aqEtOH concentration, X_2 : US frequency and X_3 : extraction duration) in the form of a polynomial equation was:

$$Y_{\text{TPC}} = 11.370 - 0.354X_1 + 0.690X_2 + 0.166X_3 - 3.554X_1^2 - 1.756X_2^2 - 1.724X_3^2 - 0.540X_1X_2 - 0.743X_1X_3 + 0.485X_2X_3 \text{ (Table 3).}$$

Table 3. Statistical analysis of the regression coefficients of USAE of TPC from AOR.

Source	Value	SD	<i>t</i>	<i>P</i> > <i>t</i>
Constant	11.370	0.059	193.27	< 0.0001***
X_1	-0.354	0.051	-6.943	0.00012***
X_2	0.690	0.051	13.543	< 0.0001***
X_3	0.166	0.051	3.253	0.011**
X_1^2	-3.554	0.069	-51.516	< 0.0001***
X_2^2	-1.756	0.069	-25.459	< 0.0001***
X_3^2	-1.724	0.069	-24.988	< 0.0001***
X_1X_2	-0.540	0.072	-7.495	< 0.0001***
X_1X_3	-0.743	0.072	-10.305	< 0.0001***
X_2X_3	0.485	0.072	6.731	0.00015***

SD standard deviation; *** significant $p < 0.001$; ** significant $p < 0.01$

The statistical analysis of the regression coefficients confirmed the relevance of our choice in the extraction variables and their respective levels for the development of the present USAE method if we refer to the level of significance with which these variables influenced the extraction (Table 3). The linear coefficients X_1 (aqEtOH concentration) and X_2 (extraction time) were statistically highly significant at $p < 0.001$, with X_1 coefficient being negative (high EtOH concentration reduced TPC) and X_2 being positive (application of US treatment had a positive effect on TPC). Extraction duration (X_3) coefficient was also significant at $p < 0.01$, but with a coefficient value close to zero indicating that a prolonged extraction period can lead to poorer extraction yield as a consequence of degradation as described in the literature [14,27,29]. All the quadratic and interaction coefficients were statistically highly significant at $p < 0.001$, but their values negative or close to zero indicated a negative or a lower impact to the extraction efficient.

The results of the analysis of variance (ANOVA) and model fitting are presented in Table 4. An elevated F-value (567.558) and low p -value ($p < 0.0001$) indicated the statistically highly significance of the model that could predict TPC as a function of the variable values with a great precision. The low non-significant value obtained for the lack of fit confirmed this trend. The value for the determination coefficient ($R^2 = 0.997$ (with adjusted value of 0.998) for the model as well as the coefficient value ($CV = 0.976$) indicated the precision of the model as well as the adequacy between the model and experimental values, respectively. The model precision in the prediction of the TPC is further depicted by the predicted *vs.* experimental TPC plot presented in Figure S1.

Table 4. ANOVA of the predicted model for used for USAE of TPC from almond residues.

Source	Sum of square	df	Mean of square	F-value	<i>p</i> -value
Model	106.071	9	11.786	567.558	< 0.0001***
Lack of fit	0.166	8	0.021	-	-

Residual	0.166	8	0.021	-	-
Pure Error	0.000	0	-	-	-
Cor. Total	106.237	17	-	-	-
R ²	0.997				
R ² adj	0.998				
CV %	0.976				

df: degree of freedom; Cor. Total: corrected total; R²: determination coefficient; R² adj: adjusted R²; CV variation coefficient value; *** significant $p < 0.001$.

To better understand the complexity of the model, 3D plots representing TPC as a function of the extraction parameters were drawn (Figure 1).

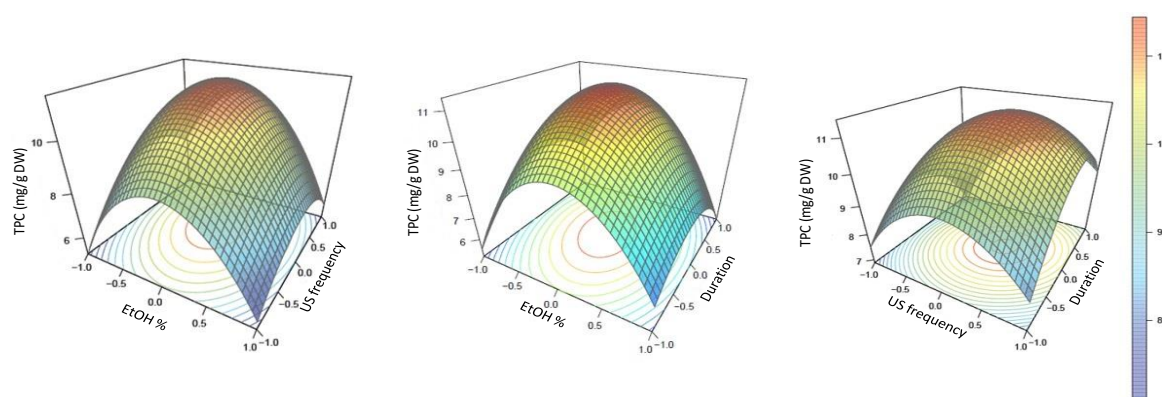


Figure 2. Predicted surface response plots of the TPC extraction yield (in mg/g DW) as a function of ethanol concentration and ultrasound frequency, ethanol concentration and extraction duration, as well as ultrasound frequency and extraction duration.

The calculated, but small, values of the linear coefficients of the second-order polynomial equation for X_2 (US frequency) and X_3 (extraction duration), as well as their interaction coefficient X_2X_3 (US frequency \times duration) indicate that a controlled increase of these parameters will have a global favorable consequences for the TPC extracted from AOR. However, their small values, in association with the negative values calculated for their quadratic coefficients (X_2^2 and X_3^2 , respectively), but also of all the coefficient involving aqEtOH concentration (*i.e.*, linear coefficient X_1 , quadratic coefficient X_1^2 , and the interaction coefficients X_1X_2 and X_1X_3), indicate that the TPC extracted from AOR according to these extraction parameters will reach a maximum value before decreasing for high values of these parameters. These considerations were clearly observed on the 3D plots (Figure 2). For each 3D plot, a first tendency was observed with a higher TPC extracted from AOR with increased aqEtOH concentration, application of US as well as prolonged extraction time. However, after reaching a maximal value for TPC extracted from AOR, a further increase in the aqEtOH concentrations as well as application of higher US frequency and/or prolonged extraction duration resulted in a pronounced drop of the TPC (Figure 2).

In various concentrations in mixture with water, aqEtOH solutions have been widely used as eco-friendly solvents to extract a wide range of polyphenols from plant matrices [13–15,24,26] including various almond products [8,16,30]. Yet, to obtain optimal results, the concentration of aqEtOH must be adapted because it is very dependent on the polyphenolic compound(s) as well as on the plant matrix considered [11,24,26]. Alongside, it is clearly established that during USAE, high US frequency associated to extended extraction duration could revealed destructive through the induction of polyphenols oxidation, in particular in presence of water [11,13,14]. Consequently, if these parameters are not finely controlled (optimized), this can lead to a sharp reduction in the extraction yield, quantitatively but also qualitatively with a drastic decay observed in the biological interest of the sample extract [12,14,15]. In our hands, using a Box-Behnken matrix for the optimization of these parameter values, and with the help of the resulting adjusted second order

polynomial equation, optimal conditions for the extraction of phenolics from our Moroccan AOR were: 53.0% (v/v) aqEtOH as solvent, 27.0 kHz for the US frequency and an extraction duration of 29.4 min. Using these optimal conditions resulted in a TPC of 11.63 ± 0.15 mg/g DW (Figure 2). The optimal aqEtOH concentration, here obtained is in line with results obtained for almond phenolics extraction very recently described [8,16]), although the starting byproduct material or the extraction method used were different from our study.

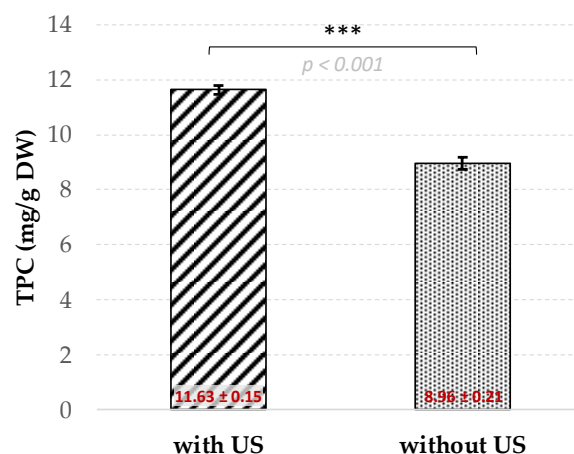


Figure 3. TPC extracted from AOR using the optimal USAE (with US) conditions and comparison with conventional heat reflux method (HRE; without US). Means \pm SD standard deviations of 3 independent extractions; *** significant at $p < 0.001$.

The present method was then validated in respect with the recommendations of the association of analytical communities (AOAC) (<http://www.aoac.org>). The parameter values of this validation procedure are satisfactory in terms of precision, repeatability and stability according to AOAC standards and are presented in Table 5. Indeed, the RSDs of both intraday and interday precisions were of 0.05 and 0.28%, respectively. The RSDs of the repeatability corresponding to five different extraction repeats of 5 samples from the same batch was of 1.30%. The recovery rates at 3 different addition levels of chlorogenic acid in the sample before extraction were between 100.26 and 101.13% reflect the accuracy of the present method.

Table 5. Validation parameters of the developed method for quantifying TPC from almond residues

Precision (%RSD)		Repeatability (%RSD)	Recovery ¹ (%)		
Intraday	Interday		0.5	1.0	2.0
0.05	0.28	1.30	100.26 \pm 0.07	100.90 \pm 0.80	101.13 \pm 0.50

¹ performed at 3 concentration level additions of gallic acid prior to extraction using optimal conditions with US (i.e. 0.5, 1.0 and 2.0 mg/g DW additions)

The efficiency of the present USAE method was compared with conventional heat reflux extraction (HRE) using the same conditions, in particular an aqEtOH concentration (53.0% (v/v) and an extraction time of 29.4 min. The difference between USAE and HRE being the application of an US frequency of 27 kHz for the present optimized UASE extraction procedure, while no US was applied for the HRE protocol operating in a classical water bath. The comparison of these extractions is depicted in Figure 2. A significant 30% gain in TPC extracted from AOR was observed with the optimized USAE (11.63 ± 0.15 mg/g DW) as compared to conventional HRE (8.96 ± 0.21 mg/g DW) (Figure 3). Increasing the extraction time for the HRE to one hour did not achieve performance levels similar to those obtained with USAE (data not shown). Consequently, it appears that the USAE method developed in the present study is of real interest according to the principles of green

chemistry [31], not only in terms of the use of a renewable green solvent, but also in terms of reducing the energy consumption. We hypothesize that this efficiency could be partly explained by the hot spot hypothesis indicating that the cavitation bubbles, after their collapse, act as a microreactor locally generating, in the surrounding solvent, a high temperature environment and pressure leading to more efficient rupture of the plant matrix subjected to extraction and increased release as well as solubilization of phenolic compounds [11].

3.2. Application to the Analysis of Samples from Different Cultivation Sites

The present USAE was then applied to the quantification of phenolics in samples from 3 different local *Beldi* genotypes cultivated at 3 different locations in Eastern Morocco. In addition to the TPC, the concentration in protocatechuic acid, *p*-hydroxybenzoic acid, chlorogenic acid and *p*-coumaric acid, reported as the main phenolic acids possibly accumulated in almond by-products [3,4,8,16,28,30,32], were also determined by HPLC after comparison with authentic commercial standards. Figure 4a shows a typical HPLC chromatogram, recorded at 325nm, of the AOR extract obtained after USAE and showing the separation of these 4 important phenolic acids: protocatechuic acid (1), *p*-hydroxybenzoic acid (2), chlorogenic acid (3) and *p*-coumaric acid (4) (Figure 4b).

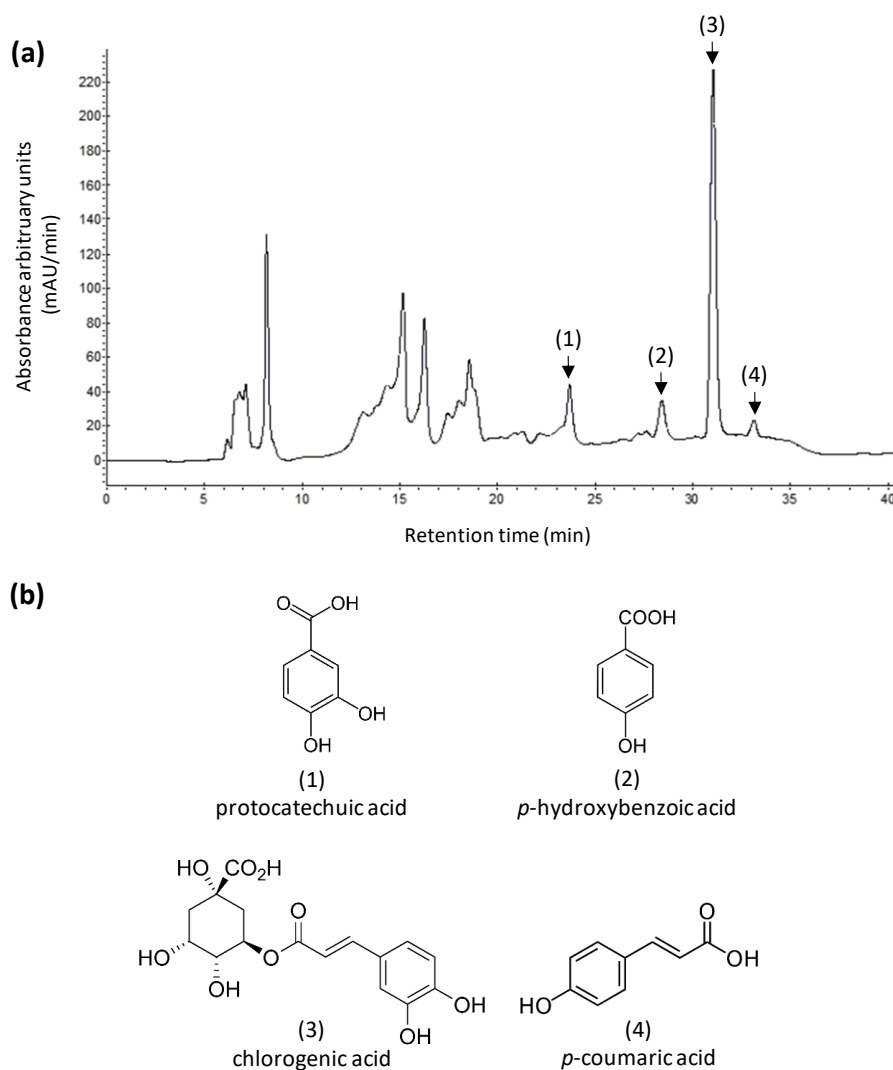


Figure 4. (a) Representative HPLC chromatogram (here with detection set at 325 nm) of an extract prepared by USAE of AOR (*Beldi* cultivar) grown in the Ain Sfa (34°46'42.4"N, 002°09'28.9"W) pilot location in the eastern Morocco. (b) Structures and their corresponding numbers on the HPLC chromatogram of the main phenolic compounds considered in this study: protocatechuic acid (1), *p*-hydroxybenzoic acid (2), chlorogenic acid (3) and *p*-coumaric acid (4).

In order to quantify these 4 phenolic compounds in different samples, 6-points calibration curves of the peak areas (y) against the injected amounts (x) of protocatechuic acid and *p*-hydroxybenzoic acid at 295 nm and chlorogenic acid and *p*-coumaric acid at 325 nm were obtained with a linearity over wide ranges from 0.5 to 200 mg/L of injected solutions and R² greater than 0.999 (Table 6). The LODs ranged from 0.12 to 0.22 mg/mL, and LOQ from 0.38 to 0.73 mg/mL, for protocatechuic acid and chlorogenic acid, respectively (Table 6).

Table 6. Quantification parameters of the HPLC method used to quantify protocatechuic acid, *p*-hydroxybenzoic acid, chlorogenic acid and *p*-coumaric acid after their USAE from AOR

Compound	RT (min)	λ_{\max} (nm)	Linear range (mg/L)	Equation	R ²	LOD (mg/L)	LOQ (mg/L)
Protocatechuic acid	23.69	295	0.5-200	$y = 3.429x + 0.814$	0.9991	0.12	0.38
<i>p</i> -Hydroxybenzoic acid	28.46	295	0.5-200	$y = 3.018x + 0.732$	0.9993	0.21	0.68
Chlorogenic acid	31.02	325	0.5-200	$y = 5.041x + 0.324$	0.9997	0.22	0.73
<i>p</i> -Coumaric acid	33.07	325	0.5-200	$y = 7.561x + 0.623$	0.9992	0.14	0.47

Applied to the quantification of TPC, protocatechuic acid, *p*-hydroxybenzoic acid, chlorogenic acid and *p*-coumaric acid in AOR resulting from samples of 3 different native *Beldi* genotypes (#1 to #3) cultivated at 3 different pilot locations in the Eastern Morocco (Sidi Bouhria (SID); Ain Sfa (AIN); Rislane (RIS)), the results are presented in Table 7.

Table 7. Variations in TPC, and protocatechuic acid, *p*-hydroxybenzoic acid, chlorogenic acid and *p*-coumaric acid contents in AOR from samples of 3 different native *Beldi* genotypes produced at 3 different pilot locations in the Eastern Morocco

Sample ID	TPC (mg/g DW)	protocatechuic acid (mg/g DW)	<i>p</i> -hydroxybenzoic acid (mg/g DW)	chlorogenic acid (mg/g DW)	<i>p</i> -coumaric acid (mg/g DW)
SID#1	9.35 ± 0.63 ^{bcd}	1.33 ± 0.10 ^{de}	0.78 ± 0.02 ^d	5.53 ± 0.13 ^e	0.26 ± 0.05 ^{ab}
SID#2	11.78 ± 1.58 ^{ab}	1.84 ± 0.07 ^{ab}	1.02 ± 0.06 ^b	7.02 ± 0.19 ^{bc}	0.29 ± 0.07 ^{ab}
SID#3	11.79 ± 1.29 ^{ab}	1.75 ± 0.09 ^b	0.98 ± 0.02 ^b	6.97 ± 0.04 ^b	0.29 ± 0.06 ^{ab}
AIN#1	8.87 ± 0.31 ^d	1.29 ± 0.06 ^e	0.75 ± 0.03 ^d	5.29 ± 0.12 ^e	0.21 ± 0.04 ^b
AIN#2	11.29 ± 1.24 ^{ab}	1.66 ± 0.10 ^{bc}	0.95 ± 0.05 ^{bc}	6.69 ± 0.06 ^c	0.28 ± 0.05 ^{ab}
AIN#3	13.86 ± 0.91 ^a	2.03 ± 0.07 ^a	1.13 ± 0.02 ^a	8.14 ± 0.10 ^a	0.26 ± 0.20 ^a
RIS#1	9.34 ± 0.27 ^{cd}	1.36 ± 0.09 ^{de}	0.75 ± 0.04 ^d	5.34 ± 0.14 ^e	0.22 ± 0.03 ^b
RIS#2	10.69 ± 0.73 ^{bc}	1.51 ± 0.05 ^{cd}	0.87 ± 0.06 ^c	6.34 ± 0.05 ^d	0.29 ± 0.02 ^a
RIS#3	11.97 ± 1.51 ^{ab}	1.76 ± 0.02 ^b	1.00 ± 0.10 ^{bc}	7.12 ± 0.08 ^b	0.30 ± 0.02 ^a

Samples were AOR from 3 different native *Beldi* genotypes (#1 to #3) cultivated at 3 different pilot locations in the Eastern Morocco: Sidi Bouhria (SID; 34°44'13.6"N, 002°20'15.0"W); Ain Sfa (AIN; 34°46'42.4"N, 002°09'28.9"W); Rislane (RIS; 34°44'59,8"N, 002°26'44.7"W). Values are means ± SD of 3 independent replicates. Different letters represent significant differences between the various extraction conditions ($p < 0.05$).

TPC ranged from 8.87 to 13.86 mg/g DW for extracts from samples AIN#1 and AIN#3, respectively; sample from genotype #3 cultivated at Ain Sfa being 56.25% richer in TPC than genotype #1 cultivated at the same location. In our hands, the 4 quantified phenolic acids occurred for approximately 80% of the TPC. In decreasing contents: 1) chlorogenic acid was the main phenolic accumulated in the sample extracts with contents ranging from 5.29 to 8.14 mg/g DW for extracts from samples AIN#1 and AIN#3, respectively (sample from genotype #3 cultivated at Ain Sfa being 53.87% richer in chlorogenic acid than genotype #1 cultivated at the same location); 2) protocatechuic acid content ranged from 1.29 to 2.03 mg/g DW for extracts from samples AIN#1 and AIN#3, respectively (sample from genotype #3 cultivated at Ain Sfa being 57.36% richer in protocatechuic acid than genotype #1 cultivated at the same location; corresponding to the highest observed variation range); 3) *p*-hydroxybenzoic acid content ranged from 0.75 to 1.13 mg/g DW for extracts from samples AIN#1 and RIS#1 for the lowest content *vs* sample AIN#3 for the highest content (sample from genotype #3 cultivated at Ain Sfa being 50.60% richer in *p*-hydroxybenzoic acid than genotype #1 cultivated both at Ain Sfa and Rislane); 4) *p*-coumaric acid content ranged from 0.21 to

0.30 mg/g DW for extracts from samples AIN#1 and AIN#3, respectively (sample from genotype #3 cultivated at Rislane being 42.85% richer in *p*-coumaric acid than genotype #1 cultivated at Ain Sfa; corresponding to the lowest observed variation range). The concentrations determined here for each phenolic compound was in the range of variations observed by Kahlaoui *et al.* [16] for different varieties of almond byproducts from Italia and Tunisia.

It is generally accepted that the genetic background, but also the environmental conditions, such as the location (*i.e.*, soil conditions) or the climate, could have a great influence on the accumulation of phenolic compounds [3,16,20,28,32,33]. The present preliminary results obtained from 3 native genotypes cultivated on the same year at 3 different location sites from Eastern Morocco, suggested a prominent influence of genetic over environment, since the impact of the genotype was more important than the influence of the cultivation site. Indeed, for each considered cultivation sites, the genotype #1 accumulated more phenolic compounds than the genotype #3, whereas both the highest and the lowest accumulation were observed on the same location (*i.e.*, Ain Sfa experimental site). Analyses of the variance (ANOVA) confirmed this absence of any significant influence of the cultivation site. Future works will be conducted with more genotypes as well as more experimental sites over several cultivation years to confirm or infirm this trend. But the prominent influence of genetic background on the accumulation of phenolic compounds in almonds was reported by several authors [3,16,28,32], whereas the influence of environmental conditions on the same genotype was less studied. Bolling *et al.* [28] reported that the cultivation season influenced less polyphenolic accumulation than the genotype. The influence of cultivation site of the same genotype will deserve further works.

3.3. Determination of the Antioxidant Potential of the Extracts and Correlation Analysis

Our next goal was to ensure that the potential biological activities is retained during the USAE procedure. For this we then determined the antioxidant potential of these 9 characterized sample extracts from AOR by using both 1) *in vitro* cell free assays based on the chemistry of the antioxidant reaction with different mechanisms - either proton transfer or electron transfer based assays; as well as 2) *in cellulo* using eukaryotic yeast cells subjected to oxidative stress induced by UV either in the presence and absence of the extracts to have an idea of their cellular antioxidant potential. Indeed, if they were preserved, this antioxidant biological activity would be of such a nature as to be of interest for both future nutraceutical and/or cosmetic applications of these AOR extracts.

The protective antioxidant action developed by plant extracts can be influenced by many internal and external factors impacting their phytochemical compositions such as genetics (the use of different genotypes in our case) but also the environment (the use of different culture sites in our case) [3,16,20,28,32,33]. Furthermore, their antioxidant activity is generally based on complex mechanisms which, in order to shorten, depending on the nature of the compounds present in the extract, can be based in particular on radical scavenging mechanisms. Here, to get an idea relating both to the antioxidant capacity but also to explore the possible mechanisms involved depending on the composition of the extract, we used three different *in vitro* cell-free assays: the DPPH (2,2-diphenyl-1-picrylhydrazyle), ABTS (2,2-azinobis(3-ethylbenzthiazoline-6-sulfonic acid)) and CUPRAC (cupric reducing antioxidant capacity) assays. These tests are based on different reaction mechanisms and could provide us a raw idea of the chemistry involved in the radical scavenging activity of the extract. Based on the chemical reaction involved, these *in vitro* cell free antioxidant assays can be roughly divided into different categories, with ABTS assay based on a hydrogen atom transfer reaction (HAT), CUPRAC assay based on an electron transfer reaction (ET), and DPPH assay being considered as a mixed assay [34,35]. The results of these antioxidant assays expressed in μM of Trolox equivalent antioxidant capacity (TEAC) per gram DW for the 9 extracts obtained after USAE of AOR are presented in Table 8.

Table 8. Variations in *in vitro* cell free (ABTS, DPPH and CUPRAC) and cellular (TBARS) antioxidant potential of extracts obtained from USAE of AOR from 3 different native *Beldi* genotypes produced at 3 different pilot locations in the Eastern Morocco

Sample ID	ABTS	DPPH	CUPRAC	TBARS
	($\mu\text{M TEAC/g DW}^1$)	($\mu\text{M TEAC/g DW}^1$)	($\mu\text{M TEAC/g DW}^1$)	(% inhibition)
SID#1	233.10 \pm 12.52 ^d	323.51 \pm 19.12 ^a	198.07 \pm 22.97 ^{ab}	51.81 \pm 1.13 ^c
SID#2	361.81 \pm 14.48 ^b	347.40 \pm 7.73 ^a	141.04 \pm 2.16 ^c	66.95 \pm 1.74 ^a
SID#3	366.49 \pm 12.97 ^b	341.17 \pm 5.49 ^{ab}	129.69 \pm 0.32 ^d	58.73 \pm 1.18 ^b
AIN#1	216.94 \pm 12.32 ^d	275.84 \pm 34.88 ^b	205.92 \pm 17.11 ^a	50.62 \pm 2.46 ^c
AIN#2	276.37 \pm 13.12 ^c	326.88 \pm 30.16 ^{ab}	164.79 \pm 14.02 ^b	51.81 \pm 1.45 ^c
AIN#3	401.52 \pm 11.44 ^a	357.33 \pm 24.24 ^a	178.73 \pm 19.10 ^{ab}	69.12 \pm 0.34 ^a
RIS#1	238.07 \pm 15.86 ^{cd}	319.73 \pm 14.74 ^b	160.93 \pm 13.74 ^{bc}	53.13 \pm 1.01 ^c
RIS#2	244.91 \pm 12.02 ^{cd}	315.60 \pm 7.43 ^b	143.76 \pm 5.24 ^{bc}	52.63 \pm 1.65 ^c
RIS#3	391.29 \pm 9.64 ^{ab}	351.07 \pm 2.89 ^a	173.47 \pm 26.29 ^{abc}	66.85 \pm 2.57 ^a

¹ TEAC: TroloxC equivalent antioxidant capacity (TEAC); Samples were AOR from 3 different native *Beldi* genotypes (#1 to #3) cultivated at 3 different pilot locations in the Eastern Morocco: Sidi Bouhria (SID); Ain Sfa (AIN); Rislane (RIS). Values are means \pm SD of 3 independent replicates. Different letters represent significant differences between the various extraction conditions ($p < 0.05$).

In our hands, antioxidant activity ranged from 216.94 to 401.52 $\mu\text{M TEAC/g DW}$ for ABTS assay, and from 275.84 to 357.33 $\mu\text{M TEAC/g DW}$ using DPPH assay. For these two *in vitro* cell free antioxidant assays, AOR extract from the genotype #3 produced at Ain Sfa showed the highest antioxidant capacity, whereas extract obtained from the genotype #1 produced at the same location displayed the lowest antioxidant values. On the contrary, results for CUPRAC assay, ranging from 129.69 to 205.92 $\mu\text{M TEAC/g DW}$, showed that this genotype #1 produced at Ain Sfa possessed the highest antioxidant capacity as compared to the genotype #3 from Sidi Bouhria.

Although interesting from a strictly predictive point of view based on chemical reactions, these *in vitro* tests do not necessarily have a great similarity with *in vivo* systems. The validity of these antioxidant data must therefore be considered as limited to an interpretation within the meaning of the chemical reactivity with respect to the considered radicals generated *in vitro*, and have to be confirmed *in vivo*. In order to have an improved understanding and better reflect the *in vivo* situation, the antioxidant activity of these 9 extracts has also been studied further for their capacity to inhibit the lipid peroxidation membrane generated by oxidative stress induced by UV-C in yeast cells. Yeast cells represent an excellent model for assessing antioxidant capacity *in vivo* in the context of cellular oxidative stress [36]. It is indeed an attractive and reliable eukaryotic model, whose defense and adaptation mechanisms to oxidative stress are well known and can be extrapolated to human cells presenting mechanisms certainly more complex but well conserved with this model [37,38]. Here, measured *in vivo* anti-lipoperoxidation activity (inhibition of malondialdehyde (MDA) formation), determined using the TBARS assay, ranged from 50.62 to 69.12%. Therefore, this *in vivo* antioxidant evaluation assay confirmed the trend observed with HAT-based *in vitro* assay, and confirmed that AOR extract from the genotype #3 produced at Ain Sfa showed the highest antioxidant capacity, particularly as compared to extracts obtained from the genotype #1 produced at the same location.

As shown in Figure 5, higher antioxidant capacity measured with HAT-based antioxidant assay appeared systematically associated with a higher accumulation of phenolics, whereas association with the ET-based antioxidant assay (*i.e.*, CUPRAC) appeared more complex and not directly linked to the accumulation of these phenolics (Figure 5a).

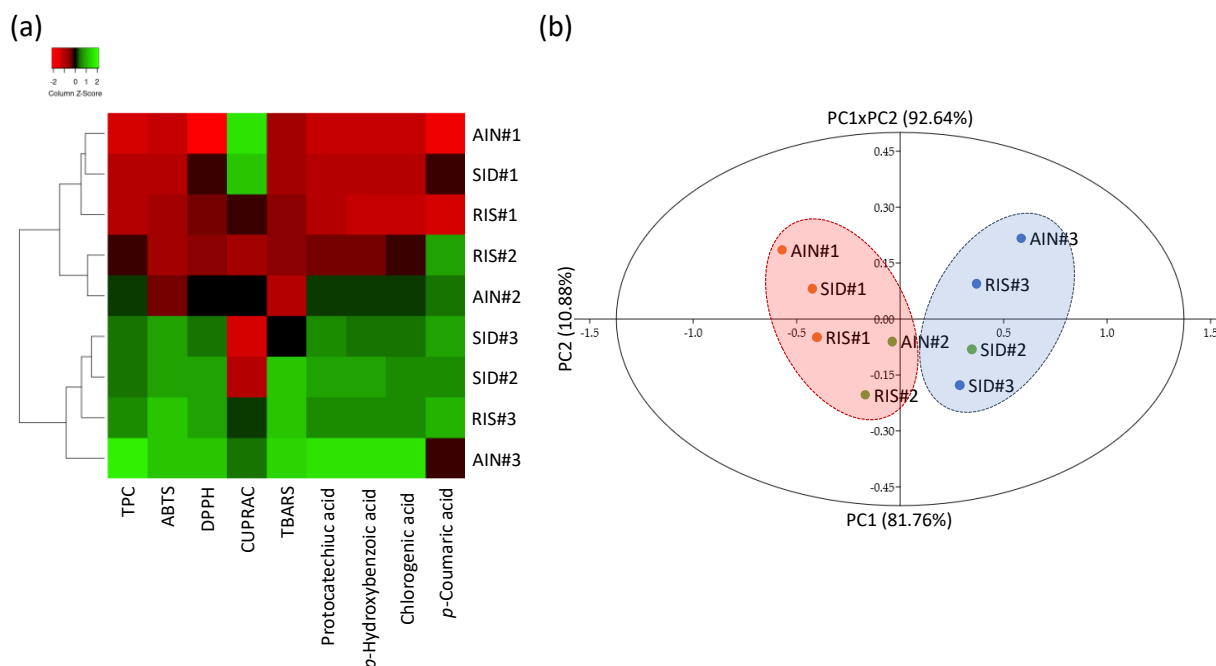


Figure 5. (a) Hierarchical clustering analysis (HCA) showing the relation between the phytochemical composition and antioxidant activity of each extracts from AOR of Eastern Morocco obtained by USAE; (b) Principal component analysis (PCA) showing the discrimination of the different extracts from AOR of Eastern Morocco obtained by USAE.

Principal component analysis was performed to further discriminate these 9 samples (Figure 5b). The resulting biplot representation accounts for 92.64% (F1 + F2) of the initial variability of the data as shown in Figure 4b. The discrimination occurs mainly in the first dimension (PC1) which explains 85.76% of the initial variability. The loading plots (Figures S2) confirmed the strong link between phytochemical composition, in particular the presence of the phenolics, and the HAT-based as well as cellular antioxidant capacity.

In order to link the antioxidant capacity to the presence of a particular phytochemical, a Pearson correlation analysis was applied (Figure 6).

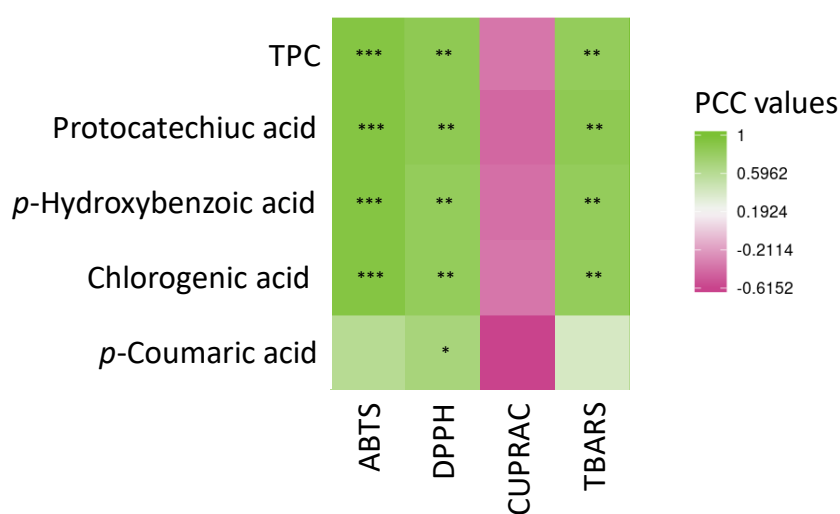


Figure 6. Pearson correlation analysis (PCC) of the relation between the main phytochemicals (protocatechiuc acid, *p*-hydroxybenzoic acid, chlorogenic acid and *p*-coumaric acid) from AOR extracts obtained after USAE and the different antioxidant assays (ABTS, DPPH, CUPRAC and TBARS). *** significant $p < 0.001$; ** significant $p < 0.01$; * significant $p < 0.05$; actual PCC values are indicated in Supplementary Materials Table S1.

This analysis evidenced the strong and highly significant correlation between both HAT-based *in vitro* assays as well as cellular antioxidant assay and TPC of the extract, in particular with the presence of chlorogenic acid, protocatechuic acid and *p*-hydroxybenzoic acid (Figure 6, Table S1). The presence of *p*-coumaric acid was significantly correlated with DPPH assay only. On the contrary, none of these phytochemicals, here analyzed, were significantly correlated with the *in vitro* ET-based antioxidant CUPRAC assay.

Altogether, these results showed a higher antioxidant activity, expressed in $\mu\text{M TEAC/g DW}$, determined with the ABTS and DPPH assays as compared to the CUPRAC assay. Therefore, these results suggested the prominence of the HAT- over the ET-based mechanism for the antioxidant action of these extracts. In good agreement, several authors have reported an antioxidant activity of extracts from various almond products based on HAT mechanism [16,28,30,32]. Similarly, a higher relation between HAT assay and phenolic acids as compared to flavonoids have been previously reported [34,39]. This observation is also in line with the results of Liang and Kitts [40] that reported a relatively stronger scavenging capacity of radicals generated by the ABTS and DPPH assays for chlorogenic acid, the main phenolic acid of our AOR extracts, and its derivatives. The authors attributed this observation to the available hydroxyl groups of these compounds. The presence of flavonoids has been also reported in almond products [8,16,28,30,32]. Here, we cannot exclude the presence of flavonoids potentially linked to the ET-based antioxidant activity evidenced by CUPRAC assay. Prgomet *et al.* [8]) have reported in the presence of flavonoids in almond skin (*i.e.*, isorhamnetin derivatives). Future works will be conducted to study in details the flavonoid fraction of our AOR extracts. Cellular antioxidant assay using yeast, further confirmed the interest of this system to study natural antioxidant from plant extract [20,34] as also previously reported for other natural antioxidants such as thiamine or and melatonin [37,38]. Natural antioxidants have aroused increasing interest over the past decade due to their possible use as an alternative to potentially dangerous synthetic antioxidants such as butylated hydroxyanisole (BHA) or butylated hydroxytoluene (BHT) in various food or cosmetic formulations [5–7]. These first results indicate a potential use of our AOR extracts generated by the present validated USAE.

4. Conclusions

P. dulcis or the so-called almond is a rich source of antioxidant phenolic compounds that are retained, after almond cold-pressed oil extraction, in a skin-enriched by-product which, thus, represent an attractive starting material for their extraction. As natural antioxidants, these phenolic compounds almond attracted much attention as alternatives to synthetic antioxidants in foods, pharmaceutical and cosmetic preparations. Here, using a multivariate Box-Behnken design coupled with surface response methodology, we proposed an optimized and validated ultrasound-assisted extraction (USAE) of these phenolic compounds from almond cold-pressed oil residue (AOR). Optimal conditions for USAE were: aqEtOH 53.0%(v/v) as green solvent, US frequency 27.0 kHz and extraction duration 29.4 min. Following its optimization, the present USAE method was validated according to international standards of the association of analytical communities (AOAC) to ensure its precision and accuracy in the quantitation of total phenolic content. The efficiency of the present USAE has allowed substantial gains in terms of extraction efficiency compared to conventional heat reflux extraction. In particular by a strong reduction in extraction time, which is of particular interest in the context of green chemistry in terms of reduction of energy consumption, together with the use of a green extraction solvent. The application of this method already makes it possible to suggest a higher impact of the genetic background than of the environment on 3 genotypes cultivated on 3 experimental sites. This method therefore opens the door to more complete studies on this subject. Finally, both *in vitro* cell free and cellular antioxidant assays revealed the great potential of valorization of these extracts as a source of natural antioxidants. To summarize, the present extraction method allows a quick, green, simple and efficient validated USAE for the possible valorization of antioxidant phenolic compounds from Moroccan almond cold-pressed oil residues,

making it possible to generate extracts with attractive antioxidant activities for future nutraceutical and/or cosmetic applications.

Supplementary Materials: The following are available online at www.mdpi.com/xxx/s1, **Figure S1:** Biplot representation of the linear relation between predicted *vs* measured TPC in the 18 Box-Behnken design sample extracts; **Figure S2:** Loading scores of the first and second axis of the principal component analysis; **Table S1:** Actual values for PCC (Pearson Correlation Coefficient) showing the relation between the different phytochemicals and antioxidant assays.

Author Contributions: Conceptualization, D.T., A.E., M.Ad. and C.H.; Methodology, D.T., A.E., M.Ab., S.D., R.K.; L.G., A.K., M.Ad. and C.H.; Software, S.D.; Validation, M.Ad., C.H., D.T.; Formal analysis, M.Ad., C.H., S.D., D.T.; Investigation, S.D., D.T. and R.K.; Resources, M.Ad. and C.H.; Data curation, M.Ad. and C.H.; Writing—original draft preparation, C.H.; Writing—review and editing, M.Ad., D.T., A.K., and C.H.; Visualization, C.H. and S.D.; Supervision, M.Ad. and C.H.; Project administration, M.Ad. and C.H.; Funding acquisition, M.Ad. and C.H.

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Conflicts of Interest: The authors declare no conflict of interest.

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