

## INFLUENCE OF THERMAL TREATMENT ON THE STABILITY OF PHENOLIC COMPOUNDS AND THE MICROBIOLOGICAL QUALITY OF SUCROSE SOLUTION FOLLOWING OSMOTIC DEHYDRATION OF Highbush BLUEBERRY FRUITS

Anna Kucner<sup>1</sup>✉, Agnieszka Papiewska<sup>1</sup>, Robert Klewicki<sup>1</sup>, Michał Sójka<sup>1</sup>, Elżbieta Klewicka<sup>2</sup>

<sup>1</sup>Institute of Chemical Technology of Food, Lodz University of Technology  
Stefanowskiego 4/10, 90-924 Łódź, Poland

<sup>2</sup>Institute of Fermentation Technology and Microbiology, Lodz University of Technology  
Wólczańska 171/173, 90-924 Łódź, Poland

### ABSTRACT

**Background.** Osmotic dehydration is a process of the partial removal of water which is based on immersion of material having cellular structure in a hypertonic solution. Osmotic dehydration is used as a pretreatment for the dehydration of foods before they are subjected to further processing such as freezing, freeze drying, vacuum drying. Management of spent syrup is one of the most important problems related to osmotic dewatering. Osmotic solutions are heavily polluted with carbohydrates, remains of the dehydrated material and microorganisms. The aim of this study was to determine the effect of thermal treatment on the content of phenolic compounds and the microbiological quality of sucrose solution used in 15 cycles of osmotic dehydration of highbush blueberry (*Vaccinium corymbosum* L.) fruits.

**Material and methods.** The tested material was 65.0 ± 0.5°Brix sucrose solution used for 15 cycles of osmotic dehydration of highbush blueberry (*Vaccinium corymbosum* L.). Osmotic dehydration was conducted at 40°C for 120 min using fruits previously subjected to enzymatic pretreatment. The thermal treatment of sucrose solution was conducted at 70, 80, 90, 100 and 115°C for 20, 40 and 60 s. The sucrose solution was analysed in terms of total polyphenols, particular polyphenols using high performance liquid chromatography and microbiological analysis was subjected.

**Results.** Thermal treatment at 70-115°C for 20 s caused degradation of 8.5% to 12.7% of polyphenols, while as much as 23.1% of polyphenols were degraded at 115°C after 60 s. The present paper proposes heating parameters that are optimal from the point of view of phenolic compound retention and microbiological quality: thermal treatment of syrup at 100°C for 40 s. Under these conditions, total polyphenols retention was 94.5%, while the retention of individual phenolic compounds varied from 89.2% to 37.2%, and that of flavan-3-ols amounted to 89.5%. The studied manner of syrup treatment eliminated the problem of syrup contamination with yeasts and molds (reducing their levels to less than 1 CFU/mL).

**Key words:** osmotic dehydration, blueberry, spent syrup, polyphenols, microbial contamination, thermal treatment

✉anna.kucner@dokt.p.lodz.pl

## INTRODUCTION

Highbush blueberry (*Vaccinium corymbosum* L.) fruits are an important source of phenolic compounds, which are thought to be natural antioxidants. According to literature data, they play a significant role in combating harmful free radicals, thus decreasing the risk of developing cardiovascular diseases [Petti and Scully 2009]. The major groups of polyphenols in highbush blueberries are phenolic acids, flavonols, anthocyanins and procyanidins [Krupa and Tomala 2007, Łata et al. 2005, Barnes et al. 2009]. Due to the seasonality of highbush blueberries, it is necessary to preserve them after harvest.

Osmotic dehydration is considered a method that minimizes adverse physical and chemical changes in foodstuffs as compared to the traditional preservation techniques used in the food industry. In the process of dehydration, the material is immersed in a solution with high osmotic pressure to induce diffusive mass transfer between the solution and the biological material being dehydrated. In industrial applications, sucrose, glucose, fructose and starch syrup solutions are most often used [Torregiani and Bertolo 2001].

The main advantage of osmotic dehydration is the fact that water is removed from the product without phase change, which significantly contributes to the preservation of the appropriate quality of the material being processed. Free intercellular spaces are filled with the osmotic substance that penetrates the product. The low-molecular weight components of vacuolar sap [organic acids, mineral salts, vitamins, phenolic compounds) may leak to the osmotic solution. Thus, the process conditions should be optimized so as to reduce the loss of valuable nutrients from fruits [Lewicki and Lenart 2006].

A factor that hampers a wider application of the discussed method is a lack of efficient ways of utilizing spent osmotic solutions. The high concentration of carbohydrates, the presence of remains of the dehydrated material, as well as the proliferation of microorganisms make such solutions difficult to reuse. Due to the high costs of purification, spent osmotic solutions are considered an industrial waste. However, as a result of the presence of valuable nutrients leaked from fruits, as well as owing to economic and environmental considerations, scientists seek ways in which at least

some of these solutions could be utilized [Dalla Rosa and Giroux 2001]. The suggestion most often made in the literature is to reuse the osmotic solution in successive cycles of the process. In a study by Valdez-Fragoso et al. [2002], osmotic dehydration of apples with a 60°Bx sucrose solution reused for 20 cycles was carried out without a significant increase in dry matter content or water loss in consecutive batches of dehydrated fruits. Uczciwek et al. [2011] have shown that one can reuse a fructooligosaccharide concentrate multiple times for dehydration of sour cherries, blackcurrants and apples.

Alternatively, spent osmotic solutions may be utilized in the production of other foodstuffs. For example, Garcia et al. [2009] recycled a sucrose solution that had been used in osmo-convective drying of mangoes as a material for wine production. According to consumers, the product was acceptable and did not differ much from commercial wines. Alternatively, spent osmotic solution may also be used as an ingredient in jams and a raw material in the production of fruit juices and natural aromas [Dalla Rosa and Giroux 2001]. Sucrose osmotic solutions may also be converted into fructooligosaccharide concentrates [Aachary and Prapulla 2009].

An important aspect of the utilization of spent osmotic solutions is microbiological contamination. Syrup microflora reflects both primary and secondary contamination, originating from the raw material and from the production process, respectively. According to literature data, spent solutions particularly susceptible to the proliferation of undesirable microorganisms are those used for the dehydration of vegetables. Solutions used for fruit dehydration are less vulnerable as their pH is usually acidic. In the literature there are no straightforward guidelines as to permissible contamination levels for spent osmotic solutions. Nevertheless, thermal treatment is proposed in order to eliminate the risks posed by excessive microbial contamination. For instance, in one study mild pasteurization (71-75°C for 30 s) reduced mold levels to 1 CFU/mL and bacteria levels to 10<sup>2</sup> CFU/mL (from 10<sup>5</sup>) [Dalla Rosa and Giroux 2001].

However, thermal treatment may lead to a reduction in the content in biologically active substances, including polyphenols, anthocyanins, phenolic acids and other important compounds, such as vitamin C.

Therefore, we sought methods that would ensure adequate microbiological purity of the syrup while retaining its desirable components.

The objective of the work was to study the influence of thermal treatment on the content of phenolic compounds and on the degradation of the microflora contaminating sucrose syrups following 15 cycles of osmotic dehydration of highbush blueberry (*Vaccinium corymbosum* L.) fruits.

## MATERIAL AND METHODS

### Material

The tested material was sucrose solution used for 15 cycles of osmotic dehydration of highbush blueberry (*Vaccinium corymbosum* L., cultivar Bluecrop) fruits. The fruits were purchased from a grower in Łódź Province (Poland) at the beginning of August 2012, and then stored at  $-20^{\circ}\text{C}$ .

### Pretreatment of fruits prior to osmotic dehydration

First,  $200 \pm 1$  g of the fruits (kept at approx.  $22^{\circ}\text{C}$  for 20 min after taking out from a freezer) was weighed into a beaker, and then 800 mL of water was added. The water (with fruits) was brought to  $\text{pH } 6.5 \pm 0.5$  with 0.1 M NaOH. Subsequently, 0.075 mL of Palatase 750-L preparation containing lipase (Novozymes A/S, Bagsvaerd, Denmark) with an enzyme activity of 750 PGU/mL was added, and the beaker was left to stand for 30 min at room temperature. Next, the fruits were separated from the solution and washed with water three times. Subsequently, the fruits (after mixing with 800 mL of water;  $\text{pH} = 4.2 \pm 0.2$ ,  $t = 25 \pm 2^{\circ}\text{C}$ ) were treated with pectinase using 1.35 mL of Pectinex Yield Mash preparation (Novozymes, Bagsvaerd, Denmark) with an enzyme activity of 46.000 PGU/mL (30 min at room temperature). Next, the fruits were separated from the solution and washed with water three times.

### Osmotic dehydration

Following enzymatic pretreatment,  $200 \pm 2$  g fruit samples were weighed into plastic containers. The dehydration process was conducted using  $65^{\circ}\text{Bx}$  sucrose solution at  $40^{\circ}\text{C}$  for 2 h with continuous vortexing (200 cycles/min). The fruit to syrup ratio was 1:4 (w/w). Following osmotic dehydration, the fruits

were drained on a sieve (rinsed with distilled water, and dried with filter paper). The syrup was reused for subsequent 14 cycles of osmotic dehydration. For each cycle, the concentration of the solution was adjusted by adding  $70 \pm 5^{\circ}\text{Bx}$  sucrose solution. A new batch of fruits was dehydrated each time.

### Thermal treatment of osmotic solution

The experiments involved thermal treatment with the use of glass capillaries with the following dimensions: an external diameter of  $d_{\text{ex}} = 1.4$  mm, an internal diameter of  $d_{\text{in}} = 0.8$  mm, and a length of 20 cm. The capillaries were filled with sucrose solution after 15 cycles of osmotic dehydration, diluted to  $15 \pm 0.5^{\circ}\text{Bx}$  with sterile distilled water. Each capillary contained 0.05 mL of solution, which amounted to approx 50% of the capillary volume. The capillaries filled in this way were sealed with the flame of a gas burner and placed in a Teflon holder made especially for that purpose. The holder was used to immerse the capillaries in an oil bath (TU-20D from Techne). Every capillary was heated with oil at a desired constant temperature: 70, 80, 90, 100 or  $115^{\circ}\text{C}$  for 20, 40 or 60 s. Following the thermal treatment, the capillaries were immediately cooled in a water bath. Subsequently, the solution was removed and subjected to microbiological analyses and tested for polyphenols.

### Microbiological analyses

- Total mesophilic bacteria count-by the pour plate method with PCA medium-incubation at  $30^{\circ}\text{C}$  for up to 3 days.
- Determination of (osmotolerant) yeasts/molds-by the pour plate method with Whaley Scar medium-incubation at  $30^{\circ}\text{C}$  for up to 5 days.
- Total thermophilic bacteria count-by the pour plate method with Cameron medium-incubation at  $55^{\circ}\text{C}$  for up to 2 days.

In the experiments, syrup samples were poured directly onto plates in two replicates for every determination. After incubation, the resulting colonies were counted and the results were given as CFU/mL of the analysed syrup solution.

### Determination of total polyphenols content

First, 0.5 mL of the sample, 0.25 mL of Folin-Ciocalteu reagent, and 2.5 mL of 20%  $\text{Na}_2\text{CO}_3$  were

placed in 25 mL volumetric flasks. Then, the flasks were filled with distilled water and the contents were mixed and incubated at room temperature for 1 h. The absorbance of the solutions was measured at 720 nm. Total polyphenols content was expressed as (-)-epicatechin equivalents [Singleton and Rossi 1965].

#### **Determination of anthocyanin content with the HPLC method**

Chromatographic analysis was performed using a Knauer HPLC chromatograph with 5uC18110A Phenomenex Gemini columns (150 × 4.60 mm) and a Security Guard Phenomenex Gemini system (4 × 3.0 mm). Separation was performed at 40°C, the flow rate was 1 mL/min and a DAD detector was used. Phase A: H<sub>2</sub>O:HCOOH (9:1, v/v), phase B: ACN:H<sub>2</sub>O:HCOOH 95:4:1, v/v) with the flow gradient: 12% B from 0 to 0.6 min, 12 to 30% B from 0.6 to 16 min, 100% B from 22 to 25 min, and 12% B from 25 to 35 min. The injection volume was 20 µL. Separation was performed on solutions obtained in point “Thermal treatment of osmotic solution”. The samples were diluted with mobile phase A (1:1 (v/v)) and centrifuged at 5000 rev./min before injection. Data were collected with Eurochrom 2000 software (Knauer, Germany). Standards obtained from Extrasynthèse (Geny, France) and Sigma-Aldrich (Steinheim, Germany), MS analyses, as well as literature data [Kalt et al. 1999, Häkkinen and Torronen 2000, Wu and Prior 2005, Castrejón et al. 2008, Lohachoompol et al. 2008, You et al. 2011] were used for the identification of anthocyanins. Quantitative results of the determinations are given as cyanidin-3-glucoside equivalents.

#### **Preparation of samples for MS analysis**

Anthocyanins were separated using a Knauer HPLC chromatograph (point “Determination of anthocyanin content with the HPLC method”) equipped with a fraction collector FOXY R1 (Teledyne ISCO Lincoln, NE, USA). Peaks were collected from ten repeated HPLC separations. The obtained samples were diluted 1:3 with distilled water and passed through the SPE columns (STRATA X, Phenomenex, Torrance, CA, USA) previously pre-conditioned with 1 mL 100% MeOH and 1 mL H<sub>2</sub>O. The retained compounds were eluted with 1 mL of 100% MeOH.

#### **LC-ESI-MS/MS analysis**

The samples prepared according to point “Preparation of samples for MS analysis” were directly injected into MS detector (LTQ VETOS, Thermo Scientific, Waltham, MA, USA). Analyses utilized the positive ion mode. The source parameters were as follows: ion spray voltage 3.00 kV, capillary temperature 325°C, sheath gas 30 units/min, auxiliary gas 10 units/min. To generate MS/MS data, the precursor ions were fragmented by helium gas collision in the ion trap by optimizing the collision energy in order to obtain an intensity of the precursor ion close to 10% of the relative scale of the spectrum.

#### **Determination of the content of flavanols (procyanidins and catechins) with the HPLC method**

First, syrups were dried in a Laboratory Freeze Dryer (Martin Christ Gefriertrocknungsanlagen, Osterode AM Harz, Germany). Subsequently, 20 mg ±1.0 mg of a sample was weighed into a 2 mL Eppendorf tube and 800 µL of methanol solution containing phloroglucinol (75 g/L) and ascorbic acid (1 g/L) was added. The reaction was started by adding 400 µL 0.4 M HCl in dry methanol. Then, the sample was mixed thoroughly and placed in a water bath at 50°C for 30 min. The samples were immediately cooled in an ice bath and the reaction was stopped by adding 600 µL 40 µM sodium acetate solution. The samples were centrifuged for 5 min at 5000 rev./min, and then diluted twice with 40 µM sodium acetate solution. Acid degradation products of polymeric proanthocyanidins were separated using a chromatograph from Knauer (Berlin, Germany) equipped with a UV-Vis detector and a fluorescence detector (FD). Separation of components was performed on C18 110A 5U Gemini columns (250 × 4.6 mm) using a gradient of two mobile phases. Phase A: 0.25% (v/v) formic acid in water; phase B: 80% (v/v) acetonitrile in water. Separation was performed at a flow rate of 1 mL/min at 25°C and a gradient of 4-7% B for 0-10 min, 7-30% B for 10-27 min, 30-70% B for 27-29 min, 70% B for 29-34 min, 70-4% B for 34-35 min, and 4% B for 35-40 min. The injection volume was 20 µL. Data were collected with Eurochrom 2000 software (Knauer). Quantitative analysis was performed using the peak area recorded by the FD detector with the excitation wavelength set

to 278 nm and the emission wavelength to 360 nm. Quantitative calculations were based on the following calibration curves: (-)-epicatechin, (+)-catechin, and (-)-epicatechin-phloroglucinol.

### Statistical analysis

The results were analysed statistically using one-way ANOVA and Duncan's multiple range test at  $p < 0.05$  with the Statistica 6.1 software (Stat Soft, Tulsa, OK, USA)

## RESULTS AND DISCUSSION

### Microbiological analysis of syrup

Experiments showed that thermal treatment of the sucrose solution led to a decrease in the number of microorganisms, the extent of which depended on the type of microbes and process parameters (Table 1). Total mesophilic bacteria count in untreated solution ranged from 4 to  $6 \times 10^2$  CFU/mL, and dropped to  $3-6 \times 10^1$  CFU/mL following heating at 70, 80 and 90°C for 20, 40 and 60 s. Within this temperature range, it would be perhaps advisable to apply longer thermal treatment to achieve a more efficient degradation of mesophilic bacteria. If the treatment is

conducted for too short a period, the bacteria may only be weakened (reversible mutations may be induced), and subsequently they may regenerate. Treatment at 100°C led to greater destruction of this group of microorganisms, especially at longer exposure times (after 20 s their survivability amounted to  $1.3 \times 10^1$  CFU/mL, while after 60 s it decreased to less than 10 CFU/mL). At a higher temperature (115°C) and longer exposure times, the survival of mesophilic bacteria in the heated syrup solution remained at less than 10 CFU/mL.

The syrup used in the experiments is susceptible to spoilage caused by yeasts and molds due to the presence of sucrose [Dalla Rosa and Giroux 2001]. Contamination with yeast and mold cells (especially those tolerating the high sugar concentrations in the starting solution) ranged from 2.5 to  $4 \times 10^1$  CFU/mL. According to the literature, an excessive accumulation of this kind of microflora may lead to adverse changes in the quality of food products [Dalla Rosa and Giroux 2001, Sapata et al. 2009]. The thermal treatment applied in our experiments, in all the tested temperature variants, resulted in an effective reduction of yeast and mold contamination to a level of less than 10 CFU/mL, and, at 100 and 115°C, to less than 1 CFU/mL.

**Table 1.** Influence of thermal treatment on the survival of microorganisms present in sucrose solution following 15 cycles of osmotic dehydration

Microbial group CFU/ml	Time s	Syrup before heat treatment	Temperature, °C				
			70	80	90	100	115
Total mesophilic bacteria	20	$5 \cdot 10^2$ <sup>a</sup>	$6 \cdot 10^1$ <sup>b</sup>	$2 \cdot 10^1$ <sup>b</sup>	$2 \cdot 10^1$ <sup>b</sup>	$1.3 \cdot 10^1$ <sup>b</sup>	<10 <sup>b</sup>
	40	$5 \cdot 10^2$ <sup>a</sup>	$2 \cdot 10^1$ <sup>b</sup>	$3 \cdot 10^1$ <sup>b</sup>	$2 \cdot 10^1$ <sup>b</sup>	<10 <sup>b</sup>	<10 <sup>b</sup>
	60	$5 \cdot 10^2$ <sup>a</sup>	$2 \cdot 10^1$ <sup>b</sup>	$2 \cdot 10^1$ <sup>b</sup>	$1.3 \cdot 10^1$ <sup>b</sup>	<10 <sup>b</sup>	<10 <sup>b</sup>
Osmotolerant yeast molds	20	$3.5 \cdot 10^1$ <sup>a</sup>	<1 <sup>b</sup>	<1 <sup>b</sup>	<10 <sup>b</sup>	<10 <sup>b</sup>	<10 <sup>b</sup>
	40	$3.5 \cdot 10^1$ <sup>a</sup>	<1 <sup>b</sup>	<1 <sup>b</sup>	<1 <sup>b</sup>	<1 <sup>b</sup>	<10 <sup>b</sup>
	60	$3.5 \cdot 10^1$ <sup>a</sup>	<10 <sup>b</sup>	<1 <sup>b</sup>	<10 <sup>b</sup>	<10 <sup>b</sup>	<1 <sup>b</sup>
Total thermophilic bacteria	20	<1 <sup>a</sup>	<10 <sup>a</sup>	<10 <sup>a</sup>	<1 <sup>a</sup>	<1 <sup>a</sup>	<1 <sup>a</sup>
	40	<1 <sup>a</sup>	<1 <sup>ab</sup>	<1 <sup>a</sup>	<10 <sup>b</sup>	<1 <sup>a</sup>	<1 <sup>a</sup>
	60	<1 <sup>a</sup>	<10 <sup>ab</sup>	<10 <sup>b</sup>	<1 <sup>a</sup>	<1 <sup>a</sup>	<1 <sup>a</sup>

The same letter in a given line indicates a lack of statistical differences ( $\alpha = 0.05$ ) between the results at particular processing times.

### Influence of thermal treatment on the content of phenolic compounds

Total polyphenols content in the starting sucrose solution following 15 cycles of dehydration was  $729.2 \pm 31.6$  mg/kg IDM. This figure is relatively low as compared to the amount of polyphenols in the studied raw material, but previous research [Kucner et al. 2012] has shown that the above-mentioned conditions of osmotic dehydration are optimal from the point of view of phenolic compound retention in fruits and osmotic dehydration effectiveness.

The data given in Table 2 indicate that thermal treatment of syrups leads to a reduction in total polyphenols as compared to the control. In all the tested temperature variants, treatment time influenced the content of the studied compounds. The longer the syrup was exposed to high temperatures, the greater the losses of the compounds were. For example, after 20 s of heating of the solution at 90°C, total polyphenols retention amounted to 96.1%, while after 60 s it dropped to 87.3%. The most dramatic decline in the content of the studied compounds was observed at 115°C: after 60 s total polyphenols content fell to 76.9% as compared to the starting syrup. After heating the solution at 100°C, total polyphenols retention (94.5% to 77.9%) was similar to that obtained at 80°C (97.8% to 77.9%). Numerous studies have shown that the main factors reducing polyphenols content in fruits during processing are temperature and process time. For instance, Kłopotek et al. [2005] reported a 27.3% loss of polyphenols following 5 min pasteurization of strawberry juice at 85°C. Blanching of highbush blueberry

fruits at 95°C for 2 min led to a total polyphenols loss of about 10% [Ścibisz and Mitek 2006]. Thus, the results obtained in our tests are similar to literature data.

A detailed phenolic profile of the highbush blueberry fruits used in the process of osmotic dehydration is given in Table 3.

Our experiments show that not all anthocyanins present in the fruits migrated to the syrup. Among all the anthocyanins identified in the syrup, malvidin derivatives were the most abundant (62.9%). The osmotic solution was found to have a relatively high content of chlorogenic acid, at a concentration of  $108.6 \pm 4.6$  mg/kg IDM. Similarly as in the case of total polyphenols content, there exists a relationship between the process temperature and time and the content of these compounds. Heating led to gradual degradation of phenolic compounds. At temperatures ranging from 70 to 115°C, chlorogenic acid content declined slightly, by up to 7%. However, after 60 s, the content of this compound was halved. This was also the case with particular anthocyanin compounds. The smallest anthocyanin losses were recorded in samples exposed to thermal treatment at 70°C and for 20 s (98.8% to 97.5% retention). The compound most vulnerable to heating was a derivative of malvidin and pentose (its exact structure was not determined), whose retention after 60 s was 24.8% and 8.2% at 90°C and 100°C, respectively. Much greater stability was revealed by a compound described as malvidin + acetoxy + hexose (I). Following 40 s of thermal treatment at 70°C its retention amounted to 75.3%, while at 115°C it was 65.5%. These results are in agreement with the observations

**Table 2.** Influence of thermal treatment on total polyphenols content in syrup following 15 dehydration cycles, mg/kg IDM

Time s	Syrup before heat treatment mg/kg IDM	Temperature, °C				
		70	80	90	100	115
20	$729.2 \pm 31.6^a$	$667.7 \pm 7.9^b$	$712.4 \pm 23.7^{ab}$	$636.7 \pm 4.3^c$	$655.5 \pm 31.8^{bc}$	$656.0 \pm 41.2^{abc}$
40	$729.2 \pm 31.6^a$	$650.9 \pm 43.3^{ab}$	$708.8 \pm 4.1^{ab}$	$700.5 \pm 45.4^{ab}$	$688.7 \pm 0.7^{ab}$	$608.4 \pm 13.5^b$
60	$729.2 \pm 31.6^a$	$595.0 \pm 15.8^c$	$567.9 \pm 6.7^c$	$667.8 \pm 23.6^b$	$567.8 \pm 3.2^c$	$561.1 \pm 8.0^c$

The same letter in a given line indicates a lack of statistical differences ( $\alpha = 0.05$ ) between the results at particular processing times.

**Table 3.** Influence of thermal treatment on the content of selected polyphenols in syrup following 15 dehydration cycles, mg/kg IDM

Compound	Syrup before heat treatment	Temperature, °C				
		70	80	90	100	115
<b>Time 20 s</b>						
Chlorogenic acid	108.57 ±4.59 <sup>a</sup>	105.94 ±1.48 <sup>a</sup>	101.22 ±0.44 <sup>a</sup>	103.12 ±5.52 <sup>a</sup>	104.21 ±6.48 <sup>a</sup>	104.31 ±1.23 <sup>a</sup>
Cyanidin-3-O-galactoside	1.60 ±0.07 <sup>a</sup>	1.58 ±0.06 <sup>a</sup>	1.59 ±0.04 <sup>a</sup>	1.51 ±0.06 <sup>a</sup>	1.55 ±0.07 <sup>a</sup>	1.49 ±0.02 <sup>a</sup>
Malvidin-3-O-glucoside	28.11 ±0.78 <sup>a</sup>	27.77 ±0.26 <sup>a</sup>	25.03 ±0.02 <sup>b</sup>	24.96 ±2.12 <sup>b</sup>	26.88 ±2.07 <sup>ab</sup>	26.38 ±0.20 <sup>ab</sup>
Malvidin-3-O-arabinoside	14.09 ±0.45 <sup>a</sup>	13.78 ±0.03 <sup>a</sup>	13.34 ±0.19 <sup>ab</sup>	13.53 ±0.74 <sup>a</sup>	13.19 ±1.98 <sup>ab</sup>	12.77 ±0.41 <sup>b</sup>
NI	22.94 ±0.69 <sup>a</sup>	22.48 ±0.11 <sup>a</sup>	17.76 ±0.41 <sup>b</sup>	18.05 ±0.82 <sup>b</sup>	20.51 ±2.46 <sup>ab</sup>	19.99 ±0.61 <sup>ab</sup>
Malvidin + pentose	0.80 ±0.05 <sup>a</sup>	0.79 ±0.01 <sup>a</sup>	0.66 ±0.06 <sup>b</sup>	0.70 ±0.06 <sup>b</sup>	0.72 ±0.02 <sup>b</sup>	0.68 ±0.02 <sup>b</sup>
Malvidin + acetyl + hexose (I)	0.12 ±0.03 <sup>a</sup>	0.11 ±0.03 <sup>a</sup>	0.08 ±0.02 <sup>a</sup>	0.09 ±0.02 <sup>a</sup>	0.11 ±0.01 <sup>a</sup>	0.06 ±0.01 <sup>b</sup>
Malvidin + acetyl + hexose (II)	0.65 ±0.09 <sup>a</sup>	0.61 ±0.01 <sup>a</sup>	0.57 ±0.17 <sup>a</sup>	0.64 ±0.07 <sup>a</sup>	0.61 ±0.06 <sup>a</sup>	0.59 ±0.00 <sup>a</sup>
Malvidin + acetyl + hexose (III)	2.85 ±0.09 <sup>a</sup>	2.78 ±0.12 <sup>a</sup>	2.64 ±0.04 <sup>b</sup>	2.70 ±0.13 <sup>a</sup>	2.58 ±0.34 <sup>b</sup>	2.50 ±0.08 <sup>b</sup>
<b>Time 40 s</b>						
Chlorogenic acid	108.57 ±4.59 <sup>a</sup>	94.12 ±3.61 <sup>b</sup>	81.35 ±9.39 <sup>b</sup>	81.37 ±9.63 <sup>b</sup>	62.96 ±3.38 <sup>c</sup>	54.11 ±5.44 <sup>c</sup>
Cyanidin-3-O-galactoside	1.60 ±0.07 <sup>a</sup>	1.17 ±0.02 <sup>b</sup>	1.24 ±0.20 <sup>b</sup>	1.57 ±0.34 <sup>a</sup>	1.15 ±0.38 <sup>b</sup>	0.81 ±0.30 <sup>b</sup>
Malvidin-3-O-glucoside	28.11 ±0.78 <sup>a</sup>	21.69 ±0.61 <sup>b</sup>	16.76 ±2.27 <sup>c</sup>	17.22 ±2.50 <sup>cd</sup>	10.66 ±2.53 <sup>d</sup>	6.51 ±1.74 <sup>c</sup>
Malvidin-3-O-arabinoside	14.09 ±0.45 <sup>a</sup>	10.96 ±0.41 <sup>b</sup>	8.78 ±0.80 <sup>c</sup>	9.54 ±1.02 <sup>c</sup>	6.06 ±0.21 <sup>d</sup>	4.14 ±0.52 <sup>c</sup>
NI	22.94 ±0.69 <sup>a</sup>	17.31 ±0.59 <sup>b</sup>	13.07 ±1.98 <sup>c</sup>	12.42 ±6.07 <sup>c</sup>	7.40 ±0.16 <sup>d</sup>	5.09 ±0.14 <sup>c</sup>
Malvidin + pentose	0.80 ±0.05 <sup>a</sup>	0.62 ±0.11 <sup>b</sup>	0.43 ±0.04 <sup>c</sup>	0.50 ±0.03 <sup>c</sup>	0.29 ±0.09 <sup>d</sup>	0.16 ±0.00 <sup>c</sup>
Malvidin + acetyl + hexose (I)	0.12 ±0.03 <sup>a</sup>	0.09 ±0.01 <sup>a</sup>	0.08 ±0.04 <sup>a</sup>	0.04 ±0.00 <sup>b</sup>	0.10 ±0.00 <sup>a</sup>	0.08 ±0.02 <sup>a</sup>
Malvidin + acetyl + hexose (II)	0.65 ±0.09 <sup>a</sup>	0.60 ±0.04 <sup>a</sup>	0.52 ±0.05 <sup>a</sup>	0.51 ±0.04 <sup>a</sup>	0.57 ±0.04 <sup>a</sup>	0.54 ±0.07 <sup>a</sup>
Malvidin + acetyl + hexose (III)	2.85 ±0.09 <sup>a</sup>	2.18 ±0.02 <sup>b</sup>	1.83 ±0.08 <sup>c</sup>	1.87 ±0.09 <sup>c</sup>	1.25 ±0.06 <sup>d</sup>	0.86 ±0.06 <sup>c</sup>
<b>Time 60 s</b>						
Chlorogenic acid	108.57 ±4.59 <sup>a</sup>	60.37 ±0.64 <sup>c</sup>	74.65 ±0.10 <sup>b</sup>	65.14 ±6.61 <sup>bc</sup>	52.44 ±2.64 <sup>d</sup>	54.14 ±0.55 <sup>d</sup>
Cyanidin-3-O-galactoside	1.60 ±0.07 <sup>a</sup>	1.04 ±0.08 <sup>b</sup>	1.33 ±0.08 <sup>c</sup>	0.86 ±0.08 <sup>d</sup>	0.91 ±0.05 <sup>b</sup>	1.23 ±0.05 <sup>c</sup>
Malvidin-3-O-glucoside	28.11 ±0.78 <sup>a</sup>	10.16 ±0.04 <sup>c</sup>	15.16 ±0.02 <sup>b</sup>	11.13 ±1.28 <sup>c</sup>	7.11 ±0.53 <sup>d</sup>	6.96 ±0.30 <sup>d</sup>
Malvidin-3-O-arabinoside	14.09 ±0.45 <sup>a</sup>	5.61 ±0.02 <sup>c</sup>	8.23 ±0.01 <sup>b</sup>	5.80 ±0.63 <sup>c</sup>	3.27 ±0.33 <sup>d</sup>	3.08 ±0.23 <sup>d</sup>
NI	22.94 ±0.69 <sup>a</sup>	7.51 ±0.09 <sup>c</sup>	11.64 ±0.05 <sup>b</sup>	8.42 ±1.08 <sup>c</sup>	4.51 ±0.31 <sup>d</sup>	4.24 ±0.29 <sup>d</sup>
Malvidin + pentose	0.80 ±0.05 <sup>a</sup>	0.29 ±0.06 <sup>c</sup>	0.41 ±0.06 <sup>b</sup>	0.20 ±0.00 <sup>d</sup>	0.17 ±0.00 <sup>d</sup>	0.22 ±0.09 <sup>cd</sup>
Malvidin + acetyl + hexose (I)	0.12 ±0.03 <sup>a</sup>	0.07 ±0.01 <sup>b</sup>	0.09 ±0.01 <sup>a</sup>	0.06 ±0.01 <sup>a</sup>	0.06 ±0.01 <sup>b</sup>	0.06 ±0.03 <sup>b</sup>
Malvidin + acetyl + hexose (II)	0.65 ±0.09 <sup>a</sup>	0.59 ±0.06 <sup>a</sup>	0.50 ±0.07 <sup>a</sup>	0.57 ±0.13 <sup>a</sup>	0.55 ±0.04 <sup>a</sup>	0.55 ±0.06 <sup>a</sup>
Malvidin + acetyl + hexose (III)	2.85 ±0.09 <sup>a</sup>	1.15 ±0.02 <sup>c</sup>	1.75 ±0.02 <sup>b</sup>	1.15 ±0.12 <sup>c</sup>	0.75 ±0.07 <sup>d</sup>	0.77 ±0.06 <sup>d</sup>

NI – not identified.

The same letter in a given line indicates a lack of statistical differences ( $\alpha = 0.05$ ).

**Table 4.** Influence of thermal treatment on the content of flavan-3-ols in syrup following 15 dehydration cycles, mg/kg IDM

Time s	Syrup before heat treatment mg/kg IDM	Temperature, °C				
		70	80	90	100	115
20	66.6 ± 1.4 <sup>a</sup>	56.7 ± 1.7 <sup>b</sup>	49.6 ± 0.6 <sup>c</sup>	54.2 ± 2.3 <sup>b</sup>	62.2 ± 0.4 <sup>a</sup>	55.2 ± 4.0 <sup>b</sup>
40	66.6 ± 1.4 <sup>a</sup>	51.5 ± 2.6 <sup>bc</sup>	46.9 ± 1.0 <sup>cd</sup>	44.2 ± 1.9 <sup>cd</sup>	59.6 ± 1.1 <sup>ab</sup>	36.7 ± 3.4 <sup>d</sup>
60	66.6 ± 1.4 <sup>a</sup>	44.4 ± 3.8 <sup>b</sup>	44.4 ± 1.3 <sup>b</sup>	41.2 ± 0.5 <sup>b</sup>	24.9 ± 4.9 <sup>c</sup>	27.6 ± 0.8 <sup>c</sup>

The same letter in a given line indicates a lack of statistical differences ( $\alpha = 0.05$ ) between the results at particular processing times.

made by Ścibisz et al. [2010], who showed that among the anthocyanins present in highbush blueberry fruit-those containing galactose or glucose (hexoses) were more resistant to 40 min treatment at 95°C than those containing arabinose (a pentose).

Anthocyanins are labile compounds that are not very resistant to the action of high temperatures. Brownmiller et al. [2008] reported anthocyanin losses of 43% during the production of highbush blueberry paste (3 min blanching at 95°C).

According to literature data, under the influence of heat glycosidic bonds in dye molecules undergo hydrolysis leading to unstable aglycones, which easily oxidize forming brown, high molecular weight compounds [Ścibisz et al. 2010].

Furthermore, the influence of thermal treatment on the content of flavan-3-ols in spent osmotic syrup was analyzed. Prior to treatment, the amount of flavan-3-ols in the syrup was 66.1 ± 1.4 mg/100 g IDM. The most dramatic drop in the content of these compounds was caused by 60 s heating at 115°C (41.5% retention). Their degradation was the smallest after 20 s of thermal treatment (over 80% retention).

Analysis of the results obtained for the influence of thermal treatment on microbial destruction and the stability of phenolic compounds shows that the optimal variant is heating the syrup at 100°C for 40 s. Such parameters enable the elimination of the vast majority of microorganisms with relatively low losses of phenolic compounds. As can be seen from Table 1, thermal treatment of the syrup at 70, 80 and 90°C does not inactivate the syrup microflora as effectively as that conducted at 100°C or more. Heating the syrup

at 100°C for 40 s results in a total polyphenols retention of 94.5%; the retention of particular phenolic compounds ranges from 89.2% to 37.2%, while that of flavan-3-ols is 89.5%. These figures are higher than those obtained during thermal treatment of the syrup at 70°C for 60 s (the corresponding results were 81.6%, from 91.4 to 32.7% and 66.6%).

## CONCLUSIONS

The present study revealed considerable variation in the content of phenolic compounds depending on the parameters of thermal treatment. Irrespective of the temperature used, after 20 s of treatment total polyphenols content did not change significantly (91.5% to 87.3% retention). Particularly sensitive to high temperatures were anthocyanin dyes and flavan-3-ols, and the content of these compounds gradually declined during heating of spent osmotic solution. The most dramatic degradation of these compounds was observed at 115°C after 60 s of thermal treatment: 78.2% of anthocyanins and 58.5% of flavan-3-ols were lost. Thermal treatment at temperatures ranging from 70 to 90°C did not reduce microbiological contamination as effectively as heating at 100 and 115°C (for 40 s). From the point of view of both the microbiological quality of the syrup and the stability of the studied phenolic compounds, the optimal treatment was heating at 100°C for 40 s. Total polyphenols retention amounted to 94.5%; that of particular phenolic compounds ranged from 89.2% to 37.2%, and that of procyanidins was 89.5%. Such process parameters made it possible to achieve considerable biological

purity-less than 10 CFU/mL for mesophilic bacteria and less than 1 CFU/mL for yeasts and molds.

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## WPŁYW OBRÓBKII TERMICZNEJ NA STABILNOŚĆ ZWIĄZKÓW FENOLOWYCH I JAKOŚĆ MIKROBIOLOGICZNĄ ROZTWORU SACHAROZY PO ODWADNIANIU OSMOTYCZNYM OWOCÓW BORÓWKI WYSOKIEJ

### STRESZCZENIE

**Wstęp.** Odwadnianie osmotyczne to proces prowadzący do częściowego zmniejszenia zawartości wody, który jest realizowany poprzez zanurzenie materiału mającego strukturę komórkową w roztworze hipertonicznym. Odwadnianie osmotyczne jest stosowane jako obróbka wstępna do odwodnienia żywności przed poddaniem jej procesom takim, jak: zamrażanie, suszenie sublimacyjne, suszenie próżniowe. Zagospodarowanie roztworów po odwadnianiu jest jednym z najważniejszych problemów związanych z odwodnieniem osmotycznym. Syropy po odwadnianiu osmotycznym są zanieczyszczone przez węglowodany, pozostałości z odwodnionego materiału i mikroorganizmy. Celem pracy było zbadanie wpływu obróbki termicznej na zawartość związków fenolowych i czystość mikrobiologiczną syropów sacharozy po 15 cyklach odwadniania osmotycznego owoców borówki wysokiej (*Vaccinium corymbosum* L.).

**Materiał i metody.** Materiałem badawczym był roztwór sacharozy – 65,0 ±0,5°Brix użyty w 15 cyklach odwadniania osmotycznego owoców borówki wysokiej (*Vaccinium corymbosum* L.) odmiany 'Bluecrop'. Odwadnianie osmotyczne prowadzono w temperaturze 40°C, przez 120 min, stosując owoce wcześniej poddawane wstępnej obróbce enzymatycznej – 70, 80, 90, 100, 115°C w czasie 20, 40 i 60 s. Termiczną obróbkę roztworu sacharozy prowadzono w temperaturze 70, 80, 90, 100 i 115°C przez 20, 40 and 60 s. W roztworze analizowano polifenole ogółem, poszczególne polifenole oraz przeprowadzono analizę mikrobiologiczną.

**Wyniki.** Termiczna obróbka w temperaturze 70-115°C w ciągu 20 s powodowała degradację od 8,5 do 12,7% polifenoli w czasie 60 s, natomiast w temperaturze 115°C uległo degradacji 23,1% polifenoli. Zaproponowano optymalne warunki z punktu widzenia związków fenolowych i jakości mikrobiologicznej: obróbka syropu w 100°C przez 40 s. W tych warunkach retencja polifenoli ogółem wynosiła 94,5%, poszczególnych związków fenolowych od 89,2 do 37,2% i flawan-3-oli 89,5%. Ten sposób utrwalania syropu pozwolił na obniżenie poziomu drożdży (poniżej 1 CFU/ml), a także bakterii termofilnych (poniżej 1 CFU/ml).

**Słowa kluczowe:** odwadnianie osmotyczne, borówka amerykańska, zużyty syrop, polifenole, zanieczyszczenie mikrobiologiczne, obróbka termiczna

Received – Przyjęto: 25.05.2013

Accepted for print – Zaakceptowano do druku: 5.11.2013

For citation – Do cytowania

Kucner A., Papiewska A., Klewicki R., Sójka M., Klewicka E., 2014. Influence of thermal treatment on the stability of phenolic compounds and the microbiological quality of sucrose solution following osmotic dehydration of highbush blueberry fruits. Acta Sci. Pol., Technol. Aliment. 13(1), 79-88.