

## ***Fusarium* species and *Fusarium* mycotoxins in grain of wheat in Poland in 2009 and 2010**

**Tomasz Góral<sup>1</sup>, Piotr Ochodźki<sup>1</sup>, Linda K. Nielsen<sup>2</sup>, Dorota Walentyn-Góral<sup>1</sup>**

<sup>1</sup>Department of Plant Pathology, Plant Breeding and Acclimatization Institute – National Research Institute, Radzików, 05-870 Błonie, Poland; t.goral@ihar.edu.pl;

<sup>2</sup>Sejlet Plant Breeding, Nørremarksvej 67, 8700 Horsens, Denmark

### **Abstract**

The aim of the study was to determine the presence *Fusarium* species and mycotoxins in winter wheat grain in Poland. Grain samples from different locations in Poland in 2009 and 2010 were analysed for the content of biomass of *Fusarium* species and mycotoxins. In 2009 biomass of *F. graminearum* and *F. poae* was present in all samples, *F. culmorum* in 82% of samples, *F. avenaceum* in 55% of samples. *F. sporotrichioides*, *F. tricinctum* and *F. equiseti* were found only in individual samples. *F. langsethiae* was not detected. In 2010, five *Fusarium* species were detected with the exception of *F. sporotrichioides*. The highest content of biomass was found for *F. graminearum* followed by *F. avenaceum*, *F. poae* and *F. langsethiae*. The amount of *F. culmorum* biomass was very low. The most frequently occurring species was *F. poae* and *F. graminearum*. In 2009, deoxynivalenol was detected in all samples. In 2010, the average content of deoxynivalenol was lower than in 2009. Nivalenol was detected at very low concentration in both years. Significant correlations between content of *F. graminearum* biomass and deoxynivalenol concentration in grain and between content of *F. poae* biomass and nivalenol concentration in grain in 2009 were found. The most important finding of this study was that main *Fusarium* species infecting wheat kernels in Poland in both years was *F. graminearum*. The amount of biomass of *F. graminearum* was the highest in both years. It was present in the most samples. The other frequently detected species was *F. poae*, which in 2010 appeared in more samples than *F. graminearum*. However, the amount of *F. poae* biomass was lower. *F. culmorum*, species that was previously dominating as wheat pathogen in Poland, was found less frequently than *F. graminearum*. The amount of biomass of this species was the lowest in 2010.

**Key words:** biomass, *Fusarium* head blight, real-time PCR, tricothecenes, zearalenone

### **Introduction**

*Fusarium* head blight (FHB) is a disease of wheat caused by a complex of toxicogenic fungi of the genus *Fusarium* (Parry *et al.*, 1995). The main species of this complex in Europe are *F. graminearum* and *F. culmorum* identified as deoxynivalenol (DON), nivalenol (NIV) and zearalenone (ZEN) producers. However, other *Fusarium* species producing mycotoxins are also prevalent: *F. avenaceum* - moniliformin, enniatins and beauvericin (BEA) producer, *F. poae* - NIV, BEA producer. *Fusarium langsethiae* and *F. sporotrichioides* - T-2 and HT-2 toxins producers are also prevalent (Bottalico, 1998; Bottalico and Perrone, 2002; Jestoi *et al.*, 2008; Somma *et al.*, 2010; Vogelgsang *et al.*, 2008). *Fusarium graminearum* and *F. culmorum* are highly pathogenic species, which can cause severe epidemics of *Fusarium* head blight. The other species are medium or weakly pathogenic, however, due to the wide prevalence, these may also cause mycotoxin contamination of grain (Nielsen *et al.*, 2011; Uhlig *et al.*, 2007; Yli-Mattila *et al.*, 2004, 2008). Because of the diversity of *Fusarium* species causing *Fusarium* head blight, monitoring of changes in the *Fusarium* population on wheat is important. Frequency of

species infecting wheat is not stable and changes depending on weather in particular year. Large differences are also observed between different regions of wheat productions in Europe. For example, the other species dominate in north-eastern Europe then in south-western part of the continent (Bottalico, 1998; Bottalico and Perrone, 2002). Species compositions changes over time, which is the results of climate warming, and changes in acreage of major cereal crops – increase of maize area. The main reported effect of the above factors is increase of *F. graminearum* and decrease of *F. culmorum* (Miller, 2008; Parikka *et al.*, 2012; Scherm *et al.*, 2013). Chandelier *et al.* (2011) analysed winter wheat samples from Belgium over 2003–2009 period. They found that main species were *F. avenaceum* and *F. graminearum*; however, their frequency changed depending on year from 20 to 100%. The frequency of *F. poae* was relatively constant over the years (about 70%). The overall incidence of *F. culmorum* decreased during the study, from 80% in 2003 to 10% over the final three years. Similarly, Isebaert *et al.* (2009) observed that *F. graminearum* and *F. culmorum* were the most important species in northern Belgium in 2002-2005. They found interesting correlation between crops and both species frequency (maize – *F. graminearum*, small grain cereals – *F. culmorum*). In Luxemburg, the most common species isolated from wheat heads were *F. graminearum*, *F. avenaceum* and *F. poae*. Increase of frequency of *F. graminearum* and decrease in *F. culmorum* were observed (Giraud *et al.*, 2010). Winter wheat cultivated in the Netherlands in 2009 was studied for *Fusarium* species and toxins (van der Fels-Klerx *et al.*, 2012). In samples collected on harvest, authors found dominance of *F. graminearum*. *F. avenaceum* and *Microdochium nivale* were also frequent. However, in the pre-harvest samples, only *F. graminearum* and *M. nivale* were present. Waalwijk *et al.* (2004) analysed wheat heads and grain collected in the Netherlands in 2001 and 2002. In 2001, in samples collected at late milk stage, *F. graminearum* was predominant; however, some samples contained also *F. avenaceum* and/or *F. culmorum*. At harvest, *F. graminearum* dominated almost completely. In 2002 the weather conditions were more favorable for FHB and they found relative dominance of *F. graminearum* in grain from the Netherlands and almost complete in samples from France. According to Birzele *et al.*, (2002) in 1997 and 1998 the dominating species in Germany in wheat grain were *F. avenaceum*, *F. poae*, *F. culmorum* and *F. graminearum*. Frequencies of two last species were similar, however percentage of *F. graminearum* increased in 1998. In Germany in 2008, *F. graminearum* sensu stricto was the predominant species followed by *F. culmorum*. Other species (*F. poae*, *F. tricinctum*, *M. nivale* etc.) were identified in small amounts (Talas *et al.*, 2011). In Hungary, in year 2010, which was very favorable for *Fusarium* head blight development, predominantly *F. graminearum* was isolated from wheat grain (Laszlo *et al.*, 2011).

Waalwijk *et al.* (2003) analysed wheat spikes with *Fusarium* head blight symptoms collected in Netherlands in 2000 and 2001. They found that *F. graminearum* was the dominating *Fusarium* species in both years. As they stated, this was significant change comparing results from the 1980s and 1990s, which showed that *F. culmorum* was the predominant species in the Netherlands. They presume that this shift could be connected with an increase in maize acreage. *F. graminearum*, unlike *F. culmorum*, is a major pathogen on maize and, can survive on maize debris. The other factor could be climate warming which favours *F. graminearum* as it has higher optimal temperature of development. The prevalence of FHB pathogens differed significantly between studied countries in 2001 and 2002 (UK, Ireland, Italy and Hungary) (Xu *et al.*, 2005). Overall, all pathogens (*F. graminearum*, *F. culmorum*, *F. avenaceum* and *F. poae*) were commonly detected in Ireland and to a lesser extent in the UK. In contrast, only two species, *F. graminearum* and *F. poae*, were regularly detected in Italy and Hungary. *Fusarium culmorum* was rarely detected except in Ireland. The later country is has the coolest summer weather among four studied countries. Authors stated that the increase of *F. graminearum*, especially in the UK, appears to have been at the expense of *F. culmorum*. The replacement of

*F. culmorum* by *F. graminearum* as the predominant FHB pathogen was also reported in Bavaria (Obst *et al.*, 1997) where the change was linked with increased maize production. It is worth to notice that in Poland grain maize acreage increased considerably from 1990 (59 000 ha) to 2017 (above 1 215 500 ha).

The *Fusarium* species can be isolated from cereal kernels and identified using classical and/or molecular methods (Wiśniewska *et al.*, 2014). The molecular method widely used for identification and quantification of *Fusarium* biomass concentration in samples is real time PCR (Horevaj *et al.*, 2011; Nicolaisen *et al.*, 2009; Nielsen *et al.*, 2011, 2013; Niessen, 2007). The aim of the present study was to determine the presence *Fusarium* species and content of mycotoxins in wheat grain in Poland to compare species frequency with earlier reported data.

## Material and methods

### Cereal grain samples

Fifty samples of wheat grain were collected during the harvesting season 2010. They originated from 25 experimental stations of COBORU (the Research Centre for Cultivar Testing) located in different regions of Poland (Figure 1). Two winter wheat cultivars ‘Bogatka’ (medium resistant to FHB) and ‘Muszelka’ (susceptible) were included. The winter wheat was grown with a moderate nitrogen input (avg. 90 kg/ha of N) and without chemical control of diseases. The grain was harvested using combine harvester.

Additionally 18 samples of wheat grain from 2009 and 2010 were analysed. Samples were collected from different locations/fields and cultivars (Table 1, Figure 1). Collected samples were stored at  $-20^{\circ}\text{C}$  before DNA and mycotoxins extraction.

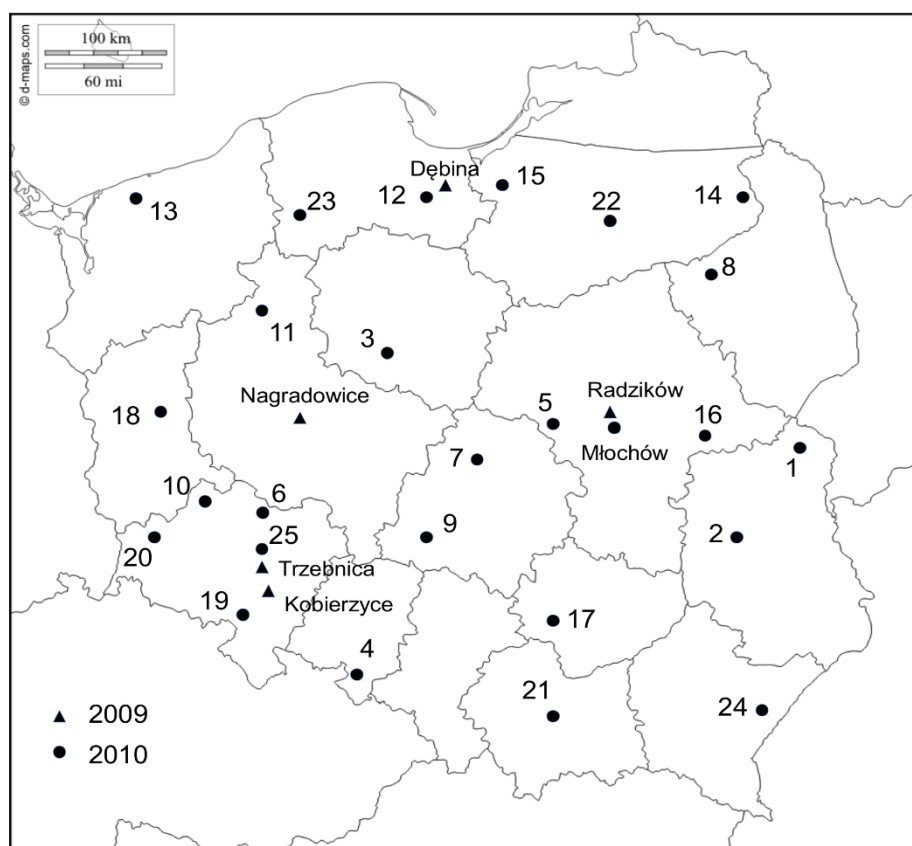


Figure 1. Map of Poland showing sample collection locations of winter and spring wheat in 2009 (triangles) and 2010 (circles). Location numbers correspond to these in Table 3

### DNA extraction and analysis

Grain samples of 300g were initially grounded with a laboratory grinder and 5 g was powdered in liquid N<sub>2</sub> with eight steel balls using Geno/Grinder 2000 (OPS Diagnostics, Bridgewater, NJ). DNA was extracted from 100 mg of that powdered sample using a modified CTAB method (<http://gmo-crl.jrc.ec.europa.eu/summaries/NK603-WEB-ProtocolValidation.pdf>) as described by Nicolaisen *et al.* (2009). DNA extracted from the wheat samples was further purified using a DNeasy kit (Qiagen) according to the manufacturer's instructions.

The *Fusarium* isolates: *F. avenaceum* 9605, *F. culmorum* 9560, *F. equiseti* 8752, *F. graminearum* 1955, *F. langsethiae* 8051, *F. poae* 8452, *F. sporotrichioides* 1926, and *F. tricinctum* 8048 were grown and extracted as described in Nielsen *et al.* (2011). They were grown on potato dextrose agar (PDA) medium at 22°C under 12 h of light and 12 h of darkness for 1–2 weeks prior to DNA extraction. PDA plates before inoculation were covered with sterile cellophane membranes (Horevaj *et al.*, 2011). Mycelium was scraped off the cellophane membrane using a spatula and ground in liquid N<sub>2</sub> with eight steel balls using a Geno/Grinder 2000 (OPS Diagnostics, Bridgewater, NJ). Powdered mycelium (100 mg) was used for DNA extraction, using the same method as for grain samples. The concentration of DNA from *Fusarium* isolates was determined using NanoDrop 1000 (Thermo Fisher Scientific, MA).

Qualitative and quantitative determinations of eight *Fusarium* species in grain were performed by real time-PCR. Primers used were based on fungal TEF-1 $\alpha$  gene sequences, designed by Nicolaisen *et al.* (2009), specific for the different *Fusarium* species: *F. avenaceum*, *F. culmorum*, *F. equiseti*, *F. graminearum*, *F. langsethiae*, *F. poae*, *F. sporotrichioides*, and *F. tricinctum*.

Real-time PCR was carried out in a total of 12.5  $\mu$ l consisting of 6.25  $\mu$ l of 2 $\times$  SYBR Green PCR Master Mix (Applied Biosystems), 250 nM each primer, bovine serum albumin at 0.5  $\mu$ g/ $\mu$ l, and 2.5  $\mu$ l of template DNA. PCR reactions were performed in duplicate on all samples. Genomic DNA from grain samples and pure cultures was diluted 1:10 before PCR.

PCR was performed on a 7900HT Sequence Detection System (Applied Biosystems) using the following cycling protocol: 2 min at 50°C; 95°C for 10 min; 40 cycles of 95°C for 15 s and 62°C for 1 min; followed by dissociation analysis at 60 to 95°C. For the plant assay annealing and extension was performed at 60 °C. Standard curves for *Fusarium* species and wheat were made of five-fold dilution series using pure fungal DNA and wheat DNA. The amount of fungal DNA was calculated from the cycle threshold (Ct) values using the standard curve. The result of each individual sample from each species-specific assay were evaluated by studying the dissociation curve and Ct value, as SYBR Green binds to all double stranded DNA and might create false positives. The plant EF1 $\alpha$  assay was used to provide a normalized measurement for *Fusarium* biomass in each sample, which was calculated as picograms of fungal DNA per micrograms of plant DNA according to Nicolaisen *et al.* (2009).

### Analysis of *Fusarium* toxins

The trichothecenes of group B - deoxynivalenol (DON), nivalenol (NIV) were quantified using gas chromatography techniques. Those mycotoxins were extracted from 5 g of ground grains using 25 ml of an aqueous solution of acetonitrile (acetonitrile: water 84: 16) in a shaker overnight. Samples were centrifuged (3000 rpm min<sup>-1</sup>, 5 min.), and the extract was purified with MycoSep® 227 Trich+ columns (Romer Labs Inc., Union, MO). One microliter of the internal standard solution (chloralose) was added to 4 ml of purified extract. The solvent was evaporated to dryness in the air stream. Mycotoxins were derivatised to the trimethylsilyl derivatives using a derivatizing agent Sylon BTZ (BSA + TMCS + TMSI, 3: 2: 3, Supelco). After dissolution of sample in isoctane, excess of derivatizing agent was decomposed and

removed with water. The organic layer was transferred to autosampler vial and analysed chromatographically with gas chromatograph SRI 8610C, with BGB-5MS column of 30 m in length, and an internal diameter of 0,25 mm.

Hydrogen was a carrier. Elution was carried out in the temperature gradient. Mycotoxin detection was carried out using electron capture detector (ECD). Identification of individual compounds was made by comparing the retention times of the pure standards of mycotoxins. The concentration of mycotoxins was established on the basis of the calibration curve, using chloralose as the internal standard.

The content of zearalenone (ZEN) was determined using a quantitative direct competitive enzyme-linked immunosorbent assay (ELISA) AgraQuant® ZON 40/1000 (LOD 10 ppb) (Romer Laboratories). In 2010, samples were analysed for T-2/HT-2 toxins using AgraQuant® T-2 Toxin 75/500 (LOD 35 ppb) assay (Romer Laboratories).

### Statistical analysis

The original *Fusarium* biomass and toxin concentrations were transformed to logarithmic values in order to obtain a normal distribution for the variables. The relationships between the results for *Fusarium* biomass and *Fusarium* toxins were investigated by Pearson correlation tests. Principal component analysis was used to analyse concentration of biomass of *Fusarium avenaceum*, *F. culmorum*, *F. graminearum*, *F. langsethiae* and *F. poae*, and concentration of biomass of *Fusarium culmorum*, *F. graminearum* and *F. poae*, and concentration of *Fusarium* toxins (DON, NIV, ZEN) in 50 grain samples of winter wheat collected from 25 locations in Poland. Significance of differences between samples from different regions or samples for different *Fusarium* species was analysed using nonparametric tests (Kruskall-Wallis test, Mann-Whitney test). The correlation and PCA analyses were performed using Microsoft® Excel 2010/XLSTAT®-Pro (Version 2013.4.07, Addinsoft, Inc., Brooklyn, NY, USA).

## Results

In wheat samples from 2009 presence of *Fusarium* DNA was found. The highest amount of *Fusarium* biomass was found in grain of spring wheat ‘Griwa’ (Radzików 1) sown after maize and winter wheat ‘Muszelka’ (Dębina 2), which is highly susceptible to *Fusarium* head blight (Table 1). The lowest amounts were detected in grain of winter wheat cultivars ‘Zawisza’ (Radzików 5) and ‘Tonacja’ (Radzików 5) and in spring wheat ‘Raweta’ (Radzików 3).

Table 1. Concentration of *Fusarium* species biomass and DON, NIV and ZEN mycotoxins levels in samples of grain of spring and winter wheat collected in 2009

No.	Sample name	<i>Fusarium</i> biomass [pg/μg] <sup>a</sup>				DON [μg/kg]	NIV [μg/kg]	ZEN [μg/kg]
		<i>Fa</i>	<i>Fc</i>	<i>Fg</i>	<i>Fp</i>			
1	Radzików 1 <sup>b</sup>	1300	153	60248	70	5719	43	63
2	Radzików 2 <sup>b</sup>	89	41	21804	0	2020	0	25
3	Radzików 3 <sup>b</sup>	53	31	911	9	104	0	0
4	Dębina 1	0	316	18966	63	2937	45	78
5	Dębina 2 <sup>c</sup>	533	8862	46102	287	7170	281	29
6	Kobierzyce	0	387	26384	949	n/a	n/a	n/a
7	Nagrodowice <sup>c</sup>	366	22949	5462	67	9239	33	230
8	Radzików 4	0	34	4277	137	658	177	0
9	Radzików 5	63	38	2115	187	213	36	12
10	Radzików 6	0	0	207	33	47	0	17
11	Trzebnica	1753	252	2044	285	123	61	0
	<b>Mean</b>	<b>378</b>	<b>3006</b>	<b>17138</b>	<b>190</b>	<b>2823</b>	<b>68</b>	<b>45</b>

<sup>a</sup> – *F. langsethiae*, *F. sporotrichioides* and *F. langsethiae* were excluded; <sup>b</sup> - spring wheat; <sup>c</sup> – grain from collected symptomatic spikes; *Fa* = *F. avenaceum*, *Fc* = *F. culmorum*, *Fg* = *F. graminearum*, *Fp* = *F. poae*

Seven *Fusarium* species, except *F. langsethiae*, were detected in wheat grain. *F. graminearum* was present in all samples, *F. poae* and *F. culmorum* in ten samples (91%), *F. avenaceum* in seven samples (64%). *F. sporotrichioides* and *F. tricinctum* were found in two individual samples. First species in sample 'Radzików 1' at 69 pg/ $\mu$ g, and the second in wheat grain from Debina ('Dębina 2') at 428 pg/ $\mu$ g. Traces of *F. equiseti* were found in two samples ('Debina 2', 'Nagradowice').

Despite large differences in *Fusarium* biomass content in grain samples, amount of *F. graminearum* DNA was the highest in eight samples (Figure 2). *F. culmorum* dominated only in a sample from Nagradowice and in sample from Trzebnica concentrations of *F. avenaceum* and *F. graminearum* biomasses were similar.

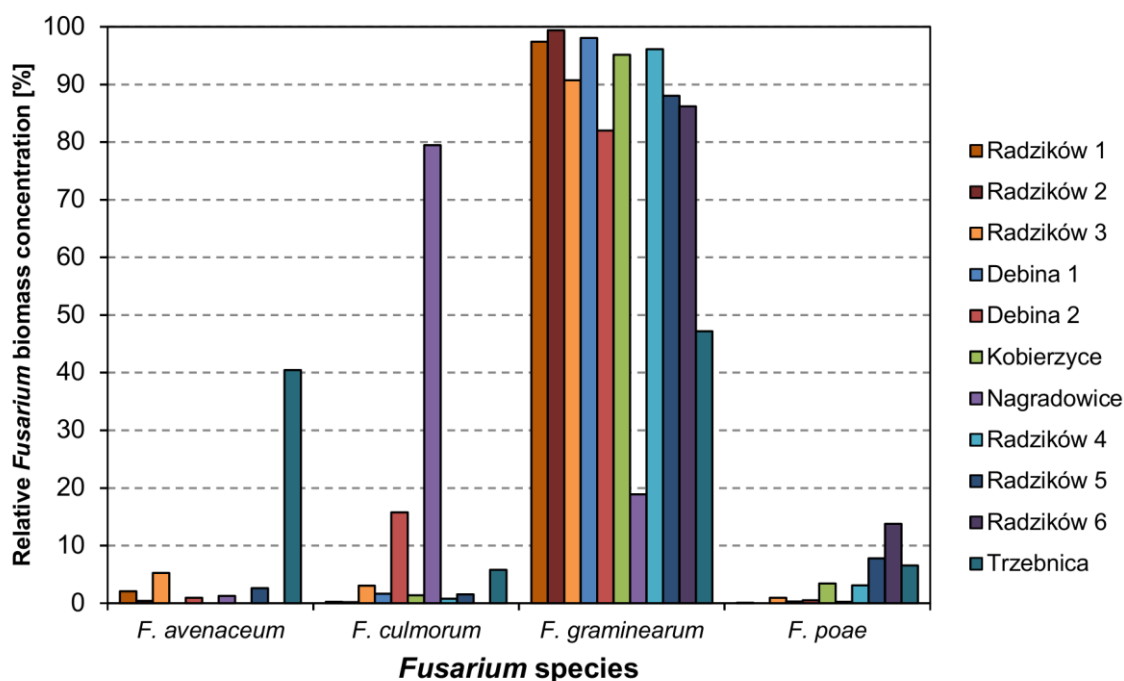


Figure 2. Relative concentration of biomasses of four *Fusarium* species in 11 samples of spring and winter wheat collected in 2009. *F. equiseti*, *F. sporotrichioides* and *F. tricinctum* were excluded

DON was detected in all samples at the average level of 2823  $\mu$ g/kg (Table 1). The most contaminated were grain samples of winter wheat 'Nagradowice' and 'Debina 2' and spring wheat 'Radzików 1'. Levels of NIV were much lower. On average, it was 74  $\mu$ g/kg. NIV was detected in seven samples. The highest concentration was in grain of winter wheat from 'Debina 2' and winter wheat 'Radzików 4'. ZEN was detected in seven samples at the average level of 51  $\mu$ g/kg. Considerable amounts of ZEN were found in samples from Nagradowice and in samples of spring wheat 'Radzików 1' and winter wheat 'Dębina 1'. DON concentration correlated significantly with total *Fusarium* biomass ( $r = 0.947$ ), for NIV and ZEN coefficients were no significant  $r = 0.537$  and  $r = 0.561$ , respectively (Table 2).

When looking at individual species, high correlation between DON and biomass of *F. graminearum* and *F. culmorum* were evident (Table 2). As regards NIV, significant correlation was observed with *F. poae* biomass.

In 2010 average concentration of *Fusarium* biomass was 1970 pg/ $\mu$ g (1430 pg/ $\mu$ g in 'Bogatka' grain and 3770 pg/ $\mu$ g in 'Muszelka' grain) (Table 3). The highest concentration of biomass was detected in grain from Zadąbrowie, South-Eastern Poland (Figure 1). The biomass was five-six times lower in grain from Czesławice (South-Eastern Poland), Rychliki, Radostowo (Northern Pl.) and Głubczyce (Southern Pl.). Very low concentration of biomass was found in grain from

Naroczyce, Nowa Wieś Ujska (western Poland), Kawęczyn (central Poland), and Rarwino (north-western Poland). At a regional scale the highest *Fusarium* biomass concentration was observed in grain from South-Eastern and North-Eastern Poland and the lowest concentrations was observed in grain from Western, north-western and central Poland (Figure 1). Concentration of *Fusarium* biomass was significantly higher in south-eastern region according to Kruskal-Wallis test.

Table 2. Coefficients of correlation between concentrations of biomass (pg/μg) of four *Fusarium* species and mycotoxins (μg/kg) DON, NIV and ZEN in grain of wheat collected in 2009.

	<i>Fusarium</i> biomass					Mycotoxin	
	<i>Fusarium</i>	<i>F. avenaceum</i>	<i>F. culmorum</i>	<i>F. graminearum</i>	<i>F. poae</i>	DON	NIV
<i>F. avenaceum</i>	0.400						
<i>F. culmorum</i>	<b>0.696</b>	0.462					
<i>F. graminearum</i>	<b>0.960</b>	0.273	0.508				
<i>F. poae</i>	0.266	0.043	0.370	0.243			
DON	<b>0.947</b>	-	<b>0.740</b>	<b>0.885</b>	0.083		
NIV	0.537	-	0.557	0.491	<b>0.875</b>	0.477	
ZEN	0.561	-	0.518	0.482	-	<b>0.706</b>	0.070

All variables were log transformed. Values in bold are different from 0 with a significance level  $P \leq 0.05$

Table 3. Concentration of total *Fusarium* biomass, and DON, NIV and ZEN mycotoxins levels in grain of winter wheat cultivars 'Bogatka' and 'Muszelka' from 2010 harvest

No.	Location	<i>Fusarium</i> biomass		DON		NIV		ZEN	
		[pg/μg] <sup>a</sup>		[μg/kg]		[μg/kg]		[μg/kg]	
		Mean	SD	Mean	SD	Mean	SD	Mean	SD
1	Cicibór	2295	620	76.3	7.6	53.7	0.4	17.9	17.9
2	Czesławice	4126	1319	181.4	37.4	63.7	0.0	93.1	35.2
3	Głębokie	842	203	63.5	7.4	59.5	1.1	18.1	18.1
4	Głubczyce	2919	969	127.2	35.5	60.7	5.6	26.8	26.8
5	Kawęczyn	55	54	61.1	2.9	51.8	1.6	0	0
6	Krościna Mała	1327	1276	61.0	7.1	52.2	0.5	0	0
7	Lućmierz	315	36	110.7	54.3	61.5	8.0	0	0
8	Marianowo	1743	948	53.3	1.4	52.1	0.1	0	0
9	Masłowice	650	203	65.3	9.4	51.7	1.7	10.6	10.6
10	Naroczyce	38	38	63.0	5.3	50.8	1.9	9.9	9.9
11	Nowa Wieś Ujska	65	65	51.9	1.7	50.3	1.1	10.1	10.1
12	Radostowo	3466	1428	58.6	1.9	53.0	0.2	0	0
13	Rarwino	108	108	53.5	3.3	52.8	1.1	0	0
14	Ruska Wieś	951	73	55.4	2.7	51.8	0.7	13.5	13.5
15	Rychliki	3230	2138	87.9	9.2	54.3	0.6	42.0	21.5
16	Seroczyn	731	731	78.1	15.0	54.6	3.4	27.6	27.6
17	Słupia	1116	474	107.3	27.7	53.6	3.7	36.9	36.9
18	Świebodzin	303	165	51.2	1.8	51.9	0.2	13.8	13.8
19	Tarnów	1213	14	89.1	28.5	54.8	4.5	0	0
20	Tomaszów Boles.	231	148	76.3	3.5	55.1	0.2	0	0
21	Węgrzce	927	451	86.9	7.2	56.5	3.3	20.1	20.1
22	Wróćkowo	1679	723	165.8	96.7	61.6	9.2	0	0
23	Wyczechy	645	288	83.0	24.0	56.7	5.4	0	0
24	Zadąbrowie	19269	6931	420.3	131.7	57.7	1.3	227.0	21.3
25	Zybiszów	1017	97	76.9	12.4	56.6	0.6	29.3	29.3
<b>Mean</b>		<b>1970</b>	-	<b>96.2</b>	-	<b>55.2</b>	-	<b>23.9</b>	-
<b>Mean 'Bogatka'</b>		<b>1430</b>	-	<b>78.2</b>	-	<b>53.4</b>	-	<b>11.4</b>	-
<b>Mean 'Muszelka'</b>		<b>3770</b>	-	<b>114.2</b>	-	<b>56.9</b>	-	<b>36.3</b>	-

<sup>a</sup> - sum of biomasses of detected *Fusarium* species

Five *Fusarium* species were detected in grain. Biomass of *F. equiseti*, *F. sporotrichioides* and *F. tricinctum* was not detected in any sample. The highest content of biomass was of *F. graminearum* (1252 pg/ $\mu$ g), then *F. avenaceum* (259 pg/ $\mu$ g), *F. langsethiae* (237 pg/ $\mu$ g) and *F. poae* (168 pg/ $\mu$ g) (Figure 3). The content of biomass of *F. culmorum* (55 pg/ $\mu$ g) was very low. The most frequently occurring species were *F. poae* (detected in 74% of samples) and *F. graminearum* (detected in 52% of samples) (Figure 3). *F. poae* was the only species found in 18% of samples. *F. langsethiae* was detected in six samples (five from three locations in northern Poland – Wyczechy, Radostowo, Rychliki). The concentration of *F. langsethiae* in these samples was relatively high 1972 pg/ $\mu$ g compared with an average for samples containing *F. graminearum* DNA – 2235 pg/ $\mu$ g. Lack of differences was confirmed by Kruskal-Wallis test.

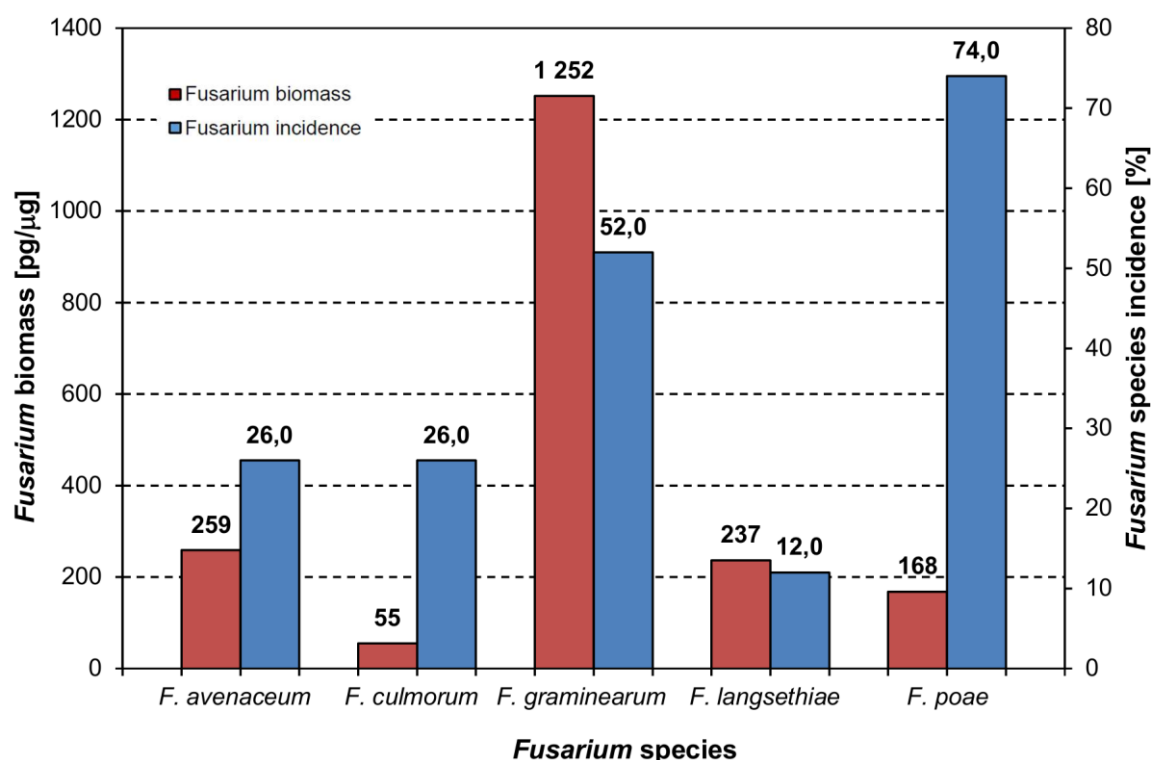


Figure 3. Average concentration of biomass (pg of fungal DNA/ $\mu$ g of wheat DNA) and incidence (percentages) of five *Fusarium* species in 50 samples of winter wheat collected in Poland in 2010

*F. poae* was detected in all samples of medium resistant cultivar ‘Bogatka’ but only in 48% of samples of susceptible ‘Muszelka’. Another species *F. avenaceum* was also found more frequently in grain of ‘Bogatka’ (32%) than ‘Muszelka’ (20%). Three other species (*F. culmorum*, *F. graminearum*, *F. langsethiae*) were detected in grain of both cultivars with similar frequency.

Amounts of biomasses of *Fusarium* species weakly correlated with each other (Table 4). Only coefficient of correlation *F. graminearum* vs. *F. culmorum* was statistically significant. Positive relationship was found between *F. avenaceum* and *F. culmorum* or *F. graminearum* as biomass of the first species was mostly detected in the same locations as the other two species - 1, 4, 14, 19, 22 (only *F. graminearum*), and 24. Biomass of *F. langsethiae* did not correlate with other species, as it was found only in six samples. Otherwise, *F. poae* biomass did not correlate with other species because as the species was present in the most of samples (74%) and in the most samples (except two) amounts of *F. poae* biomass were similar.



Table 4. Coefficients of correlation between concentrations of biomass (pg/μg) of five *Fusarium* species in grain of wheat collected in 2010 from 25 locations.

Species	<i>F. avenaceum</i>	<i>F. culmorum</i>	<i>F. graminearum</i>	<i>F. langsethiae</i>
<i>F. culmorum</i>	0.306			
<i>F. graminearum</i>	0.162	<b>0.461</b>		
<i>F. langsethiae</i>	-0.315	0.066	-0.148	
<i>F. poae</i>	-0.083	0.049	0.005	-0.058

All variables were log transformed. Values in bold are different from 0 with a significance level  $P \leq 0.05$

Biplot produced by PCA analysis on biomass concentration of five *Fusarium* species showed uneven distribution of these species in different locations (Figure 4). *F. culmorum* was present mostly in the same locations as *F. graminearum* (except 12). *F. avenaceum* was present in the same six locations as *F. culmorum* and *F. graminearum* (except 22, where only the second species was detected). In three locations (7, 8, 13) this species was accompanied only by *F. poae*. As it was mentioned earlier, *F. langsethiae* was found in four locations (12, 15, 17, 23). In Słupia (17), it was accompanied by *F. culmorum* and *F. graminearum*, in Radostowo (12) and Rychliki (15) by *F. culmorum* or *F. graminearum*, respectively.

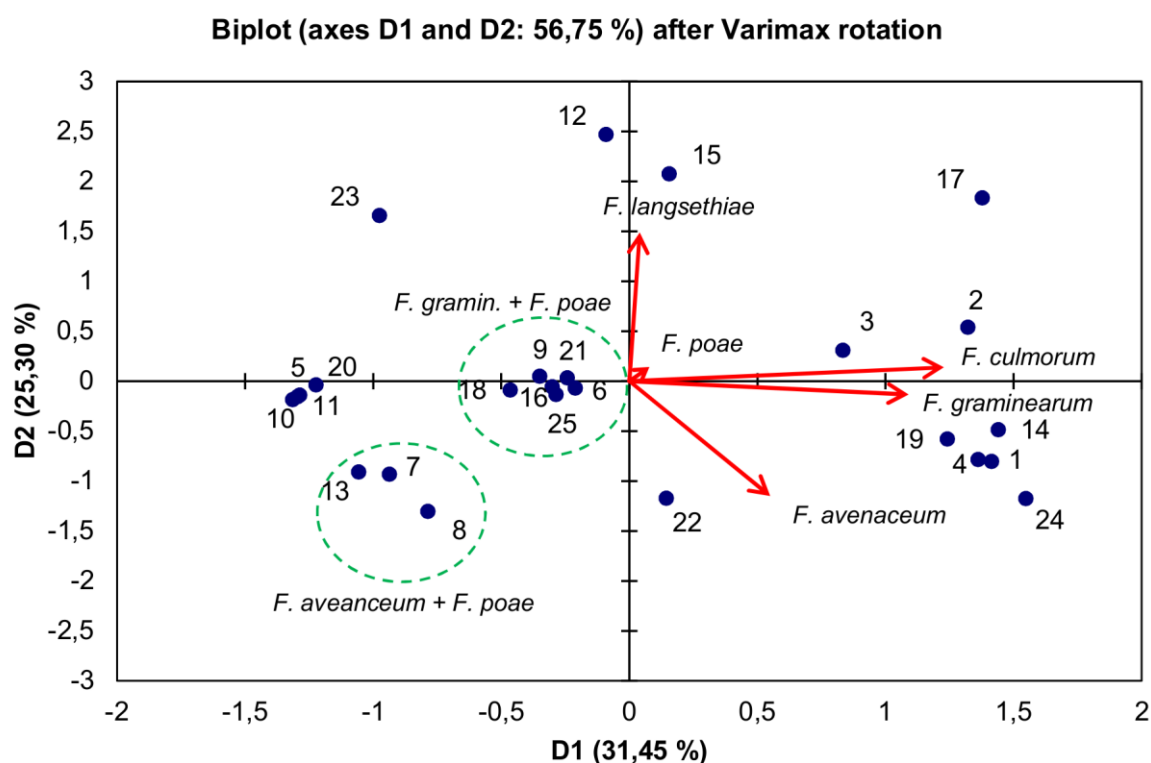


Figure 4. Principal Component Analysis based on biomass of *Fusarium avenaceum*, *F. culmorum*, *F. graminearum*, *F. langsethiae* and *F. poae* in 50 grain samples of winter wheat collected from 25 locations in Poland. Location numbers correspond to those in table 2. Variables were log transformed prior to the analysis

Average content of DON was low and amounted to 96.2 μg/kg, at a range from 49.3 to 552.0 μg/kg (Table 3). The content of NIV was very low – 55.2 μg/kg, at a range 49.2 – 70.8 μg/kg. The average content of DON for ‘Bogatka’ was 78.2 μg/kg, and 114.2 μg/kg for ‘Muszelka’. The highest concentration of DON was found in samples of both cultivars from Zadąbrowie and Czesławice, South-Eastern Poland (Figure 1). High concentration of this toxin was also found in samples of ‘Muszelka’ from Wróćikowo, Lućmierz and Głębczyce.

ZEN was detected in 12% of samples of 'Bogatka' and in 60% of samples of 'Muszelka' cultivars. Average content was 23.9  $\mu\text{g}/\text{kg}$  and was 3 times higher in grain of 'Muszelka' than in 'Bogatka'. High concentration of ZEN was present in samples of 'Muszelka' and 'Bogatka' from Zadąbrowie (248 and 206  $\mu\text{g}/\text{kg}$ , respectively) and in 'Muszelka' sample from Czesławice (128  $\mu\text{g}/\text{kg}$ ).

Table 5. Coefficients of correlation between concentration of biomass ( $\text{pg}/\mu\text{g}$ ) of three *Fusarium* species and concentration ( $\mu\text{g}/\text{kg}$ ) of mycotoxins DON, NIV and ZEN in grain of winter wheat cultivars 'Bogatka' and 'Muszelka' from 2010 harvest in 25 locations

n = 25	<i>Fusarium</i>	<i>Fg</i>	<i>Fc</i>	<i>Fg + Fc</i>	<i>F. poae</i>	DON	NIV	ZEN
DON	<b>0.622</b>	<b>0.534</b>	0.320	<b>0.509</b>	-			
NIV	<b>0.467</b>	0.381	0.242	0.354	0.300	<b>0.695</b>		
ZEN	<b>0.400</b>	<b>0.672</b>	<b>0.406</b>	<b>0.658</b>	-	<b>0.438</b>	0.186	
Toxins	<b>0.649</b>	<b>0.609</b>	<b>0.365</b>	<b>0.587</b>	-0.035	<b>0.974</b>	<b>0.643</b>	<b>0.612</b>

Values in bold are different from 0 with a significance level of  $P \leq 0.05$ ; all variables were log transformed.

*Fg* – *F. graminearum*, *Fc* – *F. culmorum*, toxins – sum of DON, NIV and ZEN.

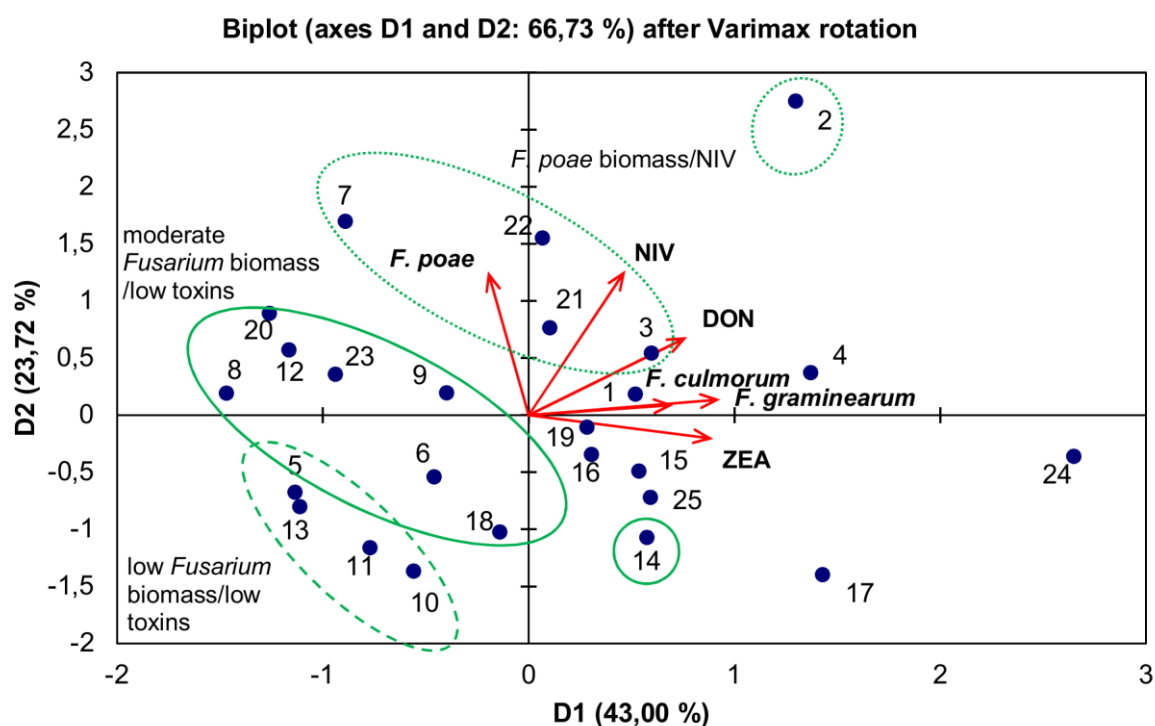


Figure 5. Principal Component Analysis based on biomass of *Fusarium culmorum*, *F. graminearum* and *F. poae*, and concentration of *Fusarium* toxins (DON, NIV, ZEN) in grain samples of winter wheat collected from 25 locations in Poland. Location numbers correspond to these in table 2. Variables were log transformed prior to the analysis.

Six samples of grain containing DNA of *F. langsethiae* were analysed for T-2/HT-2 toxins. In all samples, the total concentration of both mycotoxins was below detection limit of 35  $\mu\text{g}/\text{kg}$ . Amount of *Fusarium* biomass in grain correlated significantly with concentration of *Fusarium* toxins (DON, NIV, ZEN) (Table 5). *F. graminearum* correlated significantly with DON and ZEN concentrations, whereas *F. culmorum* with ZEN concentration only. Biomass of *F. poae* did not correlate with DON and ZEN – toxins not produced by this species. There was some positive relationship between *F. poae* and NIV concentration. Summarised amount of *F. culmorum* and *F. graminearum* biomasses did not improve the strength of correlation with

toxins. Correlation of NIV vs. *F. graminearum* + *F. poae* (possible NIV producers) was statistically significant ( $r = 0.511$ ).

Biplot produced by PCA analysis distinguished some locations based on concentrations of biomass of three *Fusarium* species and *Fusarium* toxins (Figure 5). In Zadąbrowie (24) we found the highest amount of DON and ZEN as well as amount of *F. graminearum* biomass. Grain from Czesławice (2) were characterised by the highest amounts of *F. poae* biomass and NIV but also have high concentrations of the others toxins/biomasses. On the other hand, in Słupia (17) concentration of *F. poae* biomass and NIV was low, but analysis showed high concentration of *F. culmorum* accompanied by moderate concentration of *F. graminearum* and DON.

In the five locations (5, 10, 11, 13) the concentration of biomass of three *Fusarium* species as well as the concentration of toxins were low. In another eight locations (Figure 5, solid line), concentration of toxins was low but amount of *Fusarium* biomass varied from low (23) to high (6). In Ruska Wieś (14) we found the highest concentration of *F. culmorum* biomass (511.8 pg/μg). Five locations (2, 3, 7, 21, 22) could be characterised by above average concentration of NIV and moderate to high concentration of *F. poae* biomass. This species was present at considerable amounts also in other locations (1, 8, 9, 12, 20) but NIV concentration was low.

In samples of grain of spring and winter wheat collected from Radzików and neighbouring Młochów we found more *Fusarium* biomass than in the majority of samples of ‘Bogatka’ and ‘Muszelka’ (Table 6). The highest amount of biomass was present in samples of winter wheat ‘Tonacja’ and ‘Zawisza’ (6998 pg/μg and 5738 pg/μg, respectively). In spring wheat, it was lower, except for the sample of ‘Raweta’ from Radzików (Raweta R1) (5513 pg/μg).

Four *Fusarium* species were detected in grain. *F. langsethiae* and *F. sporotrichioides* were not present. *F. avenaceum* dominated in three samples (on average 1797 pg/μg of biomass) and *F. graminearum* also in three (1432 pg/μg). In one sample (Tonacja R) amounts of biomass of these species were similar. *F. poae* was present in all samples of winter wheat (270 pg/μg). In grain of spring wheat ‘Raweta’ from Młochów only this species was present. The concentration of *F. culmorum* biomass was generally the lowest of all species (67 pg/μg).

Table 6. Concentration of biomass of four *Fusarium* species, and DON and NIV mycotoxins levels in grain of spring and winter wheat from 2010 harvest in Radzików (R) and Młochow (M)

No.	Cultivar/ location	<i>Fusarium</i> biomass [pg/μg]				<i>Fusarium</i> toxins [μg/kg]	
		<i>avenaceum</i>	<i>culmorum</i>	<i>graminearum</i>	<i>poae</i>	DON	NIV
1	Griwa (R)	587	70	1146	262	60,8	50,4
2	Parabola (R)	874	65	257	59	58,1	50,9
3	Raweta (R)	0	57	379	58	61,6	50,0
4	Raweta (R1)	3522	159	1671	161	92,1	51,1
5	Raweta (M)	0	0	0	244	53,0	49,7
6	Tonacja (R)	3223	0	2566	434	64,6	52,3
7	Tonacja (M)	6169	0	0	829	54,5	50,9
8	Zawisza (R)	0	181	5441	116	135,3	52,6
	<b>Mean</b>	<b>1797</b>	<b>67</b>	<b>1432</b>	<b>270</b>	<b>72,5</b>	<b>51,0</b>

The concentration of trichothecene toxins was low (Table 6). ZEN amount was below limit of detection. The highest concentration of DON was found in samples with high concentration of *F. graminearum* and *F. culmorum* – Zawisza R and Raweta R1. The same was true for NIV concentration in grain. No relation was found between *F. poae* and NIV; however, total concentration of *F. graminearum* and *F. poae* correlated the best with NIV amount.

## Discussion

Presence and concentration of *Fusarium* toxins in naturally infected wheat is generally in accordance with data on occurrence of *Fusarium* species on wheat in Poland. According to the published data, dominant species on wheat spikes and kernels were *F. culmorum*, *F. graminearum*, *F. avenaceum* and *F. poae* (Bottalico and Perrone, 2002; Chełkowski *et al.*, 2012; Perkowski *et al.*, 1990). Proportions of these four species changed depending on year and study. *F. graminearum* was the dominating species in wheat grain in both 2009 and 2010. In 2009 it was present in all samples, and in 2010 in 52%. Both years the average amount of *F. graminearum* biomass in grain was the highest of all investigated *Fusarium* species. *F. culmorum* was found in 2009 in 90% of samples; however, the biomass concentration of this species was approximately five times less than *F. graminearum*. *F. culmorum* prevailed only in one sample of grain. In 2010, *F. culmorum* was present in 31% of samples; however, its amount was the lowest among five identified species. Weather in 2009 was more favourable for FHB development than in 2010, which is also reflected in the difference in amount of *Fusarium* DNA and mycotoxins. In some regions (e.g. Radzików) in 2010, the drought conditions occurred in June and July with high temperatures and infrequent, heavy rainfalls. Despite differences in weather and limited number of samples in 2009, *F. graminearum* was occurring more frequently than *F. culmorum*. Amount of biomass of the first species was also higher in both years. While *F. culmorum* biomass was very low in 2010, we can conclude that dry weather is affecting to large extent presence of this species. In the Netherlands in 2009 incidence and amount of *F. culmorum* DNA was similarly low as in our study (van der Fels-Klerx *et al.*, 2012). Authors found this species only in 2% of samples and DNA concentration was 80-times lower than for *F. graminearum*.

In the paper of Wiśniewska *et al.* (2014) authors found that *F. culmorum* was the most common species on heavy infected heads of wheat in 2009. They analysed samples from six locations, and only in two in southern Poland *F. graminearum* prevailed over *F. culmorum*. These difference with our results could be effect of low number of locations sampled in (Wiśniewska *et al.*, 2014) paper as well as in present study (ten locations). The number of examined heads could be the next reason of results difference. In Radzików authors, sampled 16 symptomatic heads whereas in our research we analysed grain from five fields or plots harvested with combine and next sampled for analysis. Wiśniewska *et al.* (2014) used method of isolation of *Fusarium* species from infected kernels and next identification of pure cultures.

The issue of sampling was studied and discussed by van der Fels-Klerx *et al.* (2012). They found significant difference between pre-harvest and at harvest samples. Some species (*F. langsethiae*, *F. avenaceum*) were not found (using real-time PCR) in pre-harvest samples, but detected in harvested grain. In inoculation experiment with different species Xu *et al.* (2007) found that correlation between FHB symptoms on heads and *Fusarium* biomass in grain was weak for *F. culmorum* and *F. poae* comparing with *F. graminearum*. Authors suggest that *F. culmorum* and *F. poae* colonised mainly floral tissues and to the lesser extent kernels. On the contrary, *F. graminearum* might colonise both tissue types to a similar degree.

Tomczak *et al.* (2002) analysed *Fusarium* species causing Fusarium head blight epidemics in 1998 and 1999 in two regions of Poland. In 1998 in northern and central regions *F. avenaceum* dominated, being followed by *Fusarium graminearum* and *F. culmorum* with similar frequency. In 1999, ranking of species was the same; however, frequency of *F. graminearum* was 3-5 times higher than *F. culmorum*. Authors reminds that no *F. graminearum* was detected in the previous decade (1980's) in wheat grown in northern Poland.

Stępień and Chełkowski (2010) summarized frequencies of *Fusarium* species infecting wheat heads in Poland from 1985 to 2009. In 1985 *F. avenaceum* and *Microdochium nivale* dominated, *F. culmorum* being the third species. In 2009, *F. graminearum* dominated and *F.*

*culmorum* was the second species with about half frequency of first species. Increase of *F. graminearum* was obvious; however, differences between particular years were substantial.

*Fusarium graminearum* is the anamorph of *Gibberella zeae*, which produces sexual spores (ascospores) in perithecia. For *F. culmorum* species perfect stage is not known and fungus produces only asexual spores – macroconidia (Scherm *et al.*, 2013). Thus *F. graminearum* can disperse and infect host plants with ascospores and macroconidia, whereas *F. culmorum* only with macroconidia. Nature of *Gibberella zeae* is the homothallic which allows the production of large masses of ascospores and effectively compete against *F. culmorum* (Waalwijk *et al.*, 2003). In a recent study in Germany, the important contribution of ascospores to inoculum pressure was emphasized (Obst *et al.*, 2002). Ascospores required a relative humidity below 53%, whereas macroconidia required relative humidity of above 80% for germination, as was observed by Beyer *et al.* (2005). It can be another factor favouring *F. graminearum* over *F. culmorum* under dry conditions.

Deoxynivalenol (DON) was the most frequently detected toxin in the samples and its accumulation was the most closely associated with the presence of *Fusarium graminearum*. Coefficient was very high in 2009, because of high DON accumulation and high *F. graminearum* biomass amount in grain. In this year DON concentration correlated strongly also with *F. culmorum* biomass despite its low concentration in the most of samples. In 2010, coefficients were lower and significant only for *F. graminearum*.

Nivalenol (NIV) accumulation was much lower but significantly associated with the presence of *F. graminearum* and *F. poae* in 2009. In 2010, coefficients were insignificant but positive for all three possible NIV producers: *F. culmorum*, *F. graminearum* and *F. poae*. Xu *et al.* (2003) studied wheat grain samples harvested in 2001 from UK, Ireland, Italy and Hungary. They did not found quantitative relationships between amount of *Fusarium* DNA and the concentration of the mycotoxins in the grain. However, for total *F. graminearum* and *F. culmorum* DNA and DON concentration linear model was nearly significant. In the next survey (Xu *et al.*, 2008b) they studied *Fusarium* species frequency and mycotoxin content in wheat samples from the same countries over two years (2003-2004). They found DON being the most frequently detected toxin. DON amount correlated strongly with *F. graminearum* DNA. NIV was related significantly only to the amount of *F. culmorum* DNA. As regards ZEN, authors found strong association with both *F. culmorum* and *F. graminearum*.

In 2005 in Poland the highest amount of zearalenone was found in wheat grain infected by *F. graminearum* (Gromadzka *et al.*, 2008) In grains were *F. culmorum* was the main pathogen, zearalenone content was 10-times lower.

*Fusarium poae* was the most frequently species detected in grain (100% of samples in 2009 and 74% of samples in 2010). In 2010 in 9 samples out of 50 it was the only *Fusarium* species present. However, amount of *F. poae* biomass was about 10-times lower than *F. graminearum* biomass in dry 2010 year and up to 200 times lower in year 2009 of weather favourable for *Fusarium* head blight. According to other reports *F. poae* was frequently isolated from wheat spikes and kernels in Poland (Goliński *et al.*, 1996). This is weak pathogen of cereal spikes, however is widespread on wheat in Europe (Isebaert *et al.*, 2009; Vogelgsang *et al.*, 2008; Xu *et al.*, 2003). Audenaert *et al.* (2009) observed dominance of this species in Flanders in years 2007 and it was isolated with lower frequency in 2008. In year 2007, the infection pressure was very high as compared with 2008. Author explained this fact as a result of *F. poae* nature as a secondary pathogen. Additionally, high frequency of occurrence of *F. poae* was explained by its sporulation strategy. This species produces very large amounts of small microconidia in a dry powdery form that can easily invade cereal heads. It could be true for dry conditions and wind dispersal, because for splash dispersal Hörberg (2002) did not find any difference in patterns between *F. poae* microconidia and much larger macroconidia of *F. culmorum*.

Xu *et al.* (2008a) associated *F. poae* with relatively dry and warm conditions, whereas *F. graminearum* with warm/humid conditions. *F. avenaceum* and *F. culmorum* were both associated with niches of cooler/wet/humid conditions. Parikka *et al.* (2012) who expected increase of importance of *F. poae* (accompanied by *F. langsethiae*) in more dry conditions of Scandinavia stated the similar. The weather conditions in the most regions of Poland in 2010 were dry and warm during and after flowering. Results showed that this favoured *F. poae* spread on wheat. As only in the south/south-eastern Poland weather was warm and humid, *F. graminearum* dominated in grain from this region.

Low *F. poae* biomass in grain observed in our study could be explained by lower aggressiveness of this species as compared to *F. graminearum* (Stenglein, 2009; Vogelgsang *et al.*, 2008). It was also found that *F. poae* that predominated in wheat glumes was not detected in grain, which was infected by *F. culmorum*, *F. avenaceum* and *M. nivale* (Doohan *et al.*, 1998). Authors did not detect *F. graminearum* in wheat samples (collected in England, UK in 1994) which is good example of later *Fusarium* species shift in Europe. Polley and Turner (Polley and Turner, 1995) found that *F. poae* was associated with distinct glume spot lesions and was the most frequently isolated from glumes. Doohan *et al.* (1998) supposed that the infection process and colonization by *F. poae* differs from that of other *Fusarium* species causing FHB.

*Fusarium poae* is known as NIV producer (Schollenberger *et al.*, 2006; Thrane *et al.*, 2004). Consequently, we detected NIV in the majority of samples but at very low quantities. In Poland NIV was found primarily in oats infected by *F. poae* (Perkowski *et al.*, 1997). Edwards *et al.* (2012) found that correlation of nivalenol concentration in oat grain and *F. poae* DNA was highly significant but only accounted for 9% of the variance. It showed that other species such as *F. graminearum* and *F. culmorum* were involved in NIV production. NIV chemotypes of these species are not frequent in Poland. Stępień *et al.* (2008) found that only 12% of *F. graminearum* isolates in Poland displayed the NIV chemotype.

Besides NIV *F. poae* isolates were found to produce wide range of toxins including trichothecens of type A and B, beauvericin, enniatins, moniliformin, and others (Bottalico and Perrone, 2002; Somma *et al.*, 2010; Stenglein, 2009; Thrane *et al.*, 2004; Uhlig *et al.*, 2006). The surveys of wheat harvested in Poland in 2006 and 2007 showed that increased importance of *F. poae* in the FHB complex in Poland (Kulik and Jestoi, 2009).

In 1994 Norwegian researchers found “powdery *F. poae*” strains which were the most abundant potential producer of HT-2 and T-2 toxins in cereals (Kosiak *et al.*, 2003). In 1999 these *F. poae* strains were proved to produce T-2 toxin (Torp and Langseth, 1999). Strains originated mainly from Norwegian oats but where found also on wheat in Austria and the Netherlands. Further Torp and Nierenberg (2004) described these strains as a new species *F. langsethiae*. The species was being found primarily in northern Europe on oats and barley (Edwards *et al.*, 2012; Yli-Mattila *et al.*, 2008). Occurrence of *F. langsethiae* on wheat in Poland was confirmed in 2008 (Lukanowski *et al.*, 2008). This species was found mainly in northern Poland (including Radostowo mentioned in present study), however it was present in some samples of wheat grain from Central Poland (Lukanowski and Sadowski, 2008). In 2009 *F. langsethiae* was found on wheat grain in the Netherlands but at low level (8% of samples) (van der Fels-Klerx *et al.*, 2012). Presence of *F. langsethiae* was detected by Czaban *et al.* (2015) in years 2008 – 2010 in south-eastern Poland. Percentages of winter wheat kernels colonised by this species was low. It ranged from 0 to 2.9% in susceptible cultivar ‘Kris’ in 2010. In our research, we did not detect *F. langsethiae* in 2009, however limited number of samples was analysed. In 2010, DNA of this species was found mainly in samples from northern Poland and in only one from southern region at low concentration.

*F. langsethiae* and *F. poae* are favoured by dry conditions (Parikka *et al.*, 2012), however it seems that the first species prefer lower temperatures than the former. Kokkonen *et al.* (2012)

found that *F. langsethiae* produced the highest amount of the type-A trichothecenes at 15°C, whereas *F. poae* could produce beauvericin at both cool and warm conditions.

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