

## ELICITATION AND TREATMENT WITH PRECURSORS OF PHENOLICS SYNTHESIS IMPROVE LOW-MOLECULAR ANTIOXIDANTS AND ANTIOXIDANT CAPACITY OF BUCKWHEAT SPROUTS

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### ABSTRACT

**Background.** Recently, an increase of interest in the modification of food products on each step of production (breeding, production technology, storage condition) is observed. Nutritional properties as well as level and activity of bioactive compounds in plant-origin food may be modified using a range of technological and biotechnological practices and elicitation should be mentioned between them.

**Material and methods.** Elicitation with willow bark infusion supported by feeding with the phenylpropanoid pathway precursors were used for improving the quality of buckwheat sprouts. Special emphasis has been placed on the metabolomic and biochemical changes and the mechanism of overproduction of low-molecular antioxidants.

**Results.** The accumulation of phenolics is caused by stimulation of two main enzymes the phenylpropanoid pathway (tyrosine ammonia-lyase and phenylalanine ammonia-lyase). Tyrosine ammonia-lyase activities were effectively induced by feeding with tyrosine (about four times that of the control), whereas phenylalanine ammonia-lyase activity was the highest in the elicited control sprouts and those fed with shikimic acid (an increase by 60% compared to the control). Shikimic acid feeding (both elicited and non-elicited sprouts) effectively improved the total phenolics (by about 10% and 20%, respectively), condensed tannins (by about 30% and 28%, respectively), and flavonoids (by about 46% and 70%, respectively). Significant increase of vitexin, rutin, chlorogenic acid and isoorientin contents was also observed. The treatments increased the ascorbic acid content, too. Total antioxidant capacity of sprouts was most effectively increased by feeding with shikimic acid and further elicitation.

**Conclusion.** The studies transfer biotechnology commonly used for the induction of overproduction of secondary metabolites in plant cell line systems to low-processed food production. The obtained results could be used for better understanding of the effect of elicitation and precursor feeding on antioxidants production and contribute to improving the buckwheat sprouts quality.

**Key words:** antioxidant activity, buckwheat sprouts, elicitation, low-molecular weight antioxidants, phenylpropanoid pathway, precursor feeding

### INTRODUCTION

An increased interest has been noticed in the modification of food along each step of its production (breeding, production technology, storage condition). Briefly

speaking nutritional properties as well as the level and activity of bioactive compounds present in plant-origin food may be modified using a range of technological

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and biotechnological practices. There are two main ways of improving the antioxidant content in sprouts. The first one comprises genetic manipulation and is based on metabolic engineering (Liu et al., 2007). The second involves modification of the sprouting process, including elicitation and supported elicitation (Baenas, et al., 2014).

Elicitors are chemicals or biofactors from various sources, which can induce physiological changes in the target living organism. It may include abiotic elicitors such as metal ions, inorganic compounds, chilling etc., and biotic elicitors from fungi, bacteria, viruses, plant cell wall components, as well as chemicals that are released at the attack site by plants under a pathogen or herbivore attack (e.g., signaling compounds such as salicylic acid, jasmonic acid) (Baenas et al., 2014; Matkowski, 2008). Willow bark is an excellent source of salicylic acid and its derivatives (Durak and Gawlik-Dziki, 2014) and its infusion have already been used for induction of plants metabolism (Gawlik-Dziki et al. 2013; Świeca and Dziki, 2015). Salicylic acid (SA) is a well-known inducer of plant systematic acquired resistance (SAR) in plant-pathogen interaction. SA quickly accumulates at the site of infection during a pathogen attack, as well as during a hypersensitive reaction, and it spreads to other parts of the plant to induce a wide range of defense responses (Yuan and Lin, 2008).

The pro-health quality of low-processed food, e.g., sprouts, is strongly determined by its chemical composition (Pérez-Balibrea, et al., 2011; Świeca and Baraniak, 2014a; 2014b; 2014c). Phenolics and ascorbic acid – compounds with well-documented biological activity (Fang et al., 2002; Gawlik-Dziki et al., 2012) – play a very important role in the creation of food bioactivity. The effectiveness of antioxidants results from various mechanisms of activity: antioxidants could inhibit free-radical reaction by inhibiting lipid radical formation, disrupting propagation of chain auto-oxidation reactions, suppressing singlet oxygen, they could act as factors that aid in reducing hydrogen peroxides to stable compounds, as compounds chelating transition metal ions and as inhibitors of pro-oxidative enzymes (Carocho and Ferreira, 2013). Thus, antioxidant activity, as the fundamental property of food, is important for its health protecting ability, including its

antimutagenic, anticarcinogenic, antiobesity and antiaging effects.

So far, studies aimed to improve the antioxidant composition of sprouts have involved optimizing biosynthesis through the modifications of culture conditions (Khattak et al., 2007; Shett and McCue, 2003; Świeca et al., 2012), precursor feeding and biotransformation (Pérez-Balibrea et al., 2011; Świeca et al., 2014c), as well as elicitation and stress induced production (Gawlik-Dziki et al., 2013; Kim et al., 2011; Tsurunaga et al., 2013; Świeca, 2015). Elicitors increase the activity of key enzymes involved in the synthesis of low-molecular antioxidant e.g. activities of tyrosine/phenylalanine ammonia-lyase, chalcon synthase in the phenylpropanoid pathways (phenolics synthesis). Additionally the effectiveness of synthesis may be enhanced by enrichment of culture with precursors e.g. shikimic acid, phenylalanine and tyrosine (phenolics synthesis) (Baenas et al., 2014; Świeca et al., 2014c) or galactose, mannose (ascorbic acid synthesis) (Gallie, 2013). Most importantly, there are only a few studies concerning the mechanisms of obtaining the desirable features.

Therefore, in this study, a metabolomic and biochemical analysis (including an analysis of low-molecular antioxidants contents and activities involved in phenolic synthesis and metabolism) was performed on buckwheat sprouts. The effects of elicitation with *Salix* bark infusion and the phenylpropanoid pathway precursors feeding on the the changes in antioxidant capacity were also determined. The study aimed to widen the knowledge about the mechanisms of elicitation, as a result of which the health promoting properties of plant-food products could be strengthened.

## MATERIAL AND METHODS

### Chemicals

Ferrozine (3-(2-pyridyl)-5,6-bis-(4-phenyl-sulphonic acid)-1,2,4-triazine), ABTS (2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid), linoleic acid, ammonium thiocyanate, polyvinylpyrrolidone and haemoglobin were purchased from Sigma-Aldrich company (Poznań, Poland). All other chemicals were of analytical grade.

## Materials

Buckwheat (*Fagopyrum esculentum* Moench) seeds were purchased from PNOS S.A. in Ożarów Mazowiecki, Poland. Seeds were sterilized in 1% (v/v) sodium hypochloride for 10 min, then drained and washed with distilled water until they reached neutral pH. After that seeds were placed in distilled water (C, control and E, elicitor treatment) or phenolic precursor solution (0.1 mM shikimic acid – Sh and Sh+E 0.1 mM L-phenylalanine – Phe and Phe+E; 0.1 mM L-tyrosine – Tyr and Tyr+E) and soaked for 4 h at 25°C. Seeds were dark germinated for 3 days in a growth chamber on Petri dishes (ϕ 125 mm) lined with absorbent paper (approximately 400 seeds per dish). Seedlings were watered daily with 5 mL of Milli-Q water. For elicitor treatment (E, Sh+E, Phe+E and Tyr+E), 1-day-old sprouts were watered with 5 ml of 0.01% infusion of *Salix myrsinifolia* bark. The plates were then covered and sprouts were germinated under control conditions. Bark of willow (*Salix daphnoides*), obtained from an ecological farm in Poland, was dried and pulverized in a laboratory mill. Elicitor solution was prepared by brewing 1 g of the material with 100 mL of boiling water for 15 min.

## Analysis of low-molecular weight antioxidants

### • PHENOLICS

**Extraction procedure.** Flours (0.25 g in triplicate) were extracted three times with 4 ml of ethanol: water (80:20, v/v). After centrifugation (10 min, 6800 × g) fractions were collected, combined and used for further analysis of phenolics and antioxidant capacity.

**Total phenolics.** The amount of total phenolics was determined using Folin-Ciocalteu reagent (Singleton et al., 1974). The amount of total phenolics was calculated as a gallic acid equivalent (GAE) in mg per g of dry mass (d.m.).

**Total flavonoids** content was determined according to the method described by Lamaison and Carnat (1990). Total flavonoids content was calculated as a quercetin equivalent (QE) in mg per g of dry mass (d.m.).

**Condensed tannin** content was determined according to the method described by Sun et al. (1998). Condensed tannin content was calculated as a (+)-catechin equivalent (CE) in mg per g of dry mass (d.m.).

**Quantitative – qualitative analysis of phenolic.** Samples were analysed with a Varian ProStar high-performance liquid chromatography (HPLC) system separation module (Varian, Palo Alto, CA, USA) equipped with Varian ChromSpher C18 reverse phase column (250 mm × 4.6 mm) and ProStar DAD detector (Świeca and Baraniak, 2014b). Quantitative determinations were carried out with the external standard calculation, using calibration curves of the standards. Phenolics were expressed in µg per gram of dry mass (d.m.).

### • EXTRACTION AND DETERMINATION OF VITAMIN C CONTENT

Total ascorbic acid content was determined as a sum of ascorbic and dehydroascorbic acid, according to modified versions of the methods described earlier by Campos et al. (2009). Briefly, 500 mg of fresh sprouts was extracted two times with 2 mL of 50% (w/v) m-phosphoric acid (MPA). The mixture was centrifuged at 16,000 g, and extracts were combined and used for further determination. 500 µl of extract was mixed with 500 µl dithiothreitol 30 mM in 0.5 M Trizma buffer for 5 min. Then, 650 µl of 5% (w/v) MPA was added. Samples were filtered using a 0.45 µm Millipore and analysed with a Varian ProStar HPLC System separation module (Varian, Palo Alto, CA) equipped with a Varian ChromSpher C18 reverse phase column (250 × 4.6 mm) column and a ProStar 325 UV-Vis detector. The column thermostat was set at 25°C. The separation was performed under isocratic elution conditions using a mobile phase consisting of 30 mM KH<sub>2</sub>PO<sub>4</sub> adjusted with 5 M HCl to pH 3, at a flow-rate of 0.8 mL·min<sup>-1</sup> and detection at 245 nm. Quantitative determination was conducted with an external standard calculation, using calibration curves of the standard. Ascorbic acid content was expressed in µg per 1 g of fresh mass (FM).

### • EXTRACTION AND DETERMINATION OF CHLOROPHYLLS AND CAROTENOIDS

Chlorophylls and carotenoids were analyzed according to the method described by Lin et al. (2013). Chlorophylls and carotenoids were eluted overnight from the freeze-dried leaves samples (0.05 g) with 2 mL 80% (v/v) acetone at 4°C. The sample was then centrifuged at 13,000 g for 5 min. The supernatant was applied to determine the absorbance of chlorophyll a,

chlorophyll b and carotenoids in acetone, as measured with a spectrophotometer, at the respective wavelengths of 663, 645 and 470 nm. Concentrations of chlorophyll a, chlorophyll b and carotenoids were determined from the following equations:

$$\text{Chlorophyll a} = 12.72 A_{663} - 2.59 \times A_{645}$$

$$\text{Chlorophyll b} = 22.88 \times A_{645} - 4.67 \times A_{663}$$

$$\text{Carotenoids} = (1000 \times A_{470} - 3.27 \times \text{chlorophyll a} - 104 \times \text{chlorophyll b}) / 229$$

Carotenoids were expressed in mg/100 g d.m.

### Antioxidant activities

#### • ANTIRADICAL ACTIVITY (ABTS)

The experiments were carried out using an improved ABTS decolorization assay (Re et al., 1999). Free radical scavenging ability was expressed as a Trolox equivalent in mg per g of dry mass (d.m.).

#### • REDUCING POWER (RP)

Reducing power was determined by the method of Oyaizu (1986). Reducing power was expressed as a Trolox equivalent in mg per g of dry mass (d.m.).

#### • METAL CHELATING ACTIVITY (CHP)

Chelating power was determined by the method of Decker and Welch (1990). Chelating power was expressed as an EDTA equivalent (EDTA) in mg per g of dry mass (d.m.).

#### • INHIBITION OF LINOLEIC ACID PEROXIDATION (LPI)

The inhibition of the hemoglobin-catalyzed peroxidation of linoleic acid was determined according to Groupy et al. (2007). The activity was expressed as a Trolox equivalent (TE) in mg per g of dry mass (d.m.).

### Enzymatic activities

#### • EXTRACT PREPARATION

All enzyme extraction procedures were conducted at 4°C. For phenylalanine ammonia-lyase (PAL) and tyrosine ammonia-lyase (TAL), 200 mg of lyophilized sample were ground with 2 ml extracting buffer (0.2 M boric acid buffer containing, 1 mM EDTA, and 50 mM  $\beta$ -mercaptoethanol, pH 8.8). The extracts were then homogenized and centrifuged at 12 000  $\times$  g at 4°C for 30 min, and the supernatant was collected. For

polyphenol oxidase (PPO) and guaiacol peroxidase (POD), 200 mg of the lyophilized sample were ground with 4 ml of 100 mM sodium phosphate buffer (pH 6.4) containing 0.2 g of polyvinylpyrrolidone. The extracts were then homogenized and centrifuged at 12 000  $\times$  g at 4°C for 30 min, and the supernatants were collected.

#### • ENZYME ASSAY

**Peroxidase (POD) assay.** POD activity was determined using guaiacol as the substrate (Ippolito et al., 2000). The reaction mixture consisted of 0.1 mL of crude extract and 2 mL of guaiacol (8 mM, in 100 mM sodium phosphate buffer, pH 6.4) was incubated for 1 min at 30°C. The increase in absorbance at 460 nm was measured after 1 mL H<sub>2</sub>O<sub>2</sub> (24 mM) was added. The activity of POD was expressed as U, where U = 0.001  $\Delta$ OD<sub>460</sub>/min under the conditions of the assay. The results were presented as U per mg of protein.

**Polyphenol oxidase (PPO) assay.** For the PPO assay, 100  $\mu$ L of extract were incubated with 2 mL 0.05 M phosphate buffer (pH 7.0) and 0.5 mL 0.5 M catechol at 24°C for 5 min, and absorbance at 398 nm was measured. The PPO activity was expressed as U, where U = 0.001  $\Delta$ OD<sub>398</sub>/min under the conditions of the assay (Galeazzi et al., 1981). The results were presented as U per mg of protein.

**Tyrosine ammonia-lyase (TAL) assay.** For the TAL assay, 100  $\mu$ L of the extract were incubated with 0.9 mL 0.02 M L-tyrosine at 30°C for 60 min. After incubations, 0.5 mL 10% trichloroacetic acid (TCA) was added to stop the reaction, samples were centrifuged (15 000  $\times$  g, 10 min) and absorbance at 310 nm was measured. One unit was defined as the amount of enzyme that produced 1.0  $\mu$ g *p*-coumaric acid per min under the conditions of the assay. The results were presented as U per mg of protein (Assis et al., 2001).

**Phenylalanine ammonia-lyase (PAL) assay.** For the PAL assay, 300  $\mu$ L of the extract were incubated with 1.2 mL 0.02 M L-phenylalanine and 2 mL of the PAL extracting buffer at 30°C for 60 min. After incubations, 0.5 mL 10% TCA was added to stop the reaction, samples were centrifuged (15 000  $\times$  g, 10 min) and absorbance at 290 nm was measured. One unit

was defined as the amount of enzyme that produced 1.0 µg trans-cinnamic acid per min under the conditions of the assay. The results were presented as U per mg of protein (Assis et al., 2001).

### Protein assay

The proteins content was determined with the Bradford method (Bradford, 1976), using bovine serum albumin as the standard protein.

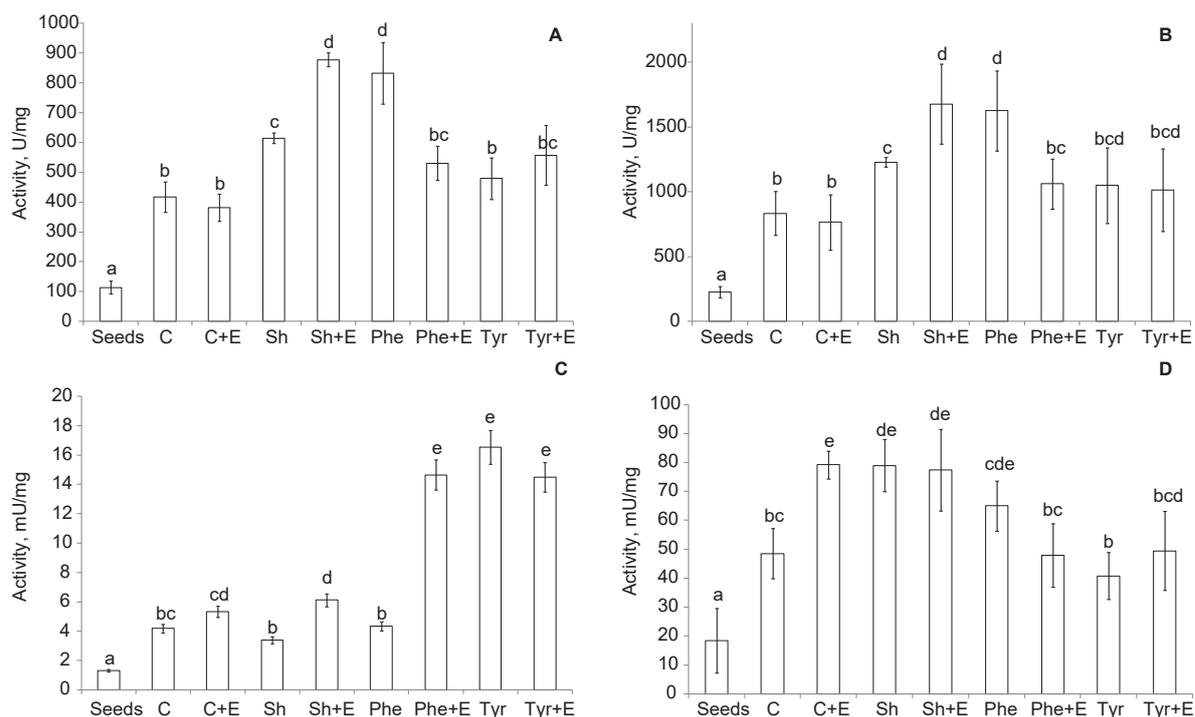
### Statistical analysis

All experimental results were mean ±S.D. of three independent experiments ( $n = 9$ ). One-way analysis of variance (ANOVA) and Turkey's post-hoc test were used to compare groups (seeds, as well as control and elicited sprouts) (STATISTICA 6, StatSoft, Inc., Tulsa, USA). Differences were considered significant at  $p < 0.05$ . Data

was evaluated using Pearson's correlation coefficients to identify relationships between selected features.

## RESULTS

The main antioxidants present in buckwheat sprouts are phenolics, in particular flavonoids. These compounds are known to be synthesized in the phenylpropanoid pathway and play a pivotal role in the plant's resistance against pathogens, radiation and environmental stresses. In Figure 1, the activities of enzymes involved in phenolic synthesis and metabolism are presented. The activity of peroxidase, an enzyme involved *inter alia* in lignin synthesis from phenolic constituents, was significantly increased by elicitation in the control sprouts (C+E), as well as those obtained from seed fed with shikimic acid (Sh and Sh+E). The



**Fig. 1.** Influence of elicitation combined with precursor feeding on the activities of enzymes involved in phenolic synthesis and metabolism: A – peroxidase, B – polyphenols oxidase, C – L-tyrosine ammonia-lyase, D – L-phenylalanine ammonia-lyase; S – seeds, C – control, C+E – elicitation, Sh – shikimic acid feeding, Sh+E – elicitation and shikimic acid feeding, Phe – phenylalanine feeding, Phe+E – elicitation and phenylalanine feeding, Tyr – tyrosine feeding, Tyr+E – elicitation and tyrosine feeding. Means in columns followed by different letters are significantly different at  $p = 0.05$ . Each value represents the mean of three independent experiments ( $\pm$ SD)

activity of polyphenols oxidase was increased in all the sprouts fed with the phenylpropanoids precursors; however, the changes were statistically significant ( $p < 0.05$ ) only in the case of Sh, Sh+E and Phe sprouts – elevations of about 47%, 101% and 95%, respectively (compared to the control). The activities of L-tyrosine ammonia-lyase (TAL) and L-phenylalanine ammonia-lyase (PAL), two enzymes directly involved in phenolic synthesis, were also diversified by the applied modification of sprouting modification. In respect to the control, TAL activity was most effectively induced by seeds soaking with tyrosine – for Tyr and Tyr+E about 4-fold and 3.5-fold, respectively. A significant increase was also determined for elicited sprouts obtained from seeds fed with phenylalanine (Phe+E). The activity of PAL was the highest in the elicited control sprouts (C+E) and sprouts obtained from seeds fed

with shikimic acid (Sh and Sh-E), where an increase of about 60% in respect to control was determined.

Germination significantly ( $p < 0.05$ ) increased the phenolic content – an about 2-fold was observed after 3 days of germination (compared to the seeds). The highest elevation was observed in the sprouts obtained from seeds enriched with shikimic acid (Sh and Sh+E) – total phenolics (by about 10% and 20%, respectively), condensed tannins (by about 30% and 28%, respectively), and flavonoids (by about 46% and 70%, respectively). An increased content of total phenolics and condensed tannins was also determined for sprouts obtained from seeds fed with phenylalanine (Phe) (Table 1). Qualitative–quantitative analysis of main sprout phenolics is presented in Figure 2. Generally, the rutin and chlorogenic acid content were comparable in all the sprouts except for the elicited sprouts obtained from seeds fed with shikimic acid, where an increase was determined. These sprouts were also characterized by significantly ( $p < 0.05$ ) increased contents of vitexin and isoorientin. Most importantly, feeding with the phenylpropanoid precursor caused a significant ( $p < 0.05$ ), about 2-fold, decrease of orientin content. Except for the Tyr and Tyr-E sprouts, the applied modification of sprouting decreased also the (+)-catechin content (Fig. 2). Analysis of the relationships between main flavonoids content and the activities of enzymes involved in phenolics synthesis, as well as their metabolism showed that both enzymes of the phenylpropanoids pathway (PAL and TAL) take a part in the synthesis of flavonoids (Table 2). Pearson correlation coefficients indicate that peroxidase and polyphenols oxidase (enzymes “consuming” phenolics) influenced negatively only orientin ( $R = -0.64$  and  $-0.65$ , respectively) and (+)-catechin ( $R = -0.49$  and  $-0.49$ , respectively) levels.

Though phenolics are the main antioxidants of buckwheat sprouts, the influence of the used treatments on the content of other low-molecular antioxidants, such as ascorbic acid and carotenoids, was also studied (Table 3). Germination significantly ( $p < 0.05$ ) increased the vitamin C and carotenoids content of sprouts. In the sprouts obtained from seeds fed with the studied precursors of phenolics phenolic the amount of ascorbic acid was significantly higher than that determined for the control sprouts (C and C+E). Most importantly, vitamin was generally present in a reduced

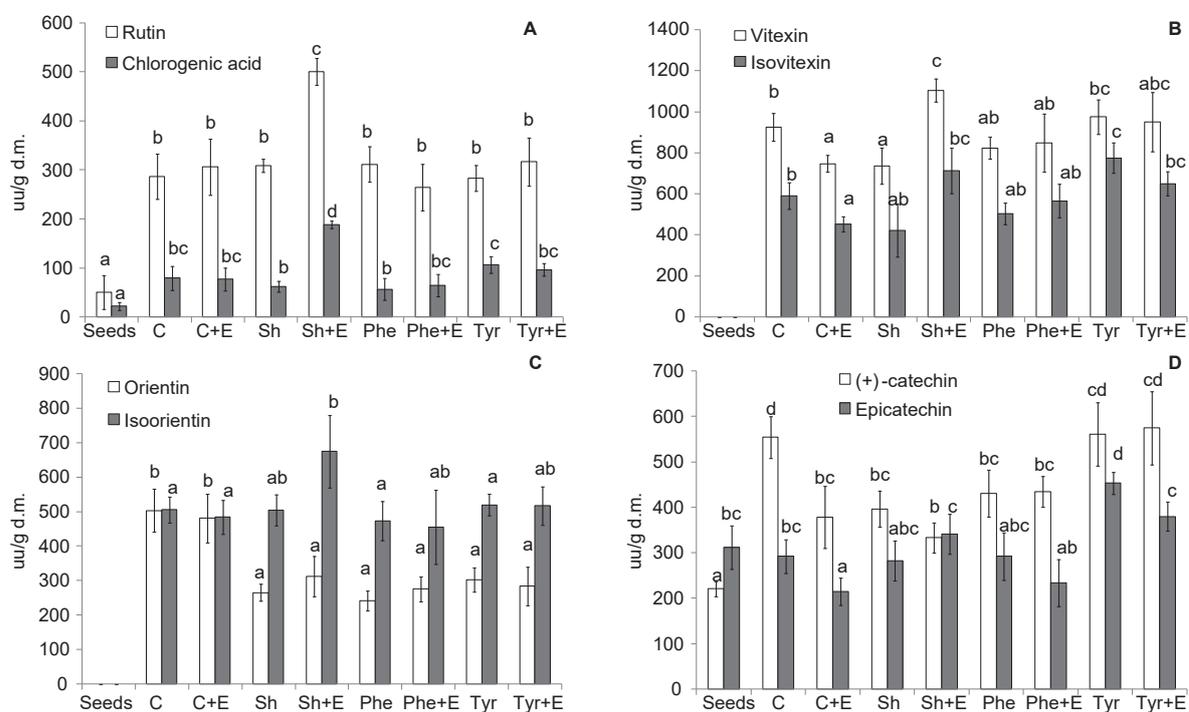
**Table 1.** Phenolic content in buckwheat sprouts – the effect of elicitation and elicitation supported by the phenylpropanoid pathway feeding

	Total phenolics mg/g d.m.	Condensed tanins mg/g d.m.	Total flavonoids mg/g d.m.
Seeds	22.83 ±3.07a	2.70 ±0.05a	1.11 ±0.36a
C	37.68 ±1.36b	7.25 ±0.33b	5.31 ±0.33b
C+E	37.99 ±2.14bc	7.25 ±0.16b	5.41 ±0.14b
Sh	41.77 ±1.08bc	9.47 ±0.19d	7.78 ±0.43c
Sh+E	45.56 ±1.25d	9.35 ±0.49cd	9.06 ±0.65d
Phe	44.45 ±1.71cd	9.12 ±0.22cd	5.97 ±0.19b
Phe+E	39.39 ±2.03bc	7.59 ±0.58bcd	5.75 ±0.18b
Tyr	38.68 ±1.29b	7.66 ±0.48bc	5.30 ±0.16b
Tyr+E	39.60 ±0.59b	8.97 ±0.20c	5.75 ±0.29b

S – seeds, C – control, C+E – elicitation, Sh – shikimic acid feeding, Sh+E – elicitation and shikimic acid feeding, Tyr – tyrosine feeding, Tyr+E – elicitation and tyrosine feeding, Phe – phenylalanine feeding, Phe+E – elicitation and phenylalanine feeding.

Means in columns followed by different letters are significantly different at  $p = 0.05$ .

Each value represents the mean of three independent experiments (±SD).



**Fig. 2.** Influence of elicitation combined with precursor feeding on the content of main phenolics in buckwheat sprouts: A – peroxidase, B – polyphenols oxidase, C – L-tyrosine ammonia-lyase, D – L-phenylalanine ammonia-lyase; S – seeds, C – control, C+E – elicitation, Sh – shikimic acid feeding, Sh+E – elicitation and shikimic acid feeding, Phe – phenylalanine feeding, Phe+E – elicitation and phenylalanine feeding, Tyr – tyrosine feeding, Tyr+E – elicitation and tyrosine feeding. Means in columns followed by different letters are significantly different at  $p = 0.05$ . Each value represents the mean of three independent experiments ( $\pm$ SD)

**Table 2.** Relationships between main flavonoids content and the activities of enzymes involved in phenolics synthesis, as well as their metabolism (Pearson correlation coefficients)

	Phenylalanine ammonia-lyase	Tyrosine ammonia-lyase	Polyphenols oxidase	Peroxidase
Isoorientin	0.33	-0.13	0.52	0.54
Orientin	0.05	-0.34	-0.65	-0.64
Isovitexin	-0.57	0.64	0.13	0.11
Vitexin	-0.31	0.34	0.34	0.36
Rutin	0.53	-0.28	0.66	0.70
(+)-catechin	-0.86	0.54	-0.49	-0.49
Epicatechin	-0.49	0.53	0.17	0.13

form (dehydroascorbic acid level did not exceed 5% of total vitamin C). Generally, the used modification of sprouting had only a slight effect on the carotenoids content – the changes did not exceed 10% (Table 3).

In plant-origin foods, the increase of low-molecular antioxidants usually improves the antioxidant capacity. Antiradical ability and the ability to inhibit lipids peroxidation were most effectively improved in the sprouts obtained from seeds fed with shikimic acid (Sh and Sh+E), where compared to the control sprouts an elevation of about 14% and 11% was observed for both variants of sprouting, respectively (Table 4). The highest reducing potential was determined for Sh+E and Phe sprouts – an increase by about 8% and 7%, respectively. The used modification of sprouting, except C+E, did not affect the chelating power of sprouts. It was observed that the total antioxidant capacity

**Table 3.** Effect of elicitation and elicitation supported by the phenylpropanoid pathway feeding on vitamin C and carotenoids contents

	Vitamin C µg/g f.m.	Ascorbic acid µg/g f.m.	Dehydroascorbic acid % of vitamin C	Carotenoids mg/100 g d.m.
Seeds	2.83 ±1.4a	1.61 ±0.88a	43.2	2.33 ±0.72a
C	65.61 ±0.63b	63.49 ±3.98b	3.23	28.51 ±3.16cd
C+E	67.12 ±1.50b	61.69 ±8.19b	8.10	25.97 ±1.20c
Sh	88.81 ±11.05c	86.15 ±13.05c	3.00	30.63 ±1.63e
Sh+E	81.42 ±2.00c	79.35 ±2.87c	2.54	27.17 ±0.95cd
Phe	85.78 ±3.56c	83.10 ±5.96c	3.12	20.67 ±0.98b
Phe+E	78.06 ±2.48c	76.81 ±1.66c	1.59	27.68 ±0.99d
Tyr	80.95 ±0.66c	79.96 ±1.01c	1.23	27.50 ±1.43cd
Tyr+E	80.77 ±2.00c	76.98 ±2.69c	4.69	26.23 ±0.82c

S – seeds, C – control, C+E – elicitation, Sh – shikimic acid feeding, Sh+E – elicitation and shikimic acid feeding, Tyr – tyrosine feeding, Tyr+E – elicitation and tyrosine feeding, Phe – phenylalanine feeding, Phe+E – elicitation and phenylalanine feeding.

Means in columns followed by different letters are significantly different at  $p = 0.05$ .

Each value represents the mean of three independent experiments ( $\pm$ SD).

**Table 4.** Effect of elicitation and elicitation supported by the phenylpropanoid pathway feeding on the antioxidant capacity of buckwheat sprouts

	Antiradical ability mg TE/g d.m.	Reducing power mg TE/g d.m.	Inhibition of lipids peroxidation mg TE/g d.m.	Chelating power mg EDTA/g d.m.
Seeds	4.94 ±0.68a	12.32 ±0.86a	8.99 ±0.45b	6.24 ±0.25a
C	12.99 ±1.01b	43.74 ±3.06bc	11.15 ±0.28d	8.11 ±0.07c
C+E	12.93 ±0.68b	43.78 ±1.49b	10.04 ±0.13c	7.75 ±0.11b
Sh	14.96 ±0.54c	44.54 ±5.79bc	12.39 ±0.39e	8.26 ±0.09c
Sh+E	14.78 ±0.47c	47.30 ±1.80c	12.45 ±0.43e	8.20 ±0.05cd
Phe	14.64 ±0.67bc	46.77 ±1.93c	9.55 ±1.14bcd	8.16 ±0.02c
Phe+E	13.45 ±0.38b	41.46 ±3.89bc	7.30 ±0.10a	8.12 ±0.11cd
Tyr	13.18 ±0.34b	41.08 ±1.96b	9.79 ±1.02bcd	8.31 ±0.05d
Tyr+E	13.44 ±0.92bc	42.92 ±4.77bc	12.39 ±0.25e	8.06 ±0.08c

S – seeds, C – control, C+E – elicitation, Sh – shikimic acid feeding, Sh+E – elicitation and shikimic acid feeding, Tyr – tyrosine feeding, Tyr+E – elicitation and tyrosine feeding, Phe – phenylalanine feeding, Phe+E – elicitation and phenylalanine feeding. TE – Trolox equivalent, EDTA – ethylenediaminetetraacetic acid.

Means in columns followed by different letters are significantly different at  $p = 0.05$ .

Each value represents the mean of three independent experiments ( $\pm$ SD).

(according to four complementary methods) of sprouts was most effectively increased by feeding seeds with shikimic acid and further elicitation of sprouts during germination (Table 4).

## DISCUSSION

Modifications of sprouting aimed at overproducing antioxidants usually involve induction of the natural mechanisms of plant resistance (Gawlik-Dziki et al., 2013; Pérez-Balibrea et al., 2011; Świeca et al., 2012; Tsurunaga et al., 2013); however, there are only a few studies concerning the mechanisms of obtaining desirable features. Elicitations usually induce a plant response cascade including *inter alia* changes in the activities of some enzymes, among which peroxidase, polyphenol oxidase and phenylalanine/tyrosine ammonia lyases should be mentioned. Peroxidase oxidizes phenolics (including flavonoids), whereby ascorbate accepts an electron from phenoxyl or flavonoid radicals. Ascorbate thereby converts to form a monodehydroascorbate radical, which subsequently can become converted to dehydroascorbate. Polyphenol oxidase catalyzes the oxidation of various phenolic substrates, whose polymerization leads to the formation of brown pigments or cell wall structures (Müller et al., 2014). Unfortunately (elicitation by producing oxidative stress) induces the activity of these enzymes, which may negatively influence antioxidant capacity by a reduction of phenolics and ascorbic acid contents (Tomás-Barberán and Espín, 2001; Van Doorn and Ketsa, 2014). Most importantly, in this study the activities of PPO and POD were increased only in the Sh, Sh+E and Phe sprouts, but according to the results (the dehydroascorbic acid level) elicitation did not affect redox status of sprouts.

In response to stress, plants overproduce *inter alia* phenolic compounds acting as antioxidants, signal compounds and/or substrates for physical barriers synthesis. Phenolics are mainly produced in the phenylpropanoid pathway, thus the induction of this metabolic pathway with elicitors seems to be an effective and simple tool for improving their level. So far, polyphenol synthesis was effectively increased by elicitation in both cell cultures (Matkowski, 2008; Zhao et al., 2014), as well as in low-processed food (Baenas et al., 2014). The effect of UV radiation (wavelengths 260–320 nm, 280–320 nm,

and 300–320 nm) on the production of flavonoid compounds in buckwheat sprouts was studied by Tsurunaga et al. (2013). The results obtained in the cited studies show that UV-B (>300 nm) irradiation is effective in the production of buckwheat sprouts with increased anthocyanins, rutin and DPPH radical scavenging activity. Contrary to this study, the increase in the ability to quench free radicals was significant (by about 35%). It may be explained by the fact that metabolism of 7-day-old buckwheat seedlings is strictly directed on flavonoid production (the effects of elicitation are clearly visible). The amount of phenolic compounds and carotenoids in the buckwheat sprouts was also improved by treatment with various concentrations of NaCl (10, 50, 100, and 200 mM) (Lim et al., 2012). In case of 3-day-old sprouts the most effective was treatment with 100 mM NaCl (which effected an almost 100% increase). The effects of elicitation with methyl jasmonate (MeJA) on phytochemical production in buckwheat sprouts cultivated under dark conditions (0, 1, 3, 5, and 7 d) were also studied (Kim et al., 2011). Similarly to this study, the cited investigators found chlorogenic acid, catechin, isoorientin, orientin, rutin, vitexin and quercitrin as the main phenolics of sprouts; however, an increase of total phenolic content in elicited sprouts was higher. The pattern of antioxidant activity reflected the changes in total phenolic content – after 3 days of sprouting differences could be observed between the antioxidant activities of the control and treated sprouts (Kim et al., 2011). These observations fully confirm the results obtained in this study. The accumulation of phenolic compounds resulted from the stimulation of the phenylpropanoid pathway (activity of PAL and TAL; Table 2, Fig. 1). An increase of enzymes involved in phenolic synthesis was also observed in sweet basil after treatment with methyl jasmonate, spermine and epibrassinolide, and in radish sprout after spraying with exogenous methyl jasmonate (Kim et al., 2006). Most importantly, in buckwheat sprouts total phenolic content is strongly, positively correlated with PAL activity (Ren and Sun, 2014). A significant increase of PAL but also TAL activity was also observed in this study after elicitation with *Salix* bark infusion; however, the effect was strongly dependent on precursor feeding, which may indicate that under stress conditions phenolic synthesis is mainly limited by the availability of their precursor. This hypothesis seems to be confirmed by other studies, e.g., improved

accumulation of phenylethanoid glycosides in suspension culture of *Cistanche salsa* (Liu et al., 2007), naringenin and phenylalanine feeding in anthocyanins synthesis in *Rudbeckia* (Luczkiewicz and Cisowski, 2001) or an effect of growth regulators and phenylalanine on phenolic compounds in sweet basil (Koca and Karaman, 2015). In the study of Świeca et al. (2014c), supplementation of seeds with shikimic acid, phenylalanine and tyrosine (combined with UV-B treatment) for improving the phenolic content and antioxidant activity of lentil sprouts was successfully applied. This strategy has also been employed for broccoli sprouts; however, feeding with methionine and tryptophan (precursors of aliphatic and indolic glucosinolates) did not result in any significant elevation of the glucosinolates content (Pérez-Balibrea et al., 2011).

## CONCLUSION

In this study buckwheat sprouts were used as a biotechnological module. Elicitation with infusion of willow bark supported by feeding with the phenylpropanoid pathway precursors is a metabolic engineering tool aimed to improve the functional quality of the sprouts – namely, an overproduction of low-molecular antioxidants. Metabolomic and biochemical analysis provides information concerning the mechanisms of desirable features gaining. Without any undesirable change in the metabolism of sprouts (redox potential, activities of polyphenol oxidase and peroxidase), the applied treatments significantly increased the low-molecular weight antioxidants content and affected the antioxidant capacity of the sprouts. This study revealed that the accumulation of these compounds was caused by the stimulation of two main enzymes of the phenylpropanoid pathway (tyrosine ammonia-lyase and phenylalanine ammonia-lyase). Consequently, the obtained results could be used to better understand the effects of elicitation and precursor feeding on the production of pro-health phytochemicals in buckwheat sprout, as well as to define how this technology improves their functional quality.

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