

Fungal pathogens and antagonists in root-soil zone in organic and integrated systems of potato production

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Abstract: Occurrence of culturable Fungi and Oomycota in root-soil habitat of potato cv. Owacja in organic and integrated production systems at Osiny (northern Poland) was compared in 2008–2010. The densities of both pathogens were significantly greater in the organic system. The eudominant fungal taxa (with frequency > 10% in at least one habitat) included species of *Fusarium* + *Gibberella* + *Haematonectria*, *Penicillium*, *Phoma* and *Trichoderma*. The dominant taxa (with frequency 5–10%) included species from 13 genera. In the rhizoplane, rhizosphere and non-rhizosphere soil, the total density of potential pathogens was greater in the integrated system, and of potential antagonists in the organic system. Among eudominant and dominant pathogens, *Fusarium oxysporum* and *Gibellulopsis nigrescens* occurred at greater density in the integrated system and *Haematonectria haematococca* and *Phoma* spp. in the organic system. Among eudominant antagonists, *Trichoderma* species occurred at greater density in the organic system. The organic system provided more disease suppressive habitat than the integrated system. The occurrence of brown leaf spot and potato blight was however similar in both systems. The mean yield of organic potatoes (24.9 t · ha⁻¹) was higher than the mean organic potato yield in Poland (21.0 t · ha⁻¹) and similar to the mean in other European countries (Germany 25.1 t · ha⁻¹, Great Britain 25.0 t · ha⁻¹). The organic system, based on a 5-year rotation, with narrow-leafed lupin, white mustard and buckwheat as a cover crop, inorganic fertilization based on ground rock phosphate + potassium sulphate, and biological and chemical control of insects and diseases (*Bacillus thuringiensis* ssp. *tenebrionis* + copper hydroxide + copper oxychloride), may be recommended for use in central Europe.

Key words: biological control, cropping system, culturable Fungi and Oomycota, potato

Introduction

Potato (*Solanum tuberosum* L.) is the world's most important non-grain food commodity and the fourth main food crop, with 368 mln t produced in 2013 (Anonymous 2014). The highest production of potatoes has been in eastern and central Europe (about 130 mln t) but the most rapid expansion over the past few decades has occurred in southern and eastern Asia (China, India produced about 135 mln t). Its status as a cheap and plentiful crop results from its ability to grow in a wide variety of climates and localities. The potato is best known for its carbohydrate (starch) content. It also contains vitamins, minerals and several phytochemicals, such as carotenoids and natural phenols.

Potato can suffer from more than 20 soil-borne diseases (O'Brien and Rich 1979; Fiers *et al.* 2012; Anonymous 2015). Some can occur on tubers while some on other parts of the plant (Gudmestad *et al.* 2007) affecting crop growth and usually resulting in yield loss.

Potato soil-borne diseases are the outcome of the compatible interactions between a susceptible host plant, soil-borne pathogen and soil environment. Knowledge of the pathogens as well as factors influencing disease

incidence and severity is needed to establish efficient control strategies.

A specific microbial biomass is associated with the potato soil environment. The structure of microbial communities varies with plant age, factors relating to cultivar, nutritional status, and biotic and abiotic stresses (Krechel *et al.* 2002; Manici and Caputo 2009). About 10⁷ bacterial colony-forming units (cfu) · g⁻¹ of soil live in the potato rhizosphere and potato geocaulosphere (soil around tubers) (Lazarovits *et al.* 2007). Microbial communities contribute to disease-suppressive effects resulting in lower disease incidence or severity despite the presence of a susceptible host plant, pathogen and favorable climatic conditions (Messiha *et al.* 2007; Steinberg *et al.* 2007; Penton *et al.* 2014). They create two classical types of soil suppressiveness, general and specific suppression, which are a result of, respectively, general activity of the whole microbial biomass, and specific activity of certain individuals or groups of microorganisms (Alabouvette *et al.* 1996; Weller *et al.* 2002).

Organic farming relies on agricultural techniques that exclude the use of synthetic chemical pesticides and

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include or recommend organic fertilization. As a result, the plant and soil microbiological environment is quite different from that created by conventional practices and may induce different disease suppression. Changing from a conventional to an organic system involves facing critical disease problems during a transition period of about 5 years, although control of soil-borne diseases can be achieved in the long-term (Bruggen and Termorshuizen 2003). A thorough understanding of how and to what extent crop plants affect the microbiota of soil, and vice versa, is the key to success in both intensive and extensive agricultural systems (Berendsen *et al.* 2012).

The objective of this study was to compare the fungal communities in potato roots, rhizosphere and non-rhizosphere soil in organic and integrated systems of production. The pathogenic and antagonistic components of the communities are recognized and their significance is estimated. The potential soil suppressiveness in each system is evaluated. The observations were supposed to help in developing strategies for planned procedures used for biological control. This contributes to the general aim of promoting crop management that will improve the soil habitat, increase the soil disease suppression and sustainability and result in greater production profitability.

Materials and Methods

Potato (*S. tuberosum*) cv. Owacja was cultivated at Osiny, Poland (52.5371200 N, 21.3282600 E) over three growing seasons (2008–2010). The 8-ha field was divided into two parts; one was under an organic system (4 ha) while another part under integrated system (4 ha). In each system, potato was cultivated on 0.125-ha experimental sub-plots, in rotation. The soil was sandy loam with characteristics presented in Table 1. The average annual temperature at Osiny was 8.5–9.6°C with the highest in 2008/2009 and the average annual precipitation was 537.2–582.8 mm, with the wettest in 2009/2010. Crop management procedures used in the organic and integrated systems of potato production at Osiny in 2008–2010 are presented in Table 2.

Basal roots from 60 randomly chosen plants and 20 samples of non-rhizosphere soil of potato were collect-

ed from the A horizon (0–20 cm deep) along a diagonal transect across each of four 0.125-ha sub-plots in each system, at the emergence and flowering stages in 2008–2010.

Isolation of microorganisms

Sixty apical parts of the basal roots were rinsed in running water for 30 min, then shaken in distilled sterile water (3 × 10 min), dried in sterilized blotting paper, cut into 1-cm-long pieces and placed on Potato Dextrose Agar (PDA; 39 g Difco PDA, 1 l distilled water, pH 5.5) in Petri dishes. For isolation of Fungi and Oomycota from the rhizoplane, 10 g of 1-cm-long apical parts of the basal roots were rinsed and shaken in a mixture of 90 ml distilled sterile water + 30 g sterile quartz sand for 10 min. The suspension was serially diluted and 1 ml from each of 10⁻² and 10⁻³ dilutions was poured into a Petri dish and covered with liquid (50°C) Johnson-Martin's agar (JMA; 10 g glucose, 5 g peptone, 1 g KH₂PO₄, 0.5 g MgSO₄·7H₂O, 0.03 g rose bengal, 0.0025 g aureomycin, 20 g agar, 1 l distilled water). For isolation of Fungi and Oomycota from rhizosphere soil, the basal roots of 60 plants were shaken for collection of soil. The soil was mixed and 10 g was shaken with 90 ml distilled sterile water for 10 min. The suspension was serially diluted and 1 ml from each of 10⁻⁴ and 10⁻⁵ dilutions was poured into a Petri dish and covered with liquid JMA. For isolation of Fungi and Oomycota from non-rhizosphere soil, 1 g of soil was mixed with 149 g of sterile quartz sand and 0.02 g of mixture was put into a Petri dish and covered with liquid JMA. Thirty replicates were used for each dilution and for non-rhizosphere soil. Plates were incubated for 10–30 days at 25°C. All microorganisms were identified on the basis of their morphology and sporulation on the specific media (Lenc *et al.* 2012), according to Pitt (1979), Domsch *et al.* (1980), Kwaśna *et al.* (1991) and Seifert *et al.* (2011).

The density of microorganisms was defined as the number of cfu in a sample. Frequency was defined as the proportion of particular isolates in the total number of isolates. The diversity of a microbial community was defined as the number of species in a sample. A species, or group of related species, was considered as: (i) eudominant

Table 1. Chemical properties of soils at Osiny in 2008–2010

| Soil characteristics | Organic system | | | Integrated system | | |
|---|----------------|-----------|----------|-------------------|----------|------------|
| | 2008 | 2009 | 2010 | 2008 | 2009 | 2010 |
| pH in H ₂ O | 6.32 | 6.53 | 6.41 | 6.38 | 6.43 | 6.76 |
| Humus content [%] | 1.73 a | 1.54 | 1.63 b | 1.26 a | 1.44 | 1.32 b |
| Extractable soil nitrogen | | | | | | |
| spring | 170.00 abe | 103.00 af | 90.00 bg | 79.00 cde | 61.00 cf | 56.00 dg |
| autumn | 132.00 abf | 90.00 acg | 51.00 bc | 124.00 d | 125.00 e | 43.00 defg |
| NO ₃ + NH ₄ [mg · kg ⁻¹]* | | | | | | |
| Extractable soil phosphorus [mg · kg ⁻¹ **] | 11.59 ac | 12.87 bd | 9.53 abe | 17.77 c | 15.80 d | 18.93 e |
| Extractable soil potassium [mg · kg ⁻¹ **] | 8.91 ab | 7.80 c | 6.83 ad | 12.43 b | 14.90 c | 11.67 d |
| Extractable soil magnesium [mg · kg ⁻¹ ***] | 12.49 abc | 9.60 ad | 8.40 be | 7.37 c | 7.33 d | 12.23 e |

*analysed with the Kjeldahl method

**analysed with the Egner-Riehm method

***analysed with the Schachtschabel method

a, b, c, d, e, f, g – the same letter in a row shows statistically significant difference between years or systems according to two-way ANOVA (at p = 0.05 or p = 0.001)

Table 2. Crop management procedures used in the organic and the integrated systems of potato production at Osiny in 2008–2010

| Treatment | Organic system | Integrated system |
|--|---|--|
| Rotation | potato spring wheat white clover + forage grasses clover + forage grasses winter wheat | potato spring wheat faba bean winter wheat |
| Cover crop | 2008 – narrow-leaved lupin (120 kg · ha ⁻¹) + white mustard (10 kg · ha ⁻¹) + buckwheat (20 kg · ha ⁻¹) 2009 – narrow-leaved lupin (70 kg · ha ⁻¹) + white mustard (20 kg · ha ⁻¹) + buckwheat (50 kg · ha ⁻¹) 2010 – narrow-leaved lupin (120 kg · ha ⁻¹) + buckwheat (40 kg · ha ⁻¹) | 2008 – white mustard (10 kg · ha ⁻¹) + buckwheat (40 kg · ha ⁻¹) – – |
| Tillage | 2007 – autumn – first plough (8–10 cm), disc harrowing (2×), winter plough (24–26 cm) 2008 – spring – disc harrowing, cultivation with aggregate Becker, U-1014, rolling, disc harrowing (2×), rolling 2008 – autumn – post-harvest tillage (gruber), disc harrowing, winter plough (24–26 cm) 2009 – spring – disc harrowing, cultivation with aggregate Becker, U-1014, cultivation with aggregate Crone, Landini, ridging (2×) 2009 – autumn – first plough (8–10 cm), disc harrowing (2×), winter plough (22–24 cm) 2010 – spring – disc harrowing, cultivation with aggregate Landini, disc harrowing (2–), ridging (2–), disc harrowing, ridging | 2009 – autumn – post-harvest tillage (grubber 16–18 m), winter plough (22–24 cm) 2010 – spring – disc harrowing, cultivation with aggregate Becker, Landini, ridging (2×) |
| Organic fertilizers | compost – under potato, in October, 25–30 dt · ha ⁻¹ | |
| Inorganic fertilizers applied late March-early April | ground phosphate rock + potassium sulphate 2008 – 72.5 kg P ₂ O ₅ + 62.5 kg K ₂ O · ha ⁻¹ 2009 – 36.5 kg P ₂ O ₅ + 62.5 kg K ₂ O · ha ⁻¹ 2010 – 29 kg P ₂ O ₅ + 62.5 kg K ₂ O · ha ⁻¹ | Polifoska 8 2008 – 22 kg N + 66 kg P ₂ O ₅ + 66 kg K ₂ O · ha ⁻¹ 2009 – 15 kg NH ₄ + 50 kg P ₂ O ₅ + 75 kg K ₂ O · ha ⁻¹ 2010 – 20 kg NH ₄ + 60 kg P ₂ O ₅ + 60 kg K ₂ O · ha ⁻¹ |
| Pesticides | <i>Bacillus thuringiensis</i> subsp. <i>tenebrionis</i> (Novodor SC ^a) 2008 – 9 l · ha ⁻¹ in 4 doses in June–July 2009 – 7.5 l · ha ⁻¹ in 3 doses in June–July 2010 – 13 l · ha ⁻¹ in 4 doses in June–July Copper hydroxide (Funguran OH 50 WP ^b) 2008–2010 – 2.0 kg h · ha ⁻¹ in July Copper oxychloride (Miedzian Extra 350 SC ^b) 2008–2009 – 4 l · ha ⁻¹ in July; 2010 – 2 l · ha ⁻¹ in July | linuron + clomazon (Afalon Dispersive 450 SC ^c + Command 480 EC ^c) 2008–2010 – 1 l + 0.1 l · ha ⁻¹ in May tiametoxam (Actara 25 WG ^a) – 2008 – 160 g · ha ⁻¹ in two doese in June and July 2009 – 2010 – 80 g · ha ⁻¹ in June– July mancozeb + metalaxyl (Ridomil Gold MZ 68 WG ^b) 2008–2010 2 kg · ha ⁻¹ in July mancozeb + cymoxanil (Curzate M 72,5 WG ^b) 2008 –2010 2 kg · ha ⁻¹ in July |
| Removal of weeds | harrowing or manually during vegetation | harrowing |

Compost included solid cattle manure enriched with grasses and clover and provided 2.8 t · ha⁻¹ of organic matter and 0.1 t · ha⁻¹ of N; 0.04 t · ha⁻¹ of P; 0.1 t · ha⁻¹ of K; 0.25 t · ha⁻¹ of Ca; 0.27 t · ha⁻¹ of Mg; 0.005 t · ha⁻¹ of Na. Before planting (end of April), the soil was ploughed and cultivated to create a deep tilth. Tubers were planted in earthed-up ridges.

^ainsecticide; ^bfungicide; ^cherbicide

– with frequency > 10%; (ii) dominant – with frequency 5–10%; (iii) subdominant – with frequency 2–5%; (iv) recedent – with frequency 1–2%; (v) sub recedent – with frequency < 1% (Tischler 1949).

Assessment of potato diseases

The occurrence of brown leaf spot [*Alternaria alternata* (Fries) Keissler] and potato blight [*Phytophthora infestans* (Mont.) de Bary] was assessed on 100 plants collected from random sites along a diagonal transect across each of the four replicate 0.125-ha sub-plots in each system. Potatoes in both systems were given values from 1 to 9 based on increasing proportion of leaf area affected. Potato tuber yield ($t \cdot ha^{-1}$) was determined for each system.

Statistical analysis

Chemical properties (pH, and content of humus, nitrogen, phosphorus, potassium and magnesium) of the soil were compared between two systems (organic and integrated) and three years (2008, 2009 and 2010) by two-way ANOVA analyses (at $p = 0.05$ or $p = 0.001$). Density and diversity of fungi and potato tuber yield were compared between two systems and three years (2008, 2009 and 2010) by χ^2 tests. Relationships between density of fungi and soil chemical properties were estimated by Pearson's correlation coefficient. All analyses were done separately for roots, rhizoplane, rhizosphere soil and non-rhizosphere soil.

Results

Densities of Fungi + Oomycota in potato roots, rhizoplane, rhizosphere soil and non-rhizosphere soil in individual years were greater (often significantly, $p = 0.05$ or $p = 0.001$) in the integrated system (Table 3). Densities of Fungi + Oomycota in rhizoplane, rhizosphere soil and non-rhizosphere (organic system) were usually significantly greater, in 2009 than in 2008 or 2010.

Fungi and Oomycota identified from each system were grouped into dominance classes (Table 4) and, where possible, affiliated according to their known ecological functions, i.e. as potential pathogens or antago-

nists. Eudominants and dominants included Zygomycota and Ascomycota fungi. Subdominants, apart from Ascomycota, included also Oomycota represented by *Pythium* spp. The total frequencies of eudominants + dominants in roots, rhizoplane, and rhizosphere and non-rhizosphere soil were similar in the two systems (77.5–98.6%).

The total density of potential potato pathogens was greater in the integrated system, significantly ($p = 0.05$ or $p = 0.001$) in all habitats except roots (Table 5). Densities of *Alternaria alternata*, *Gibberella intricans*, *Haematonectria haematococca*, *Phoma* spp. and *Rhizopus* were usually significantly greater in the organic system, and densities of *C. coccodes* (mostly in rhizoplane and rhizosphere), *Cladosporium herbarum*, *Fusarium culmorum*, *F. oxysporum*, *Gibellulopsis nigrescens*, *Pythium* spp. and *Stemphylium* sp. were greater in the integrated system (Table 4; some taxa not shown separately).

The total density of potential antagonists was significantly greater in the organic system, ($p = 0.001$) for rhizoplane and rhizosphere soil (Table 5). Densities of Mortierellales + Mucorales and *Trichoderma* spp. were usually significantly greater in the organic system, and densities of *Chaetomium* spp., *Paecilomyces* + *Purpureocillium* and *Sarocladium strictum* in the integrated system (Table 4). Densities of *Penicillium* spp. were significantly greater in the integrated system.

The ratio of pathogen density to antagonist density was less in the organic system than in the integrated system in all habitats (Table 5).

Acidity of soil was similar in the two systems (Table 1). Humus, nitrogen and magnesium contents were usually significantly higher in the organic system and phosphorus and potassium content in the integrated system (Table 1). In roots, rhizoplane, rhizosphere and non-rhizosphere soil there was no correlation between density of fungi and soil pH. There was negative correlation between mean density of fungi and humus content ($r = -0.8356, -0.6471, -0.5853, -0.7716$, respectively in roots, rhizoplane, rhizosphere and non-rhizosphere soil; $p < 0.0001$), nitrogen content ($r = -0.4522, -0.8005, -0.5615, -0.3411$; $p < 0.0001$) and magnesium content ($r = -0.0704, -0.6121, -0.6752, -0.5878$; $p < 0.0001$), and positive correlation between mean density of fungi and phosphorus content ($r = 0.7319, 0.3653,$

Table 3. Density (no. of colony forming units in a sample) of Fungi and Oomycota in potato cv. Owacja at Osiny in 2008–2010

| Habitat | Cropping system | 2008 | 2009 | 2010 | 2008–2010 total |
|----------------------|-----------------|-----------|---------|---------|-----------------|
| Roots | O | 88 | 93 | 90 | 271 |
| | I | 104 | 88 | 100 | 292 |
| Rhizoplane | O | 110 a,A,B | 189 A | 169 B | 468 |
| | I | 181 a | 172 | 171 | 524 |
| Rhizosphere soil | O | 93 a,A | 158 A,B | 109 B | 360 b |
| | I | 151 a,A | 156 B | 119 A,B | 426 b |
| Non-rhizosphere soil | O | 258 a,A | 322 A,B | 239 b,B | 819 a |
| | I | 497 a,A,B | 327 A | 302 b,B | 1126 a |

O – organic system; I – integrated system

a, b – the same lower case letter in a column shows statistically significant difference between systems according to χ^2 tests at $p = 0.001$ or $p = 0.05$ respectively

A, B – the same upper case letter in a row shows statistically significant difference between years according to χ^2 tests at $p = 0.001$ or $p = 0.05$

Table 4. Frequency (%) of taxa of fungi and Oomycota in potato cv. Owacja at Osiny in 2008–2010

| Taxon or group of taxa | Production system | In roots | In rhizoplane | In rhizosphere soil | In non-rhizosphere soil |
|---|-------------------|----------|---------------|---------------------|-------------------------|
| Eudominants (frequency >10% in at least one habitat) | | | | | |
| Ascomycota | | | | | |
| <i>Fusarium culmorum</i> (W. G. Sm.) Sacc. + <i>F. oxysporum</i> Schlecht. emend. Snyder. et Hans. + <i>F. sporotrichioides</i> Sherb. + <i>Gibberella avenacea</i> R.J. Cook + <i>G. intricans</i> Wollenw. + <i>G. tricineta</i> El-Gholl, McRitchie, Schoult. & Ridings + <i>G. zeae</i> (Schwein.) Petch + <i>Haematonectria haematococca</i> (Berk. & Broome) Samuels & Rossmann | O | 33.7 b | 29.8 a | 11.8 | 17.0 b |
| | I | 43.8 b | 41.8 a | 13.4 | 16.7 b |
| Including <i>F. oxysporum</i> | O | 7.4 a | 6.4 a | 2.2 a | 7.6 a |
| | I | 31.8 a | 29.0 a | 10.8 a | 11.5 a |
| Including <i>H. haematococca</i> | O | 25.5 a | 17.5 a | 5.3 a | 7.4 a |
| | I | 11.0 a | 2.7 a | 1.2 a | 3.6 a |
| <i>Penicillium</i> spp. | O | 6.3 a | 21.2 a | 24.7 a | 34.9 a |
| | I | 14.4 a | 33.6 a | 37.3 a | 49.6 a |
| <i>Phoma eupyrena</i> Sacc. + <i>P. glomerata</i> (Corda) Wollenw. & Hochapfel | O | 3.0 | 1.7 | 10.3 | 3.9 |
| | I | 0 | 1.1 | 7.3 | 2.1 |
| <i>Trichoderma hamatum</i> (Bon.) Bain. + <i>T. harzianum</i> Rifai + <i>T. koningii</i> Oudem. + <i>T. polysporum</i> (Link ex Pers.) Rifai + <i>T. viride</i> Pers. ex Gray | O | 51.3 b | 28.0 a | 35.0 a | 26.2 a |
| | I | 33.9 b | 11.9 a | 6.6 a | 9.7 a |
| Dominants (frequency 5–10% in at least one habitat) | | | | | |
| Zygomycota | | | | | |
| <i>Absidia glauca</i> Hagem + <i>Actinomucor elegans</i> (Eidam) C.R. Benj. & Hesselt. + <i>Mortierella alpina</i> Peyronel + <i>M. exigua</i> Linnem. + <i>M. hyalina</i> (Harz) W. Gams + <i>Mucor circinelloides</i> Tiegh. + <i>M. moelleri</i> (Vuill.) Lendn. + <i>M. mucedo</i> Fresen + <i>Rhizopus arrhizus</i> A. Fisch. | O | 0.7 | 5.3 b | 4.8 | 2.2 a |
| | I | 0.6 | 1.6 b | 3.3 | 4.3 a |
| Ascomycota | | | | | |
| <i>Clonostachys rosea</i> f. <i>catenulata</i> (J.C. Gilman & E.V. Abbott) Schroers + <i>C. rosea</i> f. <i>rosea</i> (Link) Schroers Samuels. Seifert & W. Gams + <i>C. solani</i> f. <i>solani</i> (Harting) Schroers & W. Gams | O | 3.3 | 3.6 | 4.4 | 2.1 |
| | I | 5.9 | 4.0 | 3.7 | 1.9 |
| <i>Gibellulopsis nigrescens</i> (Pethybr.) Zare, W. Gams & Summerb. | O | – | – | 1.1 a | 0.9 |
| | I | – | 0.8 | 5.9 a | 2.1 |
| Subdominants (frequency 2–5% in at least one habitat) | | | | | |
| Ascomycota | | | | | |
| <i>Alternaria alternata</i> (Fr.) Keissl. + <i>Cladosporium herbarum</i> (Pers.) Link + <i>Stemphylium</i> sp. | O | 0.7 | 1.9 | – | 1.3 |
| | I | 0.3 | 2.1 | 0.2 | 1.0 |
| <i>Chaetomium funicola</i> Cooke + <i>Ch. indicum</i> Corda | O | – | 0.2 | 0.6 | 3.4 |
| | I | – | 0.4 | 1.4 | 3.8 |
| <i>Gliomastix cerealis</i> (P. Karst.) C.H. Dickinson + <i>G. murorum</i> (Corda) S. Hughes | O | – | – | 0.3 a | 1.5 |
| | I | – | 0.4 | 3.1 a | 0.7 |
| <i>Paraconiothyrium fückelii</i> (Sacc.) Verkley & Gruyter | O | – | 2.8 | 0.3 | 0.1 |
| | I | – | – | 0.5 | 0.2 |
| <i>Paecilomyces niveus</i> Stolk & Samson + <i>P. variotii</i> Bainier + <i>Paecilomyces</i> spp. + <i>Purpureocillium lilacinum</i> (Thom) Luangsa-ard, Houbraken, Hywel-Jones & Samson | O | – | 0.2 | 0.6 a | 1.2 |
| | I | – | 1.3 | 4.0 a | 1.8 |
| <i>Sarocladium strictum</i> (W. Gams) Summerb. | O | 0.4 | 1.9 | 0.6 | 0.1 |
| | I | 1.0 | 0.2 | 2.1 | 0.9 |
| Oomycota | | | | | |
| <i>Pythium</i> spp. | O | – | – | – | – |
| | I | – | – | 2.1 | 0.1 |
| Recedents (frequency 1–2% in at least one habitat) | | | | | |
| <i>Colletotrichum coccodes</i> (Wallr.) S.J. Hughes | O | – | 0.4 | 0.6 | – |
| | I | – | 0.6 | 1.2 | – |

Table 4. Frequency (%) of taxa of fungi and Oomycota in potato cv. Owacja at Osiny in 2008–2010 – continuation

| Taxon or group of taxa | Production system | In roots | In rhizoplane | In rhizosphere soil | In non-rhizosphere soil |
|---|-------------------|----------|---------------|---------------------|-------------------------|
| <i>Myrothecium roridum</i> Tode | O | – | 0.9 | – | – |
| | I | – | 0.2 | 1.2 | – |
| Non-sporulating mycelia | O | – | 0.4 | 0.6 | 1.7 |
| | I | – | – | 0.9 | 1.1 |
| Total frequencies of eudominants + dominants | O | 98.3 | 89.6 | 92.1 | 87.2 |
| | I | 98.6 | 94.8 | 77.5 | 86.4 |
| Total frequencies of eudominants + dominants + subdominants | O | 99.4 | 96.6 | 94.5 | 94.8 |
| | I | 99.9 | 99.2 | 90.9 | 94.9 |

Subprecedents (frequency 0–1%): *Acrostalagmus luteoalbus* (Link) Zare, W. Gams & Schroers, *Arthrinium phaeospermum* (Corda) M.B. Ellis, *Arthrobotrys arthrobotryoides* (Berl.) Lindau, *Aspergillus fumigatus* Fresen., *A. niger* Tiegh., *Aspergillus* spp., *Cephalotrichum microsporium* (Sacc.) P.M. Kirk, *Dendryphon nanum* (Nees) S. Hughes, *Epicoccum nigrum* Link, *Gonytrichum macrocladum* (Sacc.) S. Hughes, *Gymnoascus reessii* Baran., *Humicola grisea* Traaen, *Leptosphaeria coniothyrium* (Fuckel) Sacc., *Melanospora* spp., *Microascus brevicaulis* S.P. Abbott, *Monographella nivalis* (Schaffnit) E. Müll., *Papulaspora irregularis* Hotson, *Pseudogymnoascus pannorum* (Link) Minnis & D.L. Lindner, *Septonema* sp., *Sporothrix* spp., *Thanatephorus cucumeris* (A.B. Frank) Donk, *Thysanophora penicillioides* (Roum.) W.B. Kendr., *Torulomyces indicus* (S.B. Saksena) M.H. Hashmi, W.B. Kendr. & E.B.G. Jones, *Trichothecium roseum* (Pers.) Link, *Umbelopsis vinacea* (Dixon-Stew.) Arx

In bold – potato pathogens: *A. alternata* + *Cladosporium* + *Stemphylium* sp., *C. coccodes* (Heilmann *et al.* 2006; Anonymous 2015), *Fusarium* spp. (Secor and Salas 2001; Peters *et al.* 2008 a, b; Estrada *et al.* 2010), *G. nigrescens* (Isaac and Harrison 1968), *Phoma* spp. (Adams 1983), *Penicillium* spp. (Rich 1983), *Pythium* spp. (O'Brien and Rich 1979), *Rhizopus* spp. (Clark and Moyer 1988).

Pathogens antagonists include: *Chaetomium* spp., *Mortierella* spp., *P. pannorum* (Tagawa *et al.* 2010), *Coniothyrium* spp. (Cheng *et al.* 2003). *S. strictum* (Rivera-Varas *et al.* 2007), *P. lilacinum* (Jacobs *et al.* 2003).

Stimulants of plant growth include *G. murorum* (Khan *et al.* 2009).

Nematophagous fungi with potential for controlling potato cyst nematodes include *Paecilomyces* + *Purpureocillium* (Jacobs *et al.* 2003).

O – organic system; I – integrated system

a, b – the same letter in a column shows statistically significant difference between systems according to χ^2 tests at $p = 0.001$ or $p = 0.05$ respectively

Table 5. Density of potential pathogens and antagonists in potato cv. Owacja at Osiny in 2008–2010

| Category | Cropping system | Roots | Rhizoplane | Rhizosphere soil | Non-rhizosphere soil |
|---|-----------------|-------|------------|------------------|----------------------|
| Density of pathogens (no. of cfu in a sample) | O | 101 | 158 a | 86 b | 189 b |
| | I | 129 | 248 a | 128 b | 248 b |
| Total frequency of pathogens [%] | O | 37.4 | 33.8 | 23.8 | 23.1 |
| | I | 44.1 | 47.3 | 30.1 | 22.0 |
| Diversity of pathogens (no. of species) | O | 8 | 12 | 14 | 9 |
| | I | 5 | 12 | 12 | 14 |
| Density of antagonists (no. of cfu in a sample) | O | 151 | 184 a | 166 a | 288 |
| | I | 121 | 102 a | 90 a | 252 |
| Total frequency of antagonists [%] | O | 55.7 | 39.2 | 46.0 | 35.2 |
| | I | 41.4 | 19.4 | 21.1 | 22.4 |
| Diversity of antagonists (no. of species) | O | 5 | 11 | 14 | 16 |
| | I | 8 | 12 | 11 | 18 |
| Ratio of pathogen: antagonist density | O | 0.7 | 0.9 | 0.5 | 0.7 |
| | I | 1.1 | 2.4 | 1.4 | 1.0 |
| Ratio of pathogen: antagonist density in cv. Drop (Lenc <i>et al.</i> 2012) | O | 8.9 | 6.0 | 1.9 | 1.5 |
| | I | 9.3 | 0.9 | 1.3 | 1.1 |

O – organic system; I – integrated system

a, b – the same letter in a column shows statistically significant difference between systems according to χ^2 tests at $p = 0.001$ or $p = 0.05$ respectively

Table 6. Incidence of diseases and yield of potato cv. Owacja at Osiny in 2008–2010

| Fungus | cropping system | Level of resistance to | | | Mean 2008–2010 |
|---|-----------------|------------------------|----------|------------|----------------|
| | | 2008 | 2009 | 2010 | |
| <i>Alternaria alternata</i> (brown leaf spot) | O | 7.0 | 8.5 | 7.5 | 7.7 |
| | I | 7.5 | 7.5 | 8.0 | 7.7 |
| <i>Phytophthora infestans</i> (potato late blight) | O | 7.5 | 7.0 | 7.5 | 7.3 |
| | I | 8.0 | 8.5 | 8.0 | 8.2 |
| Yield of potato tubers [t · ha ⁻¹] | O | 26.7 a | 26.0 a | 22.1 a | 24.9 a |
| | I | 46.6 a,A | 47.6 a,B | 25.9 b,A,B | 40.0 a |

O – organic system; I – integrated system

a, b – the same letter in a column shows statistically significant difference between systems according to χ^2 tests at $p = 0.05$

A, B – the same letter in a row shows statistically significant difference between years according to χ^2 tests at $p = 0.05$

0.4332, 0.6459; $p < 0.0001$). In rhizosphere and non-rhizosphere soil there was moderate positive correlation between mean density of fungi and potassium content ($r = 0.4747$, $r = 0.5253$; $p < 0.0001$).

Level of resistance to brown leaf spot and potato blight, based on disease evaluation, was 7.0–8.5 (Table 6). The mean level of resistance to brown leaf spot was similar in the two systems. The mean level of resistance to potato blight was higher in the integrated system. Yield in the organic system (22.1–26.7 t · ha⁻¹) was similar in all three years and lower than in the integrated system (not significantly in 2010). Yield in the integrated system was significantly lower in 2010 than in 2008 and 2009.

Discussion

Potatoes are generally an undemanding vegetable crop to grow, but they are not very easy using organic growing practices. They are susceptible to a few insect pests and diseases that can be a challenge for organic management. They also require relatively high levels of available nutrients in order to obtain good yields.

The exact mechanisms of natural and spontaneous control of root disease in organic or integrated farming are not known. It is generally assumed, however, that organic amendments suppress root diseases by: (i) increasing the general level of microbial activity, resulting in increased competition and/or antagonism in the rhizosphere and creating general and specific suppression; (ii) stimulation of plants by vesicular-arbuscular mycorrhiza (VAM) colonization of roots. Measuring microbial biomass and diversity (community structure) has been recommended as a biological indicator of soil quality and soil suppressiveness (Kennedy and Smith 1995; Yao *et al.* 2000) and has been employed in national and international monitoring programmes.

Organic farming has been shown to stimulate the growth and development of large and diverse soil microbial communities with potential for efficient nutrient cycling and disease suppression (Fraser *et al.* 1988; Mäder *et al.* 1995; Yeates *et al.* 1997; Gunapala and Scow 1998; Abawi and Widmer 2000; Shannon *et al.* 2002; Sadowski *et al.* 2002, 2003; Lenc 2006, 2009; Lenc *et al.* 2011). The increase in microbial biomass can be 10–26% (Fraser *et al.* 1994). How-

ever, the organic system described here resulted in smaller densities of Oomycota and fungi in potato roots/soil (but particularly in soil) compared with the integrated system. These results agree with earlier studies of Lenc *et al.* (2012) carried out earlier in the same location (Osiny), in 2005–2007, but on a different potato cultivar (cv. Drop). As previously reported (Lenc *et al.* 2012), a lower density of fungi was often associated with non-significantly higher soil acidity and higher amounts of humus, nitrogen and magnesium, but not of phosphorus or potassium. Results of earlier studies on effects of soil acidity on the growth of fungi are contradictory. Yamanaka (2003) reported that many saprotrophic species grew well at pH 7, while Blagodatskaya and Anderson (1998) and Bååth and Anderson (2003) found that soil pH could have either positive or negative effects on densities of fungi and bacteria.

The negative correlation between density of fungi and content of humus, nitrogen and magnesium observed agrees with some other studies in which fungal biomass was found to decrease as a consequence of application of organic and inorganic fertilization (Bardgett *et al.* 1999; Bloem and Vos 2004; Bittman *et al.* 2005; de Vries *et al.* 2006). However, a positive correlation between density of fungi and content of organic matter has also been reported (Alexander 1977; Henriksen and Breland 1999; Vinten *et al.* 2002). Differences result from the type of studied habitat. The microbiological response to organic matter depends on its C : N ratio. An increased C : N ratio stimulates, while decreased C : N ratio inhibits fungal growth, which is affected by accumulation of nitrogen. Organic carbon (C) in soil is transformed, stored and respired by microorganisms. Increased N content, leading to decreased C : N ratio, decreases microbial respiration, which is followed by decrease in microbial biomass (Spohn 2015).

The cover crop applied in Osiny included mostly narrow-leaved lupin, which, like other legumes, fixes nitrogen in a symbiotic interaction with different bacteria in its rhizosphere. This provision of nitrogen decreased the C : N ratio.

The humus content greater in the organic system than in the integrated system resulted from the type of cultivated cover crop. The succulent leaves of the annual legume (narrow-leaved lupin), white mustard and buckwheat break down quickly and release nitrogen and other

nutrients, but the tough and fibrous stems and roots, particularly of narrow-leaved lupin, after repeated cultivation (in 2008, 2009 and 2010), contributed to humus accumulation.

The inhibitory effect of magnesium and stimulatory effect of potassium on growth of fungi, suggested by the present results, has been reported previously (Jones and Jennings 1964).

White mustard used as a cover crop in the organic system may inhibit growth of fungi, including pathogenic *F. oxysporum*. It has been suggested that fungal suppression results from glucosinolate degradation into biologically active sulphur-containing thiocyanates (Porter 1995; Gardiner *et al.* 1999). White mustard suppressed potato soil-borne pathogens (*Fusarium*, *Rhizoctonia* and *Verticillium* species) in studies of Collins *et al.* (2006).

Evaluation of soil suppressiveness in each system was attempted by affiliating most microorganisms according to their known ecological functions. Potential pathogens and antagonists of potato pathogens were quantified.

Potato pathogens included *A. alternata* (brown spot and black pit), *C. coccodes* (black dot), *Fusarium* spp. (*Fusarium* dry rot and *Fusarium* wilt), *G. nigrescens* (*Verticillium* wilt), *Penicillium* spp. (blue and green mould rots), *Phoma* spp. (gangrene) and *Pythium* spp. (leak).

Fusarium oxysporum and *H. haematococca* were the most frequent identified species. These results obtained from plants during the vegetation differ from density of fungi on dry-rotting tubers, where *F. avenaceum*, *F. coeruleum* (Sacc.) Booth, *F. culmorum*, *F. oxysporum*, *F. sambucinum* Fuckel, *F. sulphureum* Schltld. were found, with *F. sulphureum* and *F. coeruleum* being the most common and considered the most important (Wojciechowska-Kot and Kiszczak 1981; Kurzawińska and Klima 1999; Kurzawińska and Gajda 2002). However, *F. oxysporum* was found to be the most frequent pathogen in tubers in Michigan, USA (Gachango *et al.* 2012).

In addition to *Fusarium* spp., *A. alternata* and *C. coccodes*, with frequencies of 24–28.8% and 4–6.6% respectively, have been reported as the most common dominants on tubers with dry rot symptoms (Kurzawińska and Gajda 2002). These species occurred with low frequencies in the present study and were among the subdominants and recedents, respectively. The frequency of *C. coccodes* was greater in the integrated system.

The rhizosphere region in the two systems was also inhabited by beneficial microorganisms known to be effective in biological control (Curl and Truelove 1986). *Clonostachys*, *Trichoderma* and *Purpureocillium lilacinum* were the most widespread. By means of competition, antibiosis and mycoparasitism they are able to decrease the severity of black dot, stem canker, black scurf (*Rhizoctonia solani* Kuhn.) and *Verticillium* wilt (*Verticillium dahliae* Kleb.), as well as nematodes (Davide and Zorilla 1983, 1985; Schirmbock *et al.* 1994; Kubicek *et al.* 2001; Howell 2003; Jacobs *et al.* 2003; Harman *et al.* 2004 and Fiers *et al.* 2012). Another fungus such as, *Gliomastix murorum*, produces gibberellin and promotes plant growth (Khan *et al.* 2009). The beneficial effects of indigenous biological control have been observed after only four years of organic farming (van Bruggen 1995).

The potential fungal antagonists were among the e-dominants, dominants and subdominants, and their high density seems to be essential for soil suppressiveness. Fungal recedents and subcedents (with individual frequency < 2%) were represented by 0.1–9.1% of species, with the largest population in the soil. They seem to be unimportant, making no contribution to the major biological interrelationships. They may, however, participate in natural and spontaneous control of diseases through optimization of nutrient cycles, particularly during environmental disturbances.

The role of some of the detected fungi is very complex. A few usually considered to be pathogenic (*Fusarium* or *Penicillium* species) may be antagonistic to *Streptomyces* spp. (potato scab) (Tagawa *et al.* 2010) or parasitic on nematodes (Martinez-Beringola *et al.* 2013) and hence contribute to soil suppressiveness.

The organic system was more suppressive to potential soil-borne pathogens than the integrated system. It supported less growth of *C. coccodes*, *F. culmorum*, *F. oxysporum*, *G. nigrescens* and *Pythium*, but not *Alternaria*, *H. haematococca*, *Phoma* or *Rhizopus*, and stimulated the growth of *Trichoderma* but not of other potential antagonists, *Mortierella*, *Paecilomyces* or *S. strictum*. The results are partly, from the specific viewpoint of stimulation and antagonism, in agreement with an earlier study, using a different potato cultivar, in which the integrated system was found to be generally more suppressive to potential soil-borne pathogens (Lenc *et al.* 2012).

Despite the overall greater suppressiveness of the organic system to soil-borne pathogens, *A. alternata* was found at similar (low) frequency in the organic and integrated systems and brown leaf spot (*A. alternata*) and late blight (*Ph. infestans*) occurred at similar frequencies in the two systems. Fungicides based on copper (Funguran OH 50 WP, Miedzian Extra 350 SC), used in the organic system to control late blight with a dose 6 kg · ha⁻¹ per year (as regulated by EU), seem to be no less effective than other, synthetic fungicides used in the integrated system.

The mean yield of potatoes grown in the organic system at Osiny (24.9 t · ha⁻¹) was greater than the average yield of organically grown potatoes in Poland (21.0 t · ha⁻¹) and similar to those in other European countries (Germany, 25.1 t · ha⁻¹, Great Britain, 25.0 t · ha⁻¹, Anonymous 2007). The mean yield per hectare for organically grown potatoes was 62.3% of the yield achieved in the integrated system. In other European countries, potatoes grown in organic systems achieved only 46% of the yield in conventional systems (Anonymous 2007). It may be assumed, because of the lower frequency of pathogens, higher frequency of antagonists, absence of disease differences that would affect yield, and acceptable tuber yield, that effective disease suppression was operating in the organic system.

Ratios of pathogen to antagonist densities were consistently smaller in the organic system than in the integrated system. The differences were small in roots and non-rhizosphere soil but more distinct in rhizoplane and rhizosphere soil. In a similar study (Lenc *et al.* 2012), the pathogen : antagonist density ratios in roots and rhizoplane were much greater (6.0–9.3; Table 5), partly as a result of the high frequency of *C. coccodes* in roots. The ratios were similar in

rhizosphere and non-rhizosphere soils (1.1–1.9). The higher pathogen : antagonist ratio resulted in much lower crop which on average was only 10.1–15.0 t · ha⁻¹ (Lenc *et al.* 2012). Since the inconsistencies between the two studies occurred in roots and rhizoplane, it seems likely that the differences were associated with plant lines and their genetic traits (particularly different susceptibilities to *C. coccodes*). However, the different results may have resulted from differences in management practices (especially if those practices affected incidence of *C. coccodes*).

Effects of plant genotype on the rhizosphere microbiota have been reported previously (Overbeek and van Elsas 2008; Weinert *et al.* 2009; İnceoğlu *et al.* 2010). Although often difficult to measure, it is plausible that these differences result from differences between plant cultivar (genotype) in their root-released products (İnceoğlu *et al.* 2011, 2013).

Comparison of the present results with those of Lenz *et al.* (2012) raises the major question 'to what extent individual cultivars affect the rhizosphere microbiota'. The answers are of particular significance for development, use and current admission procedures of new cultivars and genetically modified plants (Bruinsma and van Veen 2003) and would define a theoretical area of 'normality' for soil used for potato cropping. Such 'normality' indicated by a 'normal operating range' (NOR) could, according to İnceoğlu *et al.* (2013), be used as a standard reference for comparison of new cultivars and lines.

The classical method of profiling microbial communities based on pure culture isolation and morphotyping was used because of the scale of the experiment and the limitations of molecular methods. These limitations concern incomplete cell lysis, DNA sorption to soil surfaces, extraction of humic contaminants and inhibitors and DNA degradation (Fatima *et al.* 2014). Only culture-based techniques allow discrimination between viable members of a community (essential for the evaluation of pathogen-antagonist relationships) and inactive (dead) ones. While the possibility that culture-based techniques may mask the presence of slow-growing and non-culturable species must be considered, it may be assumed that the culturable microbial community is representative of the soil habitat complex (including taxa that can not be cultured) and can be considered valid for comparison of two environments.

Conclusions

The organic system examined for potato cultivation (5-year rotation, cover crop of narrow-leafed lupin + white mustard + buckwheat, inorganic fertilization based on ground rock phosphate + potassium sulphate, and biological and chemical control of insects and diseases based on *Bacillus thuringiensis* subsp. *tenebrionis* + copper hydroxide + copper oxychloride) resulted in a more disease-suppressive root/rhizosphere microbiota and similar incidence of leaf disease to the integrated system. The organic system also produced acceptable tuber yields. This system may therefore be suitable for recommendation for organic potato production in central Europe.

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