Grain dust originating from organic and conventional farming as a potential source of biological agents causing respiratory diseases in farmers

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Abstract

Introduction: Agricultural producers are exposed to a number of different health risks associated with their work environment.

Aim: The objective of the study was to assess the degree of colonization by fungi in terms of quantity and in terms of variety of species the samples taken from the settled dust from combine threshing of rye cultivation from organic and conventional farms in the Province of Lublin.

Material and methods: This paper is a preliminary quantitative assessment of the species of fungi colonizing the samples of settled dust collected during combine threshing from organic and conventional farms in the Province of Lublin. One of the stages of the project was the classification of biosafety BSL (biosafety level) of selected isolates and API ZYM tests to evaluate the potential ability of isolates to cause adverse health effects. To determine the concentration and composition of fungi in collected samples plate dilution method was used with two media: Malt Agar and Potato Dextrose Agar.

Results: Most commonly isolated fungi in settled dust samples collected during combine threshing from organic farms, on PDA medium were: *Alternaria alternata* and *Aureobasidium pullulans*. Cultures on MA medium were dominated by *Alternaria alternata*, *Mycelia sterilia* and *Fusarium poae*. In samples of dust from conventional crops, the predominant species was *Alternaria alternata* on PDA medium and on MA medium.

Conclusions: The obtained results show a potential risk of people involved in agricultural work.

Key words: organic agriculture, conventional crops, molds, alveolitis allergica.

Introduction

Conventional farming uses measures of industrial production, including fertilizers and chemical pesticides and industrial feed, ensuring higher productivity of plants and animals, and higher efficiency than organic farming [1]. Organic farming is a farming system based on respect for the principles of crop rotation, covering the soil with plants throughout the growing season, using mainly natural and organic fertilizers, while limiting mineral fertilizers and plant protection products, take action to protect plants, compliance with standards of space and cleanliness of the premises including the definition of livestock stocking limit on the area of agricultural land [2].

Agricultural producers are exposed to a number of different health risks associated with the work they undertake. One such factor is the dust of biological origin (organic powder). Depending on the place of origin it may contain various components of cereals, other plants, molds and fungi, mycotoxins, bacteria, dust mites, insect fragments and animal excretions of very diverse composition [3, 4].

Aim

The objective of the study was to assess the degree of colonization by fungi in terms of quantity and in terms of variety of species the samples taken from the settled

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dust from combine threshing of rye cultivation from organic and conventional farms in the Province of Lublin.

Selected genera and species of fungi were ranked in terms of health risk to people doing different types of work in agriculture – the class of Biosafety Levels (BSL). The selected isolates were also evaluated for the presence and activity of extracellular hydrolytic enzymes (API ZYM test, bioMérieux), which may indicate the potential ability of isolates to cause adverse health effects. This type of study has been designed because of the dynamic growth in the number of organic farms in Poland, a large exposure to agricultural producers to organic dusts and potential ability of fungi and their metabolites of inhaled organic dust to cause respiratory diseases.

Material and methods

The material consisted of settled dust samples collected during combine threshing in two agricultural systems: organic and conventional in the Province of Lublin (10 settled dust samples from each system of cultivation – a total of 20 samples in the months of July–August 2012).

To determine the concentration and composition of fungi in samples two media were used: Malt Agar (Malt Agar, Becton, Dickinson and Company) and Potato Dextrose Agar (PDA, Becton, Dickinson and Company) and the method of plate dilution with the addition of Tween 80. The study was conducted in two parallel repetitions. Malt Agar cultures were incubated at 30°C for 72 h and at room temperature for another 72 h and PDA cultures were incubated at 24°C for 144 h. Species composition of fungal flora was determined using macroscopic and microscopic methods and keys and atlases [5–10] and quantities were expressed in units of CFU/g.

Isolated fungi were ranked for the relevant class of biosafety BSL. In some species of fungi isolated from samples the hydrolytic enzymes activity was determined using the API ZYM test (bioMérieux). Selected strains were grown in PDA for 16 days at 24°C. Then the collected fragments of fungal culture with a diameter of 5–6 mm were placed in two liquid media: Sabouraud Dextrose Broth (Becton, Dickinson and Company) and Czapek-Dox Broth (Becton, Dickinson and Company). The cultures and controls were incubated for 16 days at 24°C. Culture supernatants after centrifugation were used in the API ZYM test. The next steps were complied with the API ZYM test manufacturer's instructions.

Results

Identification of fungi

In the analyzed samples of the rye dust from organic farms most often the following fungi species were found: Alternaria alternata, Aureobasidium pullulans, Mycelia sterilia, Fusarium poae, Cladosporium macrocarpum. The dust of rye from conventional farms was dominated by

Alternaria alternata, Cladosporium macrocarpum, Fusarium poae, Fusarium tricinctum and Mycelia sterilia. The most diverse in terms of species composition were dust samples from conventional crops isolated on MA, and the least diverse sample came from the dust samples from organic farms isolated on PDA medium, but the differences were small in number.

Most commonly isolated fungi in settled dust samples collected during combine threshing from organic farms, on PDA medium were: *Alternaria alternata* (16.1%) and *Aureobasidium pullulans* (15.3%). Cultures on MA medium were dominated by *Alternaria alternata* (24.5%), *Mycelia sterilia* (11.9%) and *Fusarium poae* (10%). In samples of dust from conventional crops, the predominant species was *Alternaria alternata* on PDA medium in 12.6%, and on MA medium – 18.2% (Table 1).

The most diverse isolates from the dust of both conventional and organic farming were cultured on PDA medium. The concentration of fungi in samples collected from dust settled during combine threshing derived from organic farms isolated on PDA were in the range from 965×10^3 to 7170×10^3 CFU/g, and isolated on MA ranged from 635×10^3 to 6060×10^3 CFU/g. The concentration of fungi in samples collected from settled dust during combine threshing from conventional farms isolated on PDA were in the range from 1300×10^3 to 13445×10^3 CFU/g, and on MA medium ranged from 1105 × 10³ to 5305 × 10³ CFU/g. In general, higher levels of fungi were found in settled dust samples collected during combine threshing from conventional farms compared to organic farms isolated on PDA medium. In organic farming, conventional antifungal preparations and plant protection products are not applied, however relevant strains for this type of cereals cultivation better deal with threats such as mold fungi [11]. When the substrate is MA, the concentration of fungi is higher in samples derived from organic farms (Table 2).

Classification of biosafety (BSL)

Shown in Table 3, isolates of fungi described due to the risk to people doing various types of work in agriculture.

This was done using the classification established by the European Confederation of Medical Mycology in 1996. The Classification of Biosafety Level (BSL) is the scale of potentially pathogenic fungi for humans and animals, and distinguishes three hazard classes represented by different species of fungi. The class BSL-1 is classified as saprophytes or plant pathogens causing superficial, non-invasive or mild threat. The class BSL-2 species is characterized by a relatively strong ability to survive in vertebrate tissues in patients with severely impaired immunity and can cause deep, opportunistic infection. The class of BSL-3 pathogens is potentially capable of inducing severe deep fungal infections in apparently healthy individuals [12].

Most frequently occurring fungi in settled dust collected during combine threshing mostly belong to the

Table 1. Fungi identified in dust samples from organic and conventional crops, isolated on Malt Agar (MA) and Potato Dextrose Agar (PDA)

Fungi	Organic fa	rming [%]	Conventional farming [%]				
Fungi	PDA	MA	PDA	MA			
Acremonium charticola	1.3	1.2	0.0	0.0			
Acremonium strictum	0.0	0.5	3.4	2.1			
Alternaria alternata	16.1	24.5	12.6	18.2			
Alternaria sp.	0.0	0.0	3.2	0.0			
Aspergillus candidus	0.0	0.0	1.0	4.1			
Aspergillus glaucus	0.0	0.0	1.2	1.0			
Aspergillus versicolor	1.5	1.6	6.1	9.4			
Aureobasidium pullulans	15.3	2.0	6.1	3.3			
Aureobasidium sp.	0.0	0.7	0.0	3.0			
Blastomyces dermatitidis	0.0	0.9	0.0	0.0			
Chaetomium atrobrunneum	0.0	0.0	0.0	0.3			
Chaetomium sp.	0.0	0.0	0.0	0.2			
Cladosporium herbarum	0.0	5.3	3.3	0.0			
Cladosporium macrocarpum	11.3	9.8	11.2	10.1			
Cladosporium sphaerospermum	9.1	6.4	5.0	4.0			
Fusarium cerealis	0.9	0.0	0.0	0.0			
Fusarium oxysporum	0.0	3.3	0.0	0.0			
Fusarium poae	12.2	10.0	10.5	10.0			
Fusarium solani	0.0	0.0	0.0	0.5			
Fusarium sp.	0.7	0.6	2.4	0.0			
Fusarium sporotrichioides	0.4	0.0	2.1	0.9			
Fusarium tricinctum	9.2	6.9	12.5	7.3			
Gonatobotrys sp.	1.0	0.9	0.7	2.3			
Madurella grisea	0.0	0.0	0.0	0.4			
Microsporum sp.	0.4	0.7	0.0	0.0			
Mucor racemosus	1.1	3.1	1.5	1.9			
Mycelia sterilia	12.7	11.9	8.4	9.9			
Paecilomyces variotii	0.0	0.6	0.0	0.0			
Penicillium chrysogenum	0.5	1.2	0.0	0.4			
Penicillium citrinum	2.1	2.1	0.4	0.5			
Penicillium diversum	0.0	0.1	0.0	0.0			
Penicillium expansum	2.0	2.9	4.1	3.4			
Penicillium gerundense	0.0	0.0	2.5	3.5			
Phialophora sp.	0.0	0.0	0.0	0.3			
Rhizopus oryzae	0.0	0.2	0.4	0.0			
Scopulariopsis brevicaulis	0.0	0.0	0.0	0.8			
Trichoderma harzianum	0.0	0.0	0.6	0.8			
Trichophyton sp.	0.5	0.0	1.0	0.2			
Ulocladium chartarum	0.0	0.0	0.0	1.2			
Ulocladium sp.	1.6	1.1	0.0	0.0			
Yeast-like fungi	0.3	1.1	0.0	0.0			
Total	100	100	100	100			

Sample no. 1 2 3 4 5 6 7	Organic farmin	g [× 10³ CFU/g]	Conventional f	arming [CFU/g]
Sample no.	PDA	MA	PDA	MA
1	1455	1575	1300	1105
2	965	635	1495	1330
3	1835	5725	11320	1705
4	7170	5400	2380	2700
5	1230	1450	3760	3135
6	5270	1910	2480	2035
7	6830	1300	13445	1420
8	5530	6060	1870	2390
9	2100	1110	2690	1985

785

2595

Table 2. Concentration of fungi in samples of settled rye dust during combine threshing

BSL-1 class causing a non-invasive or mild danger. Some fungal isolates have not yet been included in the classification of BSL, and several of them are in BSL-2 class that is particularly dangerous for people with immune deficiency.

2965

3535

10

Mean

Activity of extracellular hydrolytic enzymes API ZYM test (bioMérieux)

Isolated fungi from tested samples of dust were initially used to test with API ZYM (bioMérieux), which enables the assessment of the activity of 19 extracellular hydrolytic enzymes. It is generally believed that the extracellular hydrolytic enzymes secreted by both dermatophytes, yeasts and fungi play an important role in the pathogenesis of infection. These enzymes catalyze reactions of hydrolysis of C-O, C-N, and C-C bonds, making it easier for the fungus to invade the tissues, on the other hand they can stimulate the immune inflammation limiting the infection [13].

The greatest number of active hydrolytic enzymes among 9 selected isolates grown on Sabouraud Dextrose Broth (Becton, Dickinson and Company) showed species of Aspergillus versicolor (β -glucosidase, alkaline phosphatase) and Fusarium oxysporum (β -glucosidase, N-acetyl- β -glucosaminidase). The smallest number of active hydrolytic enzymes in these isolates was shown by the following species: Cladosporium herbarum, Fusarium cerealis and Penicillium diverse. The highest activity among the tested isolates was shown by the enzymes: β -glucosidase and N-acetyl- β -glucosaminidase.

In the studies, of the same isolates on Czapek-Dox Broth (Becton, Dickinson and Company), the highest number of active enzymes showed hydrolytic species of Aspergillus versicolor (α -galactosidase, acid phosphatase, alkaline phosphatase), and Aspergillus candidus (α -galactosidase, N-acetyl- β -glucosaminidase, acid phosphatase, β -glucosidase), and the least amount of active hydrolytic

enzymes in this medium had species of *Penicillium expansum* and *Fusarium oxysporum*. The highest activity among the isolates tested in this medium was shown by the same enzymes as in the medium Sabouraud Dextrose Broth i.e. β -glucosidase and N-acetyl- β -glucosaminidase. The results of the API ZYM test are presented in Table 4.

5305

2311

2410

4315

Discussion

Fungal species belonging to different taxonomic groups are important from the point of view of allergology [14]. Most are found in the common genera *Cladosporium* spp., *Alternaria* spp., *Aspergillus* spp., *Penicillium* spp., and *Mucor* spp. Due to the structure of their spores (low molecular weight), as compared with the pollen, they are easier lifted in the air and penetrate deeper into the respiratory tract of the human [15]. Then there are the fungi of the genera *Botrytis* spp., *Fusarium* spp., *Pullularia* spp., *Curvularia* spp., *Phoma* spp. and *Rhizopus* spp., which are among the most common allergens causing inhaled allergy [16].

The isolated fungi include species *Alternaria alternata*. Its spores, and not mycelium, tend to be the cause of allergy. Spores often float in the air, get into the nose and lungs and can cause the symptoms of hay fever and asthma [17].

Another frequently occurring fungi in the tested samples were *Fusarium* spp., which are cosmopolitan species. The best conditions for the development of *Fusarium* spp. occur in zones of tropical and temperate climate, although it is also isolated from the desert, alpine, arctic and other extreme environments. *Fusarium* spp. are common in soil, as saprotrophs. It can also grow on plant debris and various other organic substrates. Some fungi of the genus *Fusarium* are facultative parasites, which in favorable conditions can cause diseases of plants (in underground and overground parts) [18]. Mushrooms of the genus *Fusarium* are considered one

Table 3. Biosafety level (BSL) of fungi isolated from rye dust [9, 12, 28, 29]

Fungi isolated from rye dust	Biosafety level
Acremonium charticola	ND
Acremonium sp.	ND
Acremonium strictum	1
Alternaria alternata	1
Alternaria sp.	1
Aspergillus candidus	I
Aspergillus glaucus	I
Aspergillus versicolor	I
Aureobasidium pullulans	I
Aureobasidium sp.	ND
Blastomyces dermatitidis	II
Chaetomium atrobrunneum	I
Chaetomium sp.	I
Cladosporium herbarum	I
Cladosporium macrocarpum	ND
Cladosporium sphaerospermum	1
Fusarium cerealis	ND
Fusarium oxysporum	II
Fusarium poae	ND
Fusarium solani	II
Fusarium sp.	ND
- Fusarium sporotrichioides	ND
- -usarium tricinctum	ND
Gonatobotrys sp.	ND
Mycelia sterilia	ND
/east-like fungi	ND
Madurella grisea	II
Microsporum sp.	ND
Mucor racemosus	I
Paecilomyces variotii	II.
Penicillium chrysogenum	I
Penicillium citrinum	1
Penicillium diversum	ND
Penicillium expansum	1
Penicillium gerundense	ND
Phialophora sp.	ND
Rhizopus oryzae	II
Scopulariopsis brevicaulis	I
 Trichoderma harzianum	1
Trichophyton sp.	ND
Ulocladium chartarum	I
Jlocladium sp.	ND
UD is a to distance to a dista	

ND – not determined

of the deadliest pathogens of cereals, potatoes, vegetables, ornamental and forest plants causing significant economic losses [19].

In the past 20 years, the interest in fungi of the genus *Fusarium* spp. increased due to their ability to produce mycotoxins harmful to humans and animals [20]. Contamination of agricultural products with *Fusarium* mycotoxin is now a worldwide problem [21, 22]. Mycotoxins have the ability to penetrate through the digestive tract, respiratory system and skin. Mycotoxins in high concentrations can damage internal organs leading to severe clinical disease. At low concentration, which is often reported, these compounds can accumulate in human and animal tissues, and cause a variety of chronic diseases, including liver and kidney cancer. These chemicals can get into the eggs, meat and milk of animals fed contaminated feed and become indirectly a threat to human health [23].

Years of research conducted at the Institute of Rural Health have shown that exposure to organic dust can lead to many respiratory syndromes, such as allergic diseases (extrinsic allergic alveolitis – EAA – alveolitis allergica, bronchial asthma, allergic rhinitis, allergic conjunctivitis, allergic dermatitis – hives or airborne eczema), immunotoxic diseases (organic dust toxic syndrome – ODTS, byssinosis, "chronic fatigue syndrome" – "sick building syndrome"), nonspecific diseases, chronic bronchitis [24–26].

In 1932, Campbell described the clinical symptoms of EAA (farmer's lung disease) which affected farmers working with moldy hay [3]. This disease includes a number of diseases with a very similar clinical picture, but caused by different etiological factors. Due to the allergen and the nature of the exposure, it is called farmer's lung, lung threshing, malt worker's lung (brewers), mushroom worker's lung, bagassosis, cheesemaker's disease, bird breeder's lung, maple bark disease, woodworker's disease, suberosis, detergent worker's disease [24].

Another disease that occurs when organic dust is inhaled is the ODTS. Earlier this syndrome was described as mycotoxicosis (pulmonary), atypical farmer's lung, EAA without precipitins, silo unloader syndrome, toxic fever or inhalation fever. The ODTS is defined as a process covering both the airways and alveoli. Characterized by the formation of acute symptoms as the flu-like syndrome with no radiological signs of EAA after exposure to high concentrations of organic dust. The potential etiologic agents in this case are the bacteria and their products (endotoxin), fungi or their spores, mycotoxins, glucans, or other yet unidentified substances. To cause disease a single exposure is sufficient in contrast to the EAA where prior sensitization is necessary. Most cases of ODTS have been found in workers engaged in pig farming and in workers exposed to grain dust [3, 24].

In the etiology of chronic bronchitis (CB), the organic powder is one of the risk factors. The main importance

Table 4. Activity of enzymes secreted by isolated fungi on media Czapek-Dox Broth (Czapek) and Sabouraud Dextrose Broth (Sabouraud)

Isolates	Medium		Activity of enzymes*.**																	
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19
Aspergillus	Czapek	3	2	1	_	1	2	1	-	_	4	2	5	2	-	-	4	5	-	_
candidus	Sabouraud	3	-	1	_	-	1	_	-	_	1	2	3	2	-	-	4	1	2	_
Alternaria	Czapek	_	2	3	_	2	1	_	4	-	1	1	2	_	_	1	5	5	_	-
alternata	Sabouraud	-	_	-	_	1	1	_	1	-	-	1	2	-	-	1	5	2	-	_
Penicillium	Czapek	_	1	1	_	5	2	_	-	_	-	1	_	2	_	3	4	4	-	2
chrysoge- num	Sabouraud	-	-	-	_	-	_	_	_	_	_	2	_	2	_	1	3	1	-	_
Aspergillus versicolor	Czapek	4	1	2	_	1	1	_	-	_	4	2	5	3	_	_	3	3	-	_
	Sabouraud	4	2	2	_	1	2	1	-	_	1	2	1	1	_	1	5	2	2	_
Penicillium	Czapek	4	-	-	-	1	4	-	-	-	3	2	-	-	-	-	5	4	-	_
diverse	Sabouraud	2	_	-	-	3	-	-	-	-	1	2	-	-	-	-	3	3	-	_
Penicillium	Czapek	-	-	1	_	_	-	_	-	-	-	-	-	-	2	-	3	-	1	-
expansum	Sabouraud	1	_	1	_	1	_	-	-	-	_	1	-	2	-	_	3	1	-	-
	Czapek	2	2	1	_	_	_	-	-	-	1	1	3	-	_	_	4	4	-	_
sporium herbarum	Sabouraud	_	1	_	_	_	_	_	-	-	_	_	1	-	_	_	1	3	_	_
Fusarium	Czapek	_	1	2	1	3	_	_	1	-	-	_	-	1	_	_	2	2	-	_
oxysporum	Sabouraud	3	1	1	1	1	-	-	1	-	3	2	_	-	_	-	5	5	-	_
Fusarium	Czapek	2	2	2	-	1	-	-	1	-	4	_	_	-	_	-	4	-	-	_
tricinctum	Sabouraud	3	-	1	_	1	-	-	-	-	2	4	1	-	-	-	3	3	-	_
Fusarium	Czapek	1	2	2	-	1	-	-	4	_	1	1	_	1	-	-	5	_	_	_
cerealis	Sabouraud	2	1	-	_	_	_	_	_	_	1	2	_	_	_	_	3	4	_	_

^{*}Enzymes: 1) alkaline phosphatase, 2) esterase (C 4), 3) esterase lipase (C 8), 4) lipase (C 14), 5) leucine arylamidase, 6) valine arylamidase, 7) cystine arylamidase, 8) trypsin, 9) α -chymotrypsin, 10) acid phosphatase, 11) naphthol-as-bi-phosphohydrolase, 12) α -galactosidase, 13) β -galactosidase, 14) β -glucoridase, 15) α -glucosidase, 16) β -glucosidase, 17) N-acetylo- β -glucosaminidase, 18) α -mannosidase, 19) α -fucosidase; **0 – no activity, 1–5 – activity increases with color intensity (1 – low activity, 5 – high activity)

is given to compulsive smoking and genetic factors, but exposure to organic dust may increase the incidence of the disease [24].

Asthma is a chronic inflammatory disease of the airways. Bronchial asthma as a result of exposure to organic dust can occur in different environments (agriculture, food processing, contact with wood dust). The main etiological factors of asthma are dust antigens in plants: cereal, flour, wood, dust from flax and hemp, cotton, coffee, tea, tobacco, hops and dust of animal origin: fur, feathers, excrement, skin and other protein substances. Because of its insidious onset, asthma is often mistakenly diagnosed as chronic bronchitis and is therefore not treated or treated inappropriately [23, 27].

Conclusions

All the obtained results show a potential risk of people involved in agricultural work on the organic components of grain dust, which contain toxinogenic and allergenic molds, thus indicate a substantial degree of occupational exposure.

Health statistics show that most of occupational diseases reported in Polish farmers is caused by pathogens present in organic dusts. In Poland, lung diseases are more common in farmers than in the rest of the population, just as in other countries. Therefore, the problem is serious, of social background and appropriate preventive measures need to be taken.

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