



## Concentration of fungal metabolites, phenolic acids and metals in mixtures of cereals grown in organic and conventional farms

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**ABSTRACT.** In view of expanding cultivation of cereal mixtures, the study was conducted to examine the effects of organic (ORG) vs conventional (CON) farming on fungal contamination, concentrations of phenolic acids (PA) and metals in mixtures of cereals grown in northern and central regions of Poland. The analyses were performed on 10 ORG and 8 CON bulk samples of oat, wheat and barley mixtures, each one comprised samples taken from 4–5 farms. Fungal contamination was assessed as the concentration of ergosterol (ERG) and mycotoxins from trichothecene (TR) group, whereas the quantification of fungal species and TR genotypes was performed using TaqMan assay. Mean ERG content in grain produced in both systems was similar, although a markedly greater variation was recorded in ORG grain. Total TR mycotoxins concentration was, in both CON and ORG samples, low and comparable, however concentration of deoxynivalenol was considerably greater whereas that of 3-acetyl-deoxynivalenol (3-AcDON) and 15-AcDON and nivalenol were lower in ORG samples. Molecular analysis showed that the dominant fungi were *Fusarium culmorum* and *F. graminearum*. The *Tri5* gene, a precursor of TR formation, was detected in significantly greater relative amounts in ORG samples. The ORG cereal mixtures contained greater total amounts of PA and, in particular, of gallic, p-cumaric and ferulic acids. Concentrations of Cd, Cu and Mg did not differ between the ORG and CON cereals, whereas concentrations of Fe, Se and Zn were greater in CON, and those of Mn and Pb in ORG mixtures. The potentially hazardous substances were present in the samples of both CON and ORG cereal mixtures in concentrations which do not compromise the health of farm animals.

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### Introduction

Organic (ORG) farming is an alternative to the conventional (CON) cultivation system providing farm products of high quality, referred to as organic (Maeder et al., 2002). The ORG farming in Poland

is developing, as evidenced by the constantly growing number and acreage of ORG farms (Zdrojewska, 2013).

Cultivation of cereals in mixtures aims at the reduction of negative effects of pure stands, such as yield fluctuations, due to different adverse envi-

ronmental conditions (Leszczyńska, 2007; Tratwal and Walczak, 2010). The world production of cereal mixtures is approximated for 4.8 million tons per year (FAO, 2009), with Poland being a dominant producer. In 2012, the area cropped to cereal mixtures in Poland was 1.45 million hectares, with north-eastern Poland being the leading region, followed by the central and eastern parts of the country (according to Agricultural Market Agency – ARR, Poland, 2013).

The most common cereal mixtures include barley and oat sown at a 1:1 ratio, or barley, wheat and oat sown at a 1:1:1 ratio. Proportions of harvested grain may differ since growth conditions adverse for one of the components, may be advantageous for another, which gains dominance in the stand and compensates for the potentially lower yield (Sobkowicz and Lejman, 2011). After harvesting, quality of grain in mixtures should be thoroughly monitored, since susceptibility to infestation the components by different pathogens is variable (Binder et al., 2007).

Microorganisms commonly found in grain include filamentous fungi, primarily those from the genera *Aspergillus*, *Fusarium* and *Penicillium*, responsible for the accumulation of mycotoxins in grain (Pasanen et al., 1999). Contamination of feeds with metabolites of filamentous fungi constitutes a serious health hazard, both for animals and humans. The trichothecenes (TR) are a group of mycotoxins produced by a variety of different *Fusarium* species. In Poland, the most common toxins are those from the group B of TR, e.g., deoxynivalenol (DON) and nivalenol (NIV) (Perkowski et al., 2012). The content of fungal toxins in grain is inseparably connected with the presence of mycobiota. In addition, content of ergosterol (ERG) is the marker of the presence of both live and dead mycobiota (Zhao et al., 2005).

Due to the fact that in the Poland's climatic zone the most toxigenic fungi are *Fusarium culmorum* and *F. graminearum*, the presence of these fungi and also of chemotypes 3-acetyl-deoxynivalenol (3-AcDON) and 15-acetyl-deoxynivalenol (15-AcDON) in grain is increasingly often analysed. Surveys from diverse geographic areas have revealed that distribution of chemotypes among local populations of *F. graminearum*/*F. culmorum* varies and may change over time (Waalwijk et al., 2004). Rapid changes in the proportions of chemotypes within *F. graminearum*/*F. culmorum* are difficult to predict because inoculum dispersal and development in this field is affected by complex factors. For this reason, quick identification of chemotypes is imperative in order to determine the potential risk to human and animal health (Pasquali et al., 2010).

As the marker of the potential capacity for TR production, the presence of the *Tri5* gene, encoding trichodiene synthase involved in the first stage of the TR biosynthesis pathway, is assayed (Vegi and Wolf-Hall, 2013).

Phenolic compounds and certain microelements, due to their antioxidant activity, significantly stimulate the plant resistance against fungal pathogens. The elevated concentration of ferulic acid, the most abundant acid among phenolic acids present in cereal grain, may indicate a more active plant response to the environmental stress (Lempereur et al., 1997). Selenium is a major microelement exhibiting antioxidant activity (Hartikainen and Xue, 1999). Also iron, magnesium, zinc, copper and manganese and trace elements like cadmium, lead and mercury are considered as important minerals affecting both the plant growth and grain nutritional value (Kawecka et al., 2013).

The aim of the study was to compare the fungi contamination levels in mixtures of cereals grown in ORG and CON farms, using the concentrations of ERG and mycotoxins from the TR group as a criteria. The results referred to molecular analyses in order to indicate dominant fungi found most commonly in the culture environment, and to determine the presence of the *Tri5* gene responsible for the formation of TR in grain. The phenolic acids and selected metals contents were also determined.

## Material and methods

### Material

Cereal mixtures used in the study were grown in 2011 in the conventional and organic farms situated in ten communes of the two voivodeships in central (Kuyavian-Pomeranian and Greater Poland) and one in northern Poland (Warmian-Masurian Voivodeship) (Table 1).

The mixtures were composed of oat, wheat and barley sown in proportion 1:1:1, grown under monitored temperature and precipitation conditions (Table 2), harvested at full ripeness using combine harvester and stored in grain silos. The analyses were performed on the bulk samples: 8 representing conventional and 10 organic farming. Each bulk

**Table 1.** Voivodeships and number of samples

Voivodeship	Conventional farming		Organic farming	
	bulk	individual	bulk	individual
Kuyavian-Pomeranian	2	9	2	9
Warmian-Masurian	3	13	4	18
Greater Poland	3	12	4	18

**Table 2.** Values of mean temperature (° C) and precipitation totals (mm) in 2011 summer season (according to the Institute of Meteorology and Water Management – National Research Institute (Poland): <http://www.imgw.pl/klimat/>)

Month	Voivodeship		
	Kuyavian-Pomeranian	Warmian-Masurian	Greater Poland
	mean temperature, ° C // precipitation totals, mm		
May	14 // 30	13 // 50	14 // 30
June	18 // 80	17 // 60	18 // 70
July	17 // 140	17 // 180	17 // 140
August	18 // 40	17 // 60	18 // 40

sample was made of individual samples collected in 4–5 respective farms. The bulk samples were prepared in the laboratory, stored at –20 °C till botanical and chemical analysis.

### Botanical analysis of mixtures

The analysis of botanical composition was performed in three replications on 100 g of bulk samples. Wheat, barley and oat grains were separated by hand, weighed and their proportions in the harvested mixture were calculated.

### Chemical analysis

**Dry matter** of the grain was determined using the AOAC International (2002) method No. 2001.12 for determination of water/dry matter (moisture) in animal feed, grain and forage (plant tissue). The results were expressed in relation to the dry matter content.

**Ergosterol (ERG) concentration** was determined after saponification and microwave extraction. After saponification ERG was extracted using pentane and evaporated to dryness in a gentle stream of a high purity nitrogen. Samples were dissolved in methanol and analysed using a high-performance liquid chromatography (Alliance 2695) with an absorptiometric diode detector (Photodiode Array Detector 2996). Chromatographic separation was performed on a Nova-Pak C18 column (150 mm × 3.9 mm). This analysis was performed on the equipment provided by Waters Corporation (Milford, MA, USA). Methanol and acetonitrile mixture at a 90:10 ratio (v/v) was used as an eluent. ERG concentration was measured using an external standard at wavelength  $\lambda = 282$  nm. The compound was identified by comparing the retention time of the peak with the original standard and by adding a specific amount of the standard to the sample. The analysis was repeated. The limit of detection was 0.01 mg · kg<sup>-1</sup> (Perkowski et al., 2008).

**Contents of trichothecenes from group B** were determined following the extraction of the compounds with a acetonitrile-water mixture at a 82:18 ratio (v/v) and purification by solid-phase extraction on a charcoal column Mycosep® 225 Trich (Romer Labs, Getzersdorf, Austria). The TR from group B were analysed as trimethylsilyl derivatives. Chromatographic separation and detection were conducted on equipment provided by Hewlett Packard (Waldbronn, Germany) using a gas chromatograph (GC 6890) equipped with a mass detector (MS 5972 A) and capillary column (HP-5MS, 0.25 mm × 30 m). Analysis was performed in the selective ion monitoring mode (SIM). The selective ions for each examined compound were: deoxynivalenol (DON) 103 and 512, 3-acetyl-deoxynivalenol (3-AcDON) 117 and 482, 15-acetyl-deoxynivalenol (15-AcDON) 193 and 482, fusarenone X (FUS-X) 103 and 570, and for nivalenol (NIV) 191 and 600. Helium flow rate was 0.7 cm<sup>3</sup> · min<sup>-1</sup>. In order to confirm the presence of investigated toxins in the sample, the analysis was performed over the entire mass range (100–700 unified atomic mass unit). Results were processed in the MS Chem Station programme (HP G1034A; Hewlett Packard, Waldbronn, Germany). Recovery for the analysed toxins was: DON 84 ± 3.8%, 3AcDON 78 ± 4.8%, 15AcDON 74 ± 2.2%, and NIV 81 ± 3.8%. The detection limit for the analysed toxins was 0.001 mg · kg<sup>-1</sup> (Perkowski et al., 2003).

**Phenolic acids** were determined after basic and than acid hydrolysis. Ground grain (0.2 g) was hydrolysed using 1 ml of H<sub>2</sub>O and 4 ml of 2 M NaOH followed by acid hydrolysis in 3 ml of 6 M HCl. Phenolic acids were extracted from the inorganic phase using diethyl ether. Extracts were evaporated to dryness in a gentle stream of a high purity nitrogen. Contents of phenolic acids were determined using a high-performance liquid chromatography (Alliance 2695) with a photodiode array detector (Photodiode Array Detector 2996) (both from Waters Corporation, Milford, MA, USA). Chromatographic separation was performed on a RP C-18 column (250 mm × 4 mm × 5 µm; Waters, Ireland). A mixture of acetonitrile: 2% acetic acid in water (pH = 2) was applied as the eluent (gradient). Measurements of phenolic acid concentrations were performed using an external standard at wavelengths  $\lambda = 320$  nm and  $\lambda = 280$  nm. Compounds were identified on the basis of the comparison of retention times of the peaks with those of the standard and by adding a specific amount of the standard to the tested sample. The analyses were repeated. Retention times of analysed acids were as follows, min: gallic 8.85,

4-hydroxybenzoic 19.46, vanilic 24.11, caffeic 26.19, syringic 28.05, p-cumaric 40.20, synapic 48.00. Limit of detection was  $1 \text{ mg} \cdot \text{kg}^{-1}$ .

**Metal analysis.** The material was mineralized with a CEM Mars 5 Xpress (CEM, Matthews, NC, USA) microwave mineralization system (55 ml vessels) using 8 ml  $\text{HNO}_3$  (65%) and 2 ml  $\text{H}_2\text{O}_2$ , according to the programme comprising three stages: first stage – power 800 W, time 10 min, temperature  $120^\circ\text{C}$ ; second stage – power 1600 W, time 10 min, temperature  $160^\circ\text{C}$ ; third stage – power 1600 W, time 10 min, temperature  $200^\circ\text{C}$ . Materials after digestion were filtered through 45 mm filters (Qualitative Filter Papers Whatman, Grade 595:  $4-7 \mu\text{m}$ ; GE Healthcare, Buckinghamshire, UK), and filtrate completed with deionized water from Milli-Q Academic System (non-TOC (Total Organic Carbon); Millipores. A.S., Molsheim, France) to a final volume of 50 ml. Concentration of particular trace elements was analysed by the flame atomic absorption spectrometry (Cd, Cu, Fe, Mn, Pb and Zn), atomic emission spectrometry (Mg) and hydride generation atomic absorption spectrometry (Se), using an AA Duo – AA280FS/AA280Z spectrometer (Agilent Technologies, Mulgrave, Victoria, Australia), equipped with a Varian hollow-cathode lamp (HCL; Varian, Mulgrave, Victoria, Australia). Calibration curves were prepared in four replicates per each trace element concentration. Detection limit for the analysed metals was,  $\text{ng} \cdot \text{kg}^{-1}$ : Cd 0.01, Cu 0.18, Fe 0.11, Mg 0.003, Mn 0.005, Pb 0.14, Zn 0.06, Se 0.21 (Mleczek et al., 2010).

### Molecular analysis

**DNA extraction.** Grain samples (0.25 g) were homogenized (30 s at speed  $6.0 \text{ m} \cdot \text{s}^{-1}$ ) on a Fast-Prep-24 instrument in tubes with 1 mm silica spheres (Lysing matrix C) both provided by MP Biomedicals (Solon, OH, USA). Total DNA was extracted using a ChargeSwitch® gDNA Plant Kit following the manufacturer's recommendations (Invitrogen, Carlsbad, CA, USA).

**TaqMan assays** were used for quantification of *F. culmorum*, *F. graminearum* (Waalwijk et al., 2004), the *Tri5* gene and 3AcDON, 15AcDON chemotypes (Kulik, 2011). A fungal DNA assay (Kulik, 2011) was used as an internal control in order to identify samples, for which the PCR or the DNA extraction failed. TaqMan reaction conditions were used for both species, TR genotypes and fungal internal positive control (IPC) in the modified fast PCR protocol:  $95^\circ\text{C}$  for 20 s and ( $95^\circ\text{C}$  for 3 s,  $60^\circ\text{C}$  for 30 s)  $\times 40$ . For quantification of the *Tri5* gene, a PCR protocol

was used:  $95^\circ\text{C}$  for 10 min and ( $95^\circ\text{C}$  for 15 s,  $60^\circ\text{C}$  for 1 min)  $\times 40$ . TaqMan PCR mixture was composed of: 2  $\mu\text{l}$  DNA, 13  $\mu\text{l}$   $\text{H}_2\text{O}$ , 5  $\mu\text{l}$  TaqMan Fast Advanced Master Mix (Applied Biosystems, Foster City, CA, USA), 6 pM of each primer and 1.7 pM of probe. All PCR amplifications were carried out in a ViiA 7 Real-Time PCR System (Applied Biosystems, Foster City, CA, USA) in a final volume of 20  $\mu\text{l}$ . The threshold value was 0.1. To determine the efficiency and sensitivity of the assay, 21–36 ng of genomic DNA of *F. culmorum*/*F. graminearum* different isolates were serially diluted with water and used as a template. Amplification efficiencies for *F. culmorum*, *F. graminearum*, the *Tri5* gene, 3AcDON and 15AcDON assays were 97%, 101.2%, 93%, 101.9% and 99.3%, respectively. In order to eliminate false-positive results, a PCR reaction was considered positive only if the cycle threshold (Ct) value was  $<35$ .

### Statistical analysis

Results of chemical analyses were subjected to a statistical analysis with the use of STATISTICA 8.0 software, 2014 (StatSoft Polska, Kraków, Poland). Contents of individual compounds in CON and ORG mixtures were compared using Tukey's multiple comparison procedure with significant differences at level  $P = 0.05$ . An analysis of variance and assessment of the homogeneity of variance were determined. The least significant differences were determined by post-hoc LSD test. Linear regression between oat and ergosterol contents was calculated by Least Square method.

### Results and discussion

Both mean temperature and precipitation values in May, June, July and August 2011 were very similar in the two central voivodeships (Kuyavian-Pomeranian and Greater Poland) whereas in the north-eastern Warmian-Masurian Voivodeship total precipitations and particularly precipitations in July, were greater (Table 2).

The proportions of three cereals in bulk samples differed from the sown ratio but were similar in CON and ORG mixtures. Proportions of oat varied from 26 to 33% and from 23 to 30%, of wheat from 34 to 43% and from 28 to 40%, and those of barley from 28 to 39% and from 31 to 40% in CON and ORG samples, respectively (Table 3).

The recorded ERG concentration was relatively low and did not differ significantly between mixtures from CON and ORG farms ( $8.77$  vs  $7.94 \text{ mg} \cdot \text{kg}^{-1}$ ; Table 4). However, whereas the ERG content in

**Table 3.** Botanical composition of cereal mixture, %

No. of sample	Conventional farming			No. of sample	Organic farming		
	oat	wheat	barley		oat	wheat	barley
1	30	34	36	9	32	28	40
2	33	35	32	10	30	39	31
3	29	40	31	11	29	32	39
4	31	38	31	12	32	37	31
5	31	35	34	13	27	40	33
6	26	40	34	14	30	36	34
7	27	34	39	15	25	38	37
8	29	43	28	16	28	37	35
				17	23	38	39
				18	31	35	34

**Table 4.** Mean concentrations of ergosterol (ERG) and trichothecenes in mixtures of cereals grown in conventional (CON) and organic (ORG) farms

Cultivation system	n	ERG, mg · kg <sup>-1</sup>	Trichothecenes <sup>1</sup> , µg · kg <sup>-1</sup>					Total
			DON	FUS-X	3-AcDON	15-AcDON	NIV	
CON	8	8.77	12.5 <sup>b</sup>	1.0	2.8 <sup>a</sup>	4.3 <sup>a</sup>	9.8 <sup>a</sup>	30.4
ORG	10	7.94	21.2 <sup>a</sup>	0.9	0.7 <sup>b</sup>	2.4 <sup>b</sup>	5.3 <sup>b</sup>	30.5

<sup>1</sup>DON – deoxynivalenol, FUS-X – fusarenone X, 3-AcDON – 3-acetyl-deoxynivalenol, 15-AcDON – 15-acetyl-deoxynivalenol, NIV – nivalenol; <sup>ab</sup> – means with different superscripts within column are significantly different at  $P < 0.05$

CON mixtures was fairly uniform, in ORG mixtures about tenfold differences between samples No. 10 and 17 vs 12 were found (Figure 1). The greater variability of ERG concentration in ORG mixtures could be the effect of a probably less uniform cultivation measures and, consequently, plants' growing conditions under this system. Since there is no data available on the fungal contamination of grain in cereal mixtures, the results of present study were compared with the reported contamination of respective cereals grown in pure stands. Stuper and Perkowski (2008) found that the colonization of barley, wheat and oat grain by mycobiota was variable. In their study the mean ERG content in wheat,

barley and oat was 4.13, 9.21 and 16.11 mg · kg<sup>-1</sup>, respectively. When these values were applied to the cereals present in the mixture in mean proportion of 37:34:29, respectively, the calculated approximate ERG concentration of 9.3 mg · kg<sup>-1</sup> is very close to the analysed values.

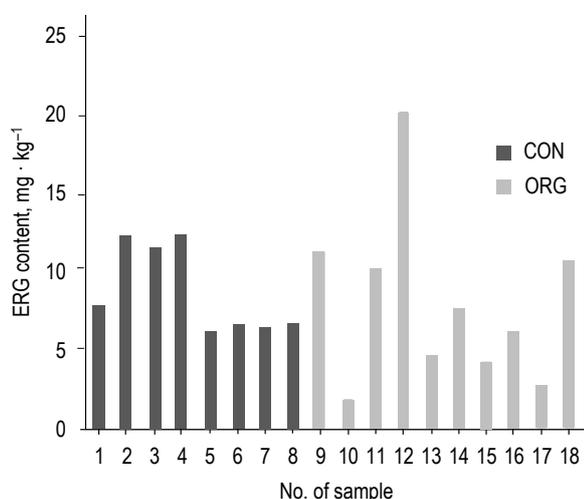
The correlation analysis between the ERG contents and proportions of wheat, barley and oat in the analysed mixtures showed that concentration of ERG is positively correlated with proportion of oat. The respective correlation formula is:

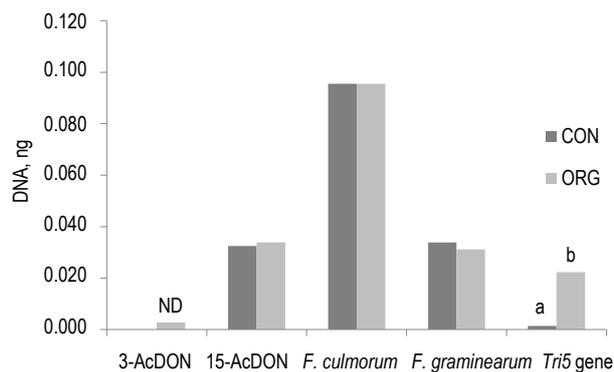
$$y = 1.113x - 24.04 \quad (r = 0.671, P = 0.05)$$

where:  $y$  – concentration of ERG (mg · kg<sup>-1</sup>),  $x$  – weight percent of oat in cereal mixtures (%).

This finding is in agreement with the results of Stuper and Perkowski (2008) and may indicate greater susceptibility of oat to fungal contamination and, in consequence, greater mycobiota concentration in the mixture with a higher proportion of oat.

The presence of mycotoxins in grain is strictly related to the content of toxigenic fungi in the agricultural environment. In Poland, the most common toxins are those from the TR group including DON and NIV (Perkowski et al., 2012). Wiśniewska and Buśko (2005) found also a strong and statistically significant correlation between DON and ERG content ( $r = 0.91$ ). Mixtures grown in the CON farms were less contaminated with DON but more contaminated with its derivatives 3-AcDON and 15-AcDON, and NIV (Table 4). Concentrations of fusarenone X (FUS-X) and total toxins did not differ between

**Figure 1.** Ergosterol (ERG) content in mixtures of cereals grown in conventional (CON) and organic (ORG) farms



**Figure 2.** Quantification of chemotypes: 3-acetyl-deoxynivalenol (3-AcDON) and 15-acetyl-deoxynivalenol (15-AcDON), fungi: *Fusarium culmorum* and *F. graminearum* and *Tri5* gene in grain of cereal mixtures from conventional (CON) and organic (ORG) cultivation systems (identification based on genetic analyses); <sup>ab</sup> – bars with different superscripts are significantly different at  $P < 0.05$  for each examined parameter separately; ND – not detected

mixtures grown under the two farming systems. Concentration of DON in the analysed samples was considerably lower than the highest admissible level set by the regulations of EU (EC 1881/2006) for  $1750 \mu\text{g} \cdot \text{kg}^{-1}$ . Moreover, the concentrations of toxins found in present study are lower than those reported in cereals grown in the same period in Europe. Šliková et al. (2013) found that the mean content of DON in wheat grain collected in 2011 was  $300 \mu\text{g} \cdot \text{kg}^{-1}$ . The similar concentration of TR was also found by Fredlund et al. (2013) in oat.

Following the analysis of individual toxins, the presence of the toxigenic fungi species and chemotypes, and of *Tri5* gene, was verified. The presence of chemotype 15-AcDON, reported in Poland for the first time in 1990 by Perkowski (1990), was found in all samples and did not differ between the ORG and CON mixtures (Figure 2). The 3-AcDON chemotype was present in minute amounts in ORG and was not detected in CON samples. The presence of *F. culmorum* was confirmed in all samples in concentrations similar for the two systems whereas *F. graminearum*

was detected almost in all samples and in smaller amounts than *F. culmorum*. The *Tri5* gene (ng DNA) was the only parameter which was affected by farming system and was greater in ORG than CON mixtures. This finding is in agreement with considerably greater DON concentration in ORG mixtures ( $21.2$  vs  $12.5 \mu\text{g} \cdot \text{kg}^{-1}$ ) but disagrees with smaller concentrations of other TR toxins. Up to our knowledge, no data on this topic is available in the literature and the effects of farming system on the concentrations of toxins and results of molecular analysis of respective fungi need more extensive studies.

Total phenolic acids concentration was considerably greater in grain from ORG than CON mixtures (Table 5). Among particular acids significantly greater were concentrations of gallic, p-cumaric and ferulic acids. Similar results were reported by Zuchowski et al. (2011) who found that concentrations of total and ferulic acids were greater in grain of spring wheat cultivated in ORG than in CON farms. Since phenolic acids are synthesized by plants in response to different types of stress (Robbins, 2003), their greater concentration may indicate that due to the lower mineral fertilization and lack of chemical protection under organic system, plants respond to the environmental stress in the enhanced manner.

The effects of farming system on the concentration of minerals were irregular (Table 6). Concentrations of iron, selenium and zinc were smaller and those of manganese and lead were greater in ORG than in CON mixtures whereas concentrations of cadmium, copper and magnesium did not differ significantly. The lower iron concentration in ORG than CON mixtures is in agreement with the results of Ryan et al. (2004) on the effect of farming system on the content of this element in wheat. In our study, the concentrations of minerals were either slightly lower (magnesium, zinc), greater (iron) or similar (manganese, copper) to those found by Suchowilska et al. (2012) in the conventionally cultivated wheat.

**Table 5.** Mean concentrations of phenolic acids in mixtures of cereals grown in conventional (CON) and organic (ORG) farms

Cultivation system	n	gallic	Phenolic acids, $\text{mg} \cdot \text{kg}^{-1}$								
			hydroxybenzoic	chlorogenic	caffeic	siringic	vanilin	p-cumaric	ferulic	synapic	total
CON	8	20 <sup>b</sup>	10	12	31	31	6	45 <sup>b</sup>	1191 <sup>b</sup>	20	1366 <sup>b</sup>
ORG	10	43 <sup>a</sup>	19	11	31	36	27	66 <sup>a</sup>	1989 <sup>a</sup>	27	2249 <sup>a</sup>

<sup>ab</sup> – means with different superscripts within column are significantly different at  $P < 0.05$

**Table 6.** Mean concentrations of mineral compounds in mixtures of cereals grown in conventional (CON) and organic (ORG) farms

Cultivation system	n	Mineral compounds, $\text{mg} \cdot \text{kg}^{-1}$							
		Cd	Cu	Fe	Mg	Mn	Pb	Se	Zn
CON	8	0.10	3.96	76.77 <sup>a</sup>	1261.00	29.69 <sup>b</sup>	0.034 <sup>b</sup>	0.28 <sup>a</sup>	27.48 <sup>a</sup>
ORG	10	0.09	5.16	54.63 <sup>b</sup>	1339.00	35.86 <sup>a</sup>	0.052 <sup>a</sup>	0.19 <sup>b</sup>	18.96 <sup>b</sup>

<sup>ab</sup> – means with different superscripts within column are significantly different at  $P < 0.05$

The cadmium concentration in the mixtures was high and not affected by cultivation system which is against the suggested effects of cadmium on the plant resistance to fungal diseases (Zhang et al., 2008). The high concentration of cadmium in grain, in spite of its generally low content in Polish soils, may be due to soil acidification and greater bio-availability of this element. Concentration of cadmium in both ORG and CON mixtures was reaching the upper limit amounting  $0.10 \text{ mg} \cdot \text{kg}^{-1}$  specified by EU (EC 1881/2006).

## Conclusions

Organic cultivation of mixture of wheat, barley and oat increases variability of ergosterol concentration, does not affect the total content but differently affects particular trichothecenes contents (increases deoxynivalenon), and increases *Tri5* gene. Ergosterol concentration in the mixture seems to be positively related to the proportion of oat. Organic cultivation stimulates the phenolic acids synthesis and has varied effect on concentration of minerals.

It may be concluded that the fungal contamination of the mixture of cereals cultivated accordingly to organic system is only slightly different from the contamination of the mixture of cereals grown conventionally and does not implicate a health hazard for animals.

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